# The OpenCV Reference Manual

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# INTRODUCTION

OpenCV (Open Source Computer Vision Library: http://opencv.willowgarage.com/wiki/) is an open-source BSD-licensed library that includes several hundreds of computer vision algorithms. The document describes the so-called OpenCV 2.x API, which is essentially a C++ API, as opposite to the C-based OpenCV 1.x API. The latter is described in opencv1x.pdf.

OpenCV has a modular structure, which means that the package includes several shared or static libraries. The following modules are available:

- **core** - a compact module defining basic data structures, including the dense multi-dimensional array `Mat` and basic functions used by all other modules.

- **imgproc** - an image processing module that includes linear and non-linear image filtering, geometrical image transformations (resize, affine and perspective warping, generic table-based remapping), color space conversion, histograms, and so on.

- **video** - a video analysis module that includes motion estimation, background subtraction, and object tracking algorithms.

- **calib3d** - basic multiple-view geometry algorithms, single and stereo camera calibration, object pose estimation, stereo correspondence algorithms, and elements of 3D reconstruction.

- **features2d** - salient feature detectors, descriptors, and descriptor matchers.

- **objdetect** - detection of objects and instances of the predefined classes (for example, faces, eyes, mugs, people, cars, and so on).

- **highgui** - an easy-to-use interface to video capturing, image and video codecs, as well as simple UI capabilities.

- **gpu** - GPU-accelerated algorithms from different OpenCV modules.

- ... some other helper modules, such as FLANN and Google test wrappers, Python bindings, and others.

The further chapters of the document describe functionality of each module. But first, make sure to get familiar with the common API concepts used thoroughly in the library.

## 1.1 API Concepts

### cv Namespace

All the OpenCV classes and functions are placed into the `cv` namespace. Therefore, to access this functionality from your code, use the `cv::` specifier or `using namespace cv;` directive:

```
#include "opencv2/core/core.hpp"
...
cv::Mat H = cv::findHomography(points1, points2, CV_RANSAC, 5);
...
```

or

```
#include "opencv2/core/core.hpp"
using namespace cv;
...
Mat H = findHomography(points1, points2, CV_RANSAC, 5 );
...
```

Some of the current or future OpenCV external names may conflict with STL or other libraries. In this case, use explicit namespace specifiers to resolve the name conflicts:

```
Mat a(100, 100, CV_32F);
randu(a, Scalar::all(1), Scalar::all(std::rand()));
cv::log(a, a);
a /= std::log(2.);
```

## Automatic Memory Management

OpenCV handles all the memory automatically.

First of all, `std::vector`, `Mat`, and other data structures used by the functions and methods have destructors that deallocate the underlying memory buffers when needed. This means that the destructors do not always deallocate the buffers as in case of `Mat`. They take into account possible data sharing. A destructor decrements the reference counter associated with the matrix data buffer. The buffer is deallocated if and only if the reference counter reaches zero, that is, when no other structures refer to the same buffer. Similarly, when a `Mat` instance is copied, no actual data is really copied. Instead, the reference counter is incremented to memorize that there is another owner of the same data. There is also the `Mat::clone` method that creates a full copy of the matrix data. See the example below:

```
// create a big 8Mb matrix
Mat A(1000, 1000, CV_64F);

// create another header for the same matrix;
// this is an instant operation, regardless of the matrix size.
Mat B = A;
// create another header for the 3-rd row of A; no data is copied either
Mat C = B.row(3);
// now create a separate copy of the matrix
Mat D = B.clone();
// copy the 5-th row of B to C, that is, copy the 5-th row of A
// to the 3-rd row of A.
B.row(5).copyTo(C);
// now let A and D share the data; after that the modified version
// of A is still referenced by B and C.
A = D;
// now make B an empty matrix (which references no memory buffers),
// but the modified version of A will still be referenced by C,
// despite that C is just a single row of the original A
B.release();

// finally, make a full copy of C. As a result, the big modified
// matrix will be deallocated, since it is not referenced by anyone
C = C.clone();
```

You see that the use of `Mat` and other basic structures is simple. But what about high-level classes or even user data types created without taking automatic memory management into account? For them, OpenCV offers the `Ptr<>` template class that is similar to `std::shared_ptr` from C++ TR1. So, instead of using plain pointers:

```
T* ptr = new T(...);
```

you can use:

```
Ptr<T> ptr = new T(...);
```

That is, `Ptr<T> ptr` encapsulates a pointer to a `T` instance and a reference counter associated with the pointer. See the `Ptr` description for details.

## Automatic Allocation of the Output Data

OpenCV deallocates the memory automatically, as well as automatically allocates the memory for output function parameters most of the time. So, if a function has one or more input arrays (`cv::Mat` instances) and some output arrays, the output arrays are automatically allocated or reallocated. The size and type of the output arrays are determined from the size and type of input arrays. If needed, the functions take extra parameters that help to figure out the output array properties.

Example:

```cpp
#include "cv.h"
#include "highgui.h"

using namespace cv;

int main(int, char**)
{
    VideoCapture cap(0);
    if(!cap.isOpened()) return -1;

    Mat frame, edges;
    namedWindow("edges",1);
    for(;;)
    {
        cap >> frame;
        cvtColor(frame, edges, CV_BGR2GRAY);
        GaussianBlur(edges, edges, Size(7,7), 1.5, 1.5);
        Canny(edges, edges, 0, 30, 3);
        imshow("edges", edges);
        if(waitKey(30) >= 0) break;
    }
    return 0;
}
```

The array `frame` is automatically allocated by the `>>` operator since the video frame resolution and the bit-depth is known to the video capturing module. The array `edges` is automatically allocated by the `cvtColor` function. It has the same size and the bit-depth as the input array. The number of channels is 1 because the color conversion code `CV_BGR2GRAY` is passed, which means a color to grayscale conversion. Note that `frame` and `edges` are allocated only once during the first execution of the loop body since all the next video frames have the same resolution. If you somehow change the video resolution, the arrays are automatically reallocated.

The key component of this technology is the `Mat::create` method. It takes the desired array size and type. If the array already has the specified size and type, the method does nothing. Otherwise, it releases the previously allocated data, if any (this part involves decrementing the reference counter and comparing it with zero), and then allocates a new

buffer of the required size. Most functions call the `Mat::create` method for each output array, and so the automatic output data allocation is implemented.

Some notable exceptions from this scheme are `cv::mixChannels`, `cv::RNG::fill`, and a few other functions and methods. They are not able to allocate the output array, so you have to do this in advance.

## Saturation Arithmetics

As a computer vision library, OpenCV deals a lot with image pixels that are often encoded in a compact, 8- or 16-bit per channel, form and thus have a limited value range. Furthermore, certain operations on images, like color space conversions, brightness/contrast adjustments, sharpening, complex interpolation (bi-cubic, Lanczos) can produce values out of the available range. If you just store the lowest 8 (16) bits of the result, this results in visual artifacts and may affect a further image analysis. To solve this problem, the so-called *saturation* arithmetics is used. For example, to store r, the result of an operation, to an 8-bit image, you find the nearest value within the 0..255 range:

$$I(x, y) = \min(\max(\text{round}(r), 0), 255)$$

Similar rules are applied to 8-bit signed, 16-bit signed and unsigned types. This semantics is used everywhere in the library. In C++ code, it is done using the `saturate_cast<>` functions that resemble standard C++ cast operations. See below the implementation of the formula provided above:

```
I.at<uchar>(y, x) = saturate_cast<uchar>(r);
```

where `cv::uchar` is an OpenCV 8-bit unsigned integer type. In the optimized SIMD code, such SSE2 instructions as `paddusb`, `packuswb`, and so on are used. They help achieve exactly the same behavior as in C++ code.

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**Note:** Saturation is not applied when the result is 32-bit integer.

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## Fixed Pixel Types. Limited Use of Templates

Templates is a great feature of C++ that enables implementation of very powerful, efficient and yet safe data structures and algorithms. However, the extensive use of templates may dramatically increase compilation time and code size. Besides, it is difficult to separate an interface and implementation when templates are used exclusively. This could be fine for basic algorithms but not good for computer vision libraries where a single algorithm may span thousands lines of code. Because of this and also to simplify development of bindings for other languages, like Python, Java, Matlab that do not have templates at all or have limited template capabilities, the current OpenCV implementation is based on polymorphism and runtime dispatching over templates. In those places where runtime dispatching would be too slow (like pixel access operators), impossible (generic `Ptr<>` implementation), or just very inconvenient (`saturate_cast<>()`) the current implementation introduces small template classes, methods, and functions. Anywhere else in the current OpenCV version the use of templates is limited.

Consequently, there is a limited fixed set of primitive data types the library can operate on. That is, array elements should have one of the following types:

- 8-bit unsigned integer (uchar)
- 8-bit signed integer (schar)
- 16-bit unsigned integer (ushort)
- 16-bit signed integer (short)
- 32-bit signed integer (int)
- 32-bit floating-point number (float)
- 64-bit floating-point number (double)

- a tuple of several elements where all elements have the same type (one of the above). An array whose elements are such tuples, are called multi-channel arrays, as opposite to the single-channel arrays, whose elements are scalar values. The maximum possible number of channels is defined by the `CV_CN_MAX` constant, which is currently set to 512.

For these basic types, the following enumeration is applied:

```
enum { CV_8U=0, CV_8S=1, CV_16U=2, CV_16S=3, CV_32S=4, CV_32F=5, CV_64F=6 };
```

Multi-channel (n-channel) types can be specified using the following options:

- `CV_8UC1` ... `CV_64FC4` constants (for a number of channels from 1 to 4)

- `CV_8UC(n)` ... `CV_64FC(n)` or `CV_MAKETYPE(CV_8U, n)` ... `CV_MAKETYPE(CV_64F, n)` macros when the number of channels is more than 4 or unknown at the compilation time.

---

**Note:** `CV_32FC1 == CV_32F`, `CV_32FC2 == CV_32FC(2) == CV_MAKETYPE(CV_32F, 2)`, and `CV_MAKETYPE(depth, n) == ((x&7)<<3) + (n-1)`. This means that the constant type is formed from the `depth`, taking the lowest 3 bits, and the number of channels minus 1, taking the next `log2(CV_CN_MAX)` bits.

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Examples:

```
Mat mtx(3, 3, CV_32F); // make a 3x3 floating-point matrix
Mat cmtx(10, 1, CV_64FC2); // make a 10x1 2-channel floating-point
                           // matrix (10-element complex vector)
Mat img(Size(1920, 1080), CV_8UC3); // make a 3-channel (color) image
                                    // of 1920 columns and 1080 rows.
Mat grayscale(image.size(), CV_MAKETYPE(image.depth(), 1)); // make a 1-channel image of
                                                            // the same size and same
                                                            // channel type as img
```

Arrays with more complex elements cannot be constructed or processed using OpenCV. Furthermore, each function or method can handle only a subset of all possible array types. Usually, the more complex the algorithm is, the smaller the supported subset of formats is. See below typical examples of such limitations:

- The face detection algorithm only works with 8-bit grayscale or color images.

- Linear algebra functions and most of the machine learning algorithms work with floating-point arrays only.

- Basic functions, such as `cv::add`, support all types.

- Color space conversion functions support 8-bit unsigned, 16-bit unsigned, and 32-bit floating-point types.

The subset of supported types for each function has been defined from practical needs and could be extended in future based on user requests.

## InputArray and OutputArray

Many OpenCV functions process dense 2-dimensional or multi-dimensional numerical arrays. Usually, such functions take cpp:class:*Mat* as parameters, but in some cases it's more convenient to use `std::vector<>` (for a point set, for example) or `Matx<>` (for 3x3 homography matrix and such). To avoid many duplicates in the API, special "proxy" classes have been introduced. The base "proxy" class is `InputArray`. It is used for passing read-only arrays on a function input. The derived from `InputArray` class `OutputArray` is used to specify an output array for a function. Normally, you should not care of those intermediate types (and you should not declare variables of those types explicitly) - it will all just work automatically. You can assume that instead of `InputArray`/`OutputArray` you can always use Mat, `std::vector<>`, `Matx<>`, `Vec<>` or `Scalar`. When a function has an optional input or output array, and you do not have or do not want one, pass `cv::noArray()`.

## Error Handling

OpenCV uses exceptions to signal critical errors. When the input data has a correct format and belongs to the specified value range, but the algorithm cannot succeed for some reason (for example, the optimization algorithm did not converge), it returns a special error code (typically, just a boolean variable).

The exceptions can be instances of the `cv::Exception` class or its derivatives. In its turn, `cv::Exception` is a derivative of `std::exception`. So it can be gracefully handled in the code using other standard C++ library components.

The exception is typically thrown either using the `CV_Error(errcode, description)` macro, or its printf-like `CV_Error_(errcode, printf-spec, (printf-args))` variant, or using the `CV_Assert(condition)` macro that checks the condition and throws an exception when it is not satisfied. For performance-critical code, there is `CV_DbgAssert(condition)` that is only retained in the Debug configuration. Due to the automatic memory management, all the intermediate buffers are automatically deallocated in case of a sudden error. You only need to add a try statement to catch exceptions, if needed:

```cpp
try
{
    ... // call OpenCV
}
catch( cv::Exception& e )
{
    const char* err_msg = e.what();
    std::cout << "exception caught: " << err_msg << std::endl;
}
```

## Multi-threading and Re-enterability

The current OpenCV implementation is fully re-enterable. That is, the same function, the same *constant* method of a class instance, or the same *non-constant* method of different class instances can be called from different threads. Also, the same `cv::Mat` can be used in different threads because the reference-counting operations use the architecture-specific atomic instructions.

# CORE. THE CORE FUNCTIONALITY

## 2.1 Basic Structures

### DataType

**class DataType**

Template "trait" class for OpenCV primitive data types. A primitive OpenCV data type is one of `unsigned char`, `bool`, `signed char`, `unsigned short`, `signed short`, `int`, `float`, `double`, or a tuple of values of one of these types, where all the values in the tuple have the same type. Any primitive type from the list can be defined by an identifier in the form `CV_<bit-depth>{U|S|F}C(<number_of_channels>)`, for example: `uchar ~ CV_8UC1`, 3-element floating-point tuple ~ `CV_32FC3`, and so on. A universal OpenCV structure that is able to store a single instance of such a primitive data type is `Vec`. Multiple instances of such a type can be stored in a `std::vector`, `Mat`, `Mat_`, `SparseMat`, `SparseMat_`, or any other container that is able to store `Vec` instances.

The `DataType` class is basically used to provide a description of such primitive data types without adding any fields or methods to the corresponding classes (and it is actually impossible to add anything to primitive C/C++ data types). This technique is known in C++ as class traits. It is not `DataType` itself that is used but its specialized versions, such as:

```
template<> class DataType<uchar>
{
    typedef uchar value_type;
    typedef int work_type;
    typedef uchar channel_type;
    enum { channel_type = CV_8U, channels = 1, fmt='u', type = CV_8U };
};
...
template<typename _Tp> DataType<std::complex<_Tp> >
{
    typedef std::complex<_Tp> value_type;
    typedef std::complex<_Tp> work_type;
    typedef _Tp channel_type;
    // DataDepth is another helper trait class
    enum { depth = DataDepth<_Tp>::value, channels=2,
        fmt=(channels-1)*256+DataDepth<_Tp>::fmt,
        type=CV_MAKETYPE(depth, channels) };
};
...
```

The main purpose of this class is to convert compilation-time type information to an OpenCV-compatible data type identifier, for example:

```
// allocates a 30x40 floating-point matrix
Mat A(30, 40, DataType<float>::type);

Mat B = Mat_<std::complex<double> >(3, 3);
// the statement below will print 6, 2 /*, that is depth == CV_64F, channels == 2 */
cout << B.depth() << ", " << B.channels() << endl;
```

So, such traits are used to tell OpenCV which data type you are working with, even if such a type is not native to OpenCV. For example, the matrix B initialization above is compiled because OpenCV defines the proper specialized template class `DataType<complex<_Tp> >`. This mechanism is also useful (and used in OpenCV this way) for generic algorithms implementations.

## Point_

**class** `Point_`

Template class for 2D points specified by its coordinates $x$ and $y$. An instance of the class is interchangeable with C structures, `CvPoint` and `CvPoint2D32f`. There is also a cast operator to convert point coordinates to the specified type. The conversion from floating-point coordinates to integer coordinates is done by rounding. Commonly, the conversion uses this operation for each of the coordinates. Besides the class members listed in the declaration above, the following operations on points are implemented:

```
pt1 = pt2 + pt3;
pt1 = pt2 - pt3;
pt1 = pt2 * a;
pt1 = a * pt2;
pt1 += pt2;
pt1 -= pt2;
pt1 *= a;
double value = norm(pt); // L2 norm
pt1 == pt2;
pt1 != pt2;
```

For your convenience, the following type aliases are defined:

```
typedef Point_<int> Point2i;
typedef Point2i Point;
typedef Point_<float> Point2f;
typedef Point_<double> Point2d;
```

Example:

```
Point2f a(0.3f, 0.f), b(0.f, 0.4f);
Point pt = (a + b)*10.f;
cout << pt.x << ", " << pt.y << endl;
```

## Point3_

**class** `Point3_`

Template class for 3D points specified by its coordinates $x$, $y$ and $z$. An instance of the class is interchangeable with the C structure `CvPoint2D32f`. Similarly to `Point_`, the coordinates of 3D points can be converted to another type. The vector arithmetic and comparison operations are also supported.

The following `Point3_<>` aliases are available:

```
typedef Point3_<int> Point3i;
typedef Point3_<float> Point3f;
typedef Point3_<double> Point3d;
```

## Size_

**class Size_**

Template class for specifying the size of an image or rectangle. The class includes two members called width and height. The structure can be converted to and from the old OpenCV structures CvSize and CvSize2D32f . The same set of arithmetic and comparison operations as for Point_ is available.

OpenCV defines the following Size_<> aliases:

```
typedef Size_<int> Size2i;
typedef Size2i Size;
typedef Size_<float> Size2f;
```

## Rect_

**class Rect_**

Template class for 2D rectangles, described by the following parameters:

- Coordinates of the top-left corner. This is a default interpretation of Rect_::x and Rect_::y in OpenCV. Though, in your algorithms you may count x and y from the bottom-left corner.

- Rectangle width and height.

OpenCV typically assumes that the top and left boundary of the rectangle are inclusive, while the right and bottom boundaries are not. For example, the method Rect_::contains returns true if

$$x \le pt.x < x + \texttt{width}, y \le pt.y < y + \texttt{height}$$

Virtually every loop over an image ROI in OpenCV (where ROI is specified by Rect_<int> ) is implemented as:

```
for(int y = roi.y; y < roi.y + rect.height; y++)
    for(int x = roi.x; x < roi.x + rect.width; x++)
    {
        // ...
    }
```

In addition to the class members, the following operations on rectangles are implemented:

- rect = rect ± point (shifting a rectangle by a certain offset)

- rect = rect ± size (expanding or shrinking a rectangle by a certain amount)

- rect += point, rect -= point, rect += size, rect -= size (augmenting operations)

- rect = rect1 & rect2 (rectangle intersection)

- rect = rect1 | rect2 (minimum area rectangle containing rect2 and rect3 )

- rect &= rect1, rect |= rect1 (and the corresponding augmenting operations)

- rect == rect1, rect != rect1 (rectangle comparison)

This is an example how the partial ordering on rectangles can be established (rect1 ⊆ rect2):

```
template<typename _Tp> inline bool
operator <= (const Rect_<_Tp>& r1, const Rect_<_Tp>& r2)
{
    return (r1 & r2) == r1;
}
```

For your convenience, the `Rect_<>` alias is available:

```
typedef Rect_<int> Rect;
```

## RotatedRect

**class RotatedRect**

The class represents rotated (i.e. not up-right) rectangles on a plane. Each rectangle is specified by the center point (mass center), length of each side (represented by cv::Size2f structure) and the rotation angle in degrees.

> **C++:** `RotatedRect::RotatedRect()`

> **C++:** `RotatedRect::RotatedRect`(const Point2f& **center**, const Size2f& **size**, float **angle**)

> **C++:** `RotatedRect::RotatedRect`(const CvBox2D& **box**)

>> **Parameters**

>>> **center** – The rectangle mass center.

>>> **size** – Width and height of the rectangle.

>>> **angle** – The rotation angle in a clockwise direction. When the angle is 0, 90, 180, 270 etc., the rectangle becomes an up-right rectangle.

>>> **box** – The rotated rectangle parameters as the obsolete CvBox2D structure.

> **C++:** void `RotatedRect::points`(Point2f **pts**[]) const

> **C++:** Rect `RotatedRect::boundingRect`() const

> **C++:** `RotatedRect::operator` CvBox2D() const

>> **Parameters**

>>> **pts** – The points array for storing rectangle vertices.
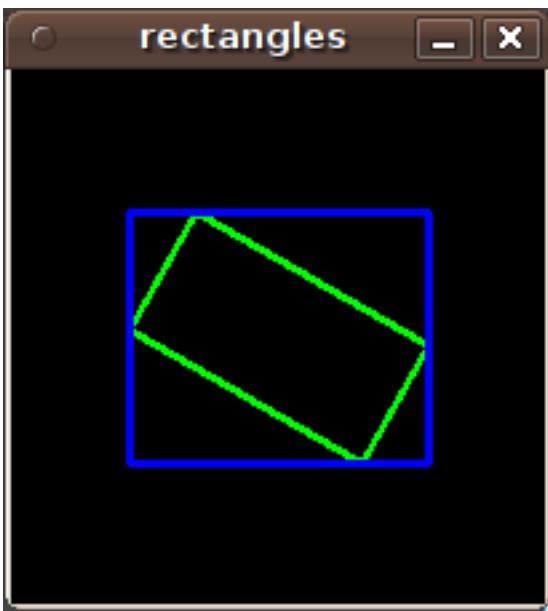
The sample below demonstrates how to use RotatedRect:

```
Mat image(200, 200, CV_8UC3, Scalar(0));
RotatedRect rRect = RotatedRect(Point2f(100,100), Size2f(100,50), 30);

Point2f vertices[4];
rRect.points(vertices);
for (int i = 0; i < 4; i++)
    line(image, vertices[i], vertices[(i+1)%4], Scalar(0,255,0));

Rect brect = rRect.boundingRect();
rectangle(image, brect, Scalar(255,0,0));

imshow("rectangles", image);
waitKey(0);
```

**See Also:**

CamShift(), fitEllipse(), minAreaRect(), CvBox2D

## TermCriteria

**class TermCriteria**

Template class defining termination criteria for iterative algorithms.

## Matx

**class Matx**

Template class for small matrices whose type and size are known at compilation time:

```cpp
template<typename _Tp, int m, int n> class Matx {...};

typedef Matx<float, 1, 2> Matx12f;
typedef Matx<double, 1, 2> Matx12d;
...
typedef Matx<float, 1, 6> Matx16f;
typedef Matx<double, 1, 6> Matx16d;

typedef Matx<float, 2, 1> Matx21f;
typedef Matx<double, 2, 1> Matx21d;
...
typedef Matx<float, 6, 1> Matx61f;
typedef Matx<double, 6, 1> Matx61d;

typedef Matx<float, 2, 2> Matx22f;
typedef Matx<double, 2, 2> Matx22d;
...
typedef Matx<float, 6, 6> Matx66f;
typedef Matx<double, 6, 6> Matx66d;
```

If you need a more flexible type, use `Mat` . The elements of the matrix M are accessible using the `M(i,j)` notation. Most of the common matrix operations (see also *Matrix Expressions* ) are available. To do an operation on `Matx` that is not implemented, you can easily convert the matrix to `Mat` and backwards.

```
Matx33f m(1, 2, 3,
          4, 5, 6,
          7, 8, 9);
cout << sum(Mat(m*m.t())) << endl;
```

## Vec

**class Vec**

Template class for short numerical vectors, a partial case of `Matx`:

```
template<typename _Tp, int n> class Vec : public Matx<_Tp, n, 1> {...};

typedef Vec<uchar, 2> Vec2b;
typedef Vec<uchar, 3> Vec3b;
typedef Vec<uchar, 4> Vec4b;

typedef Vec<short, 2> Vec2s;
typedef Vec<short, 3> Vec3s;
typedef Vec<short, 4> Vec4s;

typedef Vec<int, 2> Vec2i;
typedef Vec<int, 3> Vec3i;
typedef Vec<int, 4> Vec4i;

typedef Vec<float, 2> Vec2f;
typedef Vec<float, 3> Vec3f;
typedef Vec<float, 4> Vec4f;
typedef Vec<float, 6> Vec6f;

typedef Vec<double, 2> Vec2d;
typedef Vec<double, 3> Vec3d;
typedef Vec<double, 4> Vec4d;
typedef Vec<double, 6> Vec6d;
```

It is possible to convert `Vec<T,2>` to/from `Point_`, `Vec<T,3>` to/from `Point3_` , and `Vec<T,4>` to `CvScalar` or `Scalar_`. Use `operator[]` to access the elements of `Vec`.

All the expected vector operations are also implemented:

- `v1 = v2 + v3`

- `v1 = v2 - v3`

- `v1 = v2 * scale`

- `v1 = scale * v2`

- `v1 = -v2`

- `v1 += v2` and other augmenting operations

- `v1 == v2, v1 != v2`

- `norm(v1)` (euclidean norm)

The `Vec` class is commonly used to describe pixel types of multi-channel arrays. See `Mat` for details.

## Scalar_

**class Scalar_**

Template class for a 4-element vector derived from Vec.

```cpp
template<typename _Tp> class Scalar_ : public Vec<_Tp, 4> { ... };
```

```cpp
typedef Scalar_<double> Scalar;
```

Being derived from Vec<_Tp, 4> , Scalar_ and Scalar can be used just as typical 4-element vectors. In addition, they can be converted to/from CvScalar . The type Scalar is widely used in OpenCV to pass pixel values.

## Range

**class Range**

Template class specifying a continuous subsequence (slice) of a sequence.

```cpp
class Range
{
public:
    ...
    int start, end;
};
```

The class is used to specify a row or a column span in a matrix ( Mat ) and for many other purposes. Range(a,b) is basically the same as a:b in Matlab or a..b in Python. As in Python, start is an inclusive left boundary of the range and end is an exclusive right boundary of the range. Such a half-opened interval is usually denoted as $[start, end)$ .

The static method Range::all() returns a special variable that means "the whole sequence" or "the whole range", just like " : " in Matlab or " ... " in Python. All the methods and functions in OpenCV that take Range support this special Range::all() value. But, of course, in case of your own custom processing, you will probably have to check and handle it explicitly:

```cpp
void my_function(..., const Range& r, ....)
{
    if(r == Range::all()) {
        // process all the data
    }
    else {
        // process [r.start, r.end)
    }
}
```

## Ptr

**class Ptr**

Template class for smart reference-counting pointers

```cpp
template<typename _Tp> class Ptr
{
public:
    // default constructor
    Ptr();
    // constructor that wraps the object pointer
```

```
    Ptr(_Tp* _obj);
    // destructor: calls release()
    ~Ptr();
    // copy constructor; increments ptr's reference counter
    Ptr(const Ptr& ptr);
    // assignment operator; decrements own reference counter
    // (with release()) and increments ptr's reference counter
    Ptr& operator = (const Ptr& ptr);
    // increments reference counter
    void addref();
    // decrements reference counter; when it becomes 0,
    // delete_obj() is called
    void release();
    // user-specified custom object deletion operation.
    // by default, "delete obj;" is called
    void delete_obj();
    // returns true if obj == 0;
    bool empty() const;

    // provide access to the object fields and methods
    _Tp* operator -> ();
    const _Tp* operator -> () const;

    // return the underlying object pointer;
    // thanks to the methods, the Ptr<_Tp> can be
    // used instead of _Tp*
    operator _Tp* ();
    operator const _Tp*() const;
protected:
    // the encapsulated object pointer
    _Tp* obj;
    // the associated reference counter
    int* refcount;
};
```

The `Ptr<_Tp>` class is a template class that wraps pointers of the corresponding type. It is similar to `shared_ptr` that is part of the Boost library ( http://www.boost.org/doc/libs/1_40_0/libs/smart_ptr/shared_ptr.htm ) and also part of the C++0x standard.

This class provides the following options:

- Default constructor, copy constructor, and assignment operator for an arbitrary C++ class or a C structure. For some objects, like files, windows, mutexes, sockets, and others, a copy constructor or an assignment operator are difficult to define. For some other objects, like complex classifiers in OpenCV, copy constructors are absent and not easy to implement. Finally, some of complex OpenCV and your own data structures may be written in C. However, copy constructors and default constructors can simplify programming a lot. Besides, they are often required (for example, by STL containers). By wrapping a pointer to such a complex object `TObj` to `Ptr<TObj>`, you automatically get all of the necessary constructors and the assignment operator.

- *O(1)* complexity of the above-mentioned operations. While some structures, like `std::vector`, provide a copy constructor and an assignment operator, the operations may take a considerable amount of time if the data structures are large. But if the structures are put into `Ptr<>` , the overhead is small and independent of the data size.

- Automatic destruction, even for C structures. See the example below with `FILE*` .

- Heterogeneous collections of objects. The standard STL and most other C++ and OpenCV containers can store only objects of the same type and the same size. The classical solution to store objects of different types in the same container is to store pointers to the base class `base_class_t*` instead but then you loose the automatic

memory management. Again, by using Ptr<base_class_t>() instead of the raw pointers, you can solve the problem.

The Ptr class treats the wrapped object as a black box. The reference counter is allocated and managed separately. The only thing the pointer class needs to know about the object is how to deallocate it. This knowledge is encapsulated in the Ptr::delete_obj() method that is called when the reference counter becomes 0. If the object is a C++ class instance, no additional coding is needed, because the default implementation of this method calls delete obj; . However, if the object is deallocated in a different way, the specialized method should be created. For example, if you want to wrap FILE , the delete_obj may be implemented as follows:

```cpp
template<> inline void Ptr<FILE>::delete_obj()
{
    fclose(obj); // no need to clear the pointer afterwards,
                 // it is done externally.
}
...


// now use it:
Ptr<FILE> f(fopen("myfile.txt", "r"));
if(f.empty())
    throw ...;
fprintf(f, ....);
...
// the file will be closed automatically by the Ptr<FILE> destructor.
```

---

**Note:** The reference increment/decrement operations are implemented as atomic operations, and therefore it is normally safe to use the classes in multi-threaded applications. The same is true for Mat and other C++ OpenCV classes that operate on the reference counters.

---

## Mat

**class Mat**

OpenCV C++ n-dimensional dense array class

```cpp
class CV_EXPORTS Mat
{
public:
    // ... a lot of methods ...
    ...

    /*! includes several bit-fields:
         - the magic signature
         - continuity flag
         - depth
         - number of channels
     */
    int flags;
    //! the array dimensionality, >= 2
    int dims;
    //! the number of rows and columns or (-1, -1) when the array has more than 2 dimensions
    int rows, cols;
    //! pointer to the data
    uchar* data;

    //! pointer to the reference counter;
```

```
    // when array points to user-allocated data, the pointer is NULL
    int* refcount;

    // other members
    ...
};
```

The class `Mat` represents an n-dimensional dense numerical single-channel or multi-channel array. It can be used to store real or complex-valued vectors and matrices, grayscale or color images, voxel volumes, vector fields, point clouds, tensors, histograms (though, very high-dimensional histograms may be better stored in a `SparseMat`). The data layout of the array $M$ is defined by the array `M.step[]`, so that the address of element $(i_0, ..., i_{M.dims-1})$, where $0 \le i_k < M.size[k]$, is computed as:

$$addr(M_{i_0,...,i_{M.dims-1}}) = M.data + M.step[0] * i_0 + M.step[1] * i_1 + ... + M.step[M.dims - 1] * i_{M.dims-1}$$

In case of a 2-dimensional array, the above formula is reduced to:

$$addr(M_{i,j}) = M.data + M.step[0] * i + M.step[1] * j$$

Note that `M.step[i] >= M.step[i+1]` (in fact, `M.step[i] >= M.step[i+1]*M.size[i+1]`). This means that 2-dimensional matrices are stored row-by-row, 3-dimensional matrices are stored plane-by-plane, and so on. `M.step[M.dims-1]` is minimal and always equal to the element size `M.elemSize()`.

So, the data layout in `Mat` is fully compatible with `CvMat`, `IplImage`, and `CvMatND` types from OpenCV 1.x. It is also compatible with the majority of dense array types from the standard toolkits and SDKs, such as Numpy (ndarray), Win32 (independent device bitmaps), and others, that is, with any array that uses *steps* (or *strides*) to compute the position of a pixel. Due to this compatibility, it is possible to make a `Mat` header for user-allocated data and process it in-place using OpenCV functions.

There are many different ways to create a `Mat` object. The most popular options are listed below:

- Use the `create(nrows, ncols, type)` method or the similar `Mat(nrows, ncols, type[, fillValue])` constructor. A new array of the specified size and type is allocated. `type` has the same meaning as in the `cvCreateMat` method. For example, `CV_8UC1` means a 8-bit single-channel array, `CV_32FC2` means a 2-channel (complex) floating-point array, and so on.

  ```
  // make a 7x7 complex matrix filled with 1+3j.
  Mat M(7,7,CV_32FC2,Scalar(1,3));
  // and now turn M to a 100x60 15-channel 8-bit matrix.
  // The old content will be deallocated
  M.create(100,60,CV_8UC(15));
  ```

  As noted in the introduction to this chapter, `create()` allocates only a new array when the shape or type of the current array are different from the specified ones.

- Create a multi-dimensional array:

  ```
  // create a 100x100x100 8-bit array
  int sz[] = {100, 100, 100};
  Mat bigCube(3, sz, CV_8U, Scalar::all(0));
  ```

  It passes the number of dimensions =1 to the `Mat` constructor but the created array will be 2-dimensional with the number of columns set to 1. So, `Mat::dims` is always >= 2 (can also be 0 when the array is empty).

- Use a copy constructor or assignment operator where there can be an array or expression on the right side (see below). As noted in the introduction, the array assignment is an O(1) operation because it only copies the header and increases the reference counter. The `Mat::clone()` method can be used to get a full (deep) copy of the array when you need it.

- Construct a header for a part of another array. It can be a single row, single column, several rows, several columns, rectangular region in the array (called a *minor* in algebra) or a diagonal. Such operations are also O(1) because the new header references the same data. You can actually modify a part of the array using this feature, for example:

```
// add the 5-th row, multiplied by 3 to the 3rd row
M.row(3) = M.row(3) + M.row(5)*3;

// now copy the 7-th column to the 1-st column
// M.col(1) = M.col(7); // this will not work
Mat M1 = M.col(1);
M.col(7).copyTo(M1);

// create a new 320x240 image
Mat img(Size(320,240),CV_8UC3);
// select a ROI
Mat roi(img, Rect(10,10,100,100));
// fill the ROI with (0,255,0) (which is green in RGB space);
// the original 320x240 image will be modified
roi = Scalar(0,255,0);
```

Due to the additional `datastart` and `dataend` members, it is possible to compute a relative sub-array position in the main *container* array using `locateROI()`:

```
Mat A = Mat::eye(10, 10, CV_32S);
// extracts A columns, 1 (inclusive) to 3 (exclusive).
Mat B = A(Range::all(), Range(1, 3));
// extracts B rows, 5 (inclusive) to 9 (exclusive).
// that is, C ~ A(Range(5, 9), Range(1, 3))
Mat C = B(Range(5, 9), Range::all());
Size size; Point ofs;
C.locateROI(size, ofs);
// size will be (width=10,height=10) and the ofs will be (x=1, y=5)
```

As in case of whole matrices, if you need a deep copy, use the `clone()` method of the extracted sub-matrices.

- Make a header for user-allocated data. It can be useful to do the following:

  1. Process "foreign" data using OpenCV (for example, when you implement a DirectShow* filter or a processing module for `gstreamer`, and so on). For example:

  ```
  void process_video_frame(const unsigned char* pixels,
                           int width, int height, int step)
  {
      Mat img(height, width, CV_8UC3, pixels, step);
      GaussianBlur(img, img, Size(7,7), 1.5, 1.5);
  }
  ```

  2. Quickly initialize small matrices and/or get a super-fast element access.

  ```
  double m[3][3] = {{a, b, c}, {d, e, f}, {g, h, i}};
  Mat M = Mat(3, 3, CV_64F, m).inv();
  ```

  Partial yet very common cases of this *user-allocated data* case are conversions from `CvMat` and `IplImage` to `Mat`. For this purpose, there are special constructors taking pointers to `CvMat` or `IplImage` and the optional flag indicating whether to copy the data or not.

  Backward conversion from `Mat` to `CvMat` or `IplImage` is provided via cast operators `Mat::operator CvMat() const` and `Mat::operator IplImage()`. The operators do NOT copy the data.

```
IplImage* img = cvLoadImage("greatwave.jpg", 1);
Mat mtx(img); // convert IplImage* -> Mat
CvMat oldmat = mtx; // convert Mat -> CvMat
CV_Assert(oldmat.cols == img->width && oldmat.rows == img->height &&
    oldmat.data.ptr == (uchar*)img->imageData && oldmat.step == img->widthStep);
```

- Use MATLAB-style array initializers, `zeros()`, `ones()`, `eye()`, for example:

  ```
  // create a double-precision identity martix and add it to M.
  M += Mat::eye(M.rows, M.cols, CV_64F);
  ```

- Use a comma-separated initializer:

  ```
  // create a 3x3 double-precision identity matrix
  Mat M = (Mat_<double>(3,3) << 1, 0, 0, 0, 1, 0, 0, 0, 1);
  ```

  With this approach, you first call a constructor of the `Mat_` class with the proper parameters, and then you just put `<<` operator followed by comma-separated values that can be constants, variables, expressions, and so on. Also, note the extra parentheses required to avoid compilation errors.

Once the array is created, it is automatically managed via a reference-counting mechanism. If the array header is built on top of user-allocated data, you should handle the data by yourself. The array data is deallocated when no one points to it. If you want to release the data pointed by a array header before the array destructor is called, use `Mat::release()`.

The next important thing to learn about the array class is element access. This manual already described how to compute an address of each array element. Normally, you are not required to use the formula directly in the code. If you know the array element type (which can be retrieved using the method `Mat::type()` ), you can access the element $M_{ij}$ of a 2-dimensional array as:

```
M.at<double>(i,j) += 1.f;
```

assuming that M is a double-precision floating-point array. There are several variants of the method `at` for a different number of dimensions.

If you need to process a whole row of a 2D array, the most efficient way is to get the pointer to the row first, and then just use the plain C operator `[]` :

```
// compute sum of positive matrix elements
// (assuming that M isa double-precision matrix)
double sum=0;
for(int i = 0; i < M.rows; i++)
{
    const double* Mi = M.ptr<double>(i);
    for(int j = 0; j < M.cols; j++)
        sum += std::max(Mi[j], 0.);
}
```

Some operations, like the one above, do not actually depend on the array shape. They just process elements of an array one by one (or elements from multiple arrays that have the same coordinates, for example, array addition). Such operations are called *element-wise*. It makes sense to check whether all the input/output arrays are continuous, namely, have no gaps at the end of each row. If yes, process them as a long single row:

```
// compute the sum of positive matrix elements, optimized variant
double sum=0;
int cols = M.cols, rows = M.rows;
if(M.isContinuous())
{
    cols *= rows;
    rows = 1;
```

```
}
for(int i = 0; i < rows; i++)
{
    const double* Mi = M.ptr<double>(i);
    for(int j = 0; j < cols; j++)
        sum += std::max(Mi[j], 0.);
}
```

In case of the continuous matrix, the outer loop body is executed just once. So, the overhead is smaller, which is especially noticeable in case of small matrices.

Finally, there are STL-style iterators that are smart enough to skip gaps between successive rows:

```
// compute sum of positive matrix elements, iterator-based variant
double sum=0;
MatConstIterator_<double> it = M.begin<double>(), it_end = M.end<double>();
for(; it != it_end; ++it)
    sum += std::max(*it, 0.);
```

The matrix iterators are random-access iterators, so they can be passed to any STL algorithm, including `std::sort()`.

## Matrix Expressions

This is a list of implemented matrix operations that can be combined in arbitrary complex expressions (here A, B stand for matrices ( `Mat` ), s for a scalar ( `Scalar` ), alpha for a real-valued scalar ( `double` )):

- Addition, subtraction, negation: `A+B, A-B, A+s, A-s, s+A, s-A, -A`

- Scaling: `A*alpha`

- Per-element multiplication and division: `A.mul(B), A/B, alpha/A`

- Matrix multiplication: `A*B`

- Transposition: `A.t()` (means $A^T$)

- Matrix inversion and pseudo-inversion, solving linear systems and least-squares problems:

  `A.inv([method])` (~ $A^{-1}$) , `A.inv([method])*B` (~ X: AX=B)

- Comparison: `A cmpop B, A cmpop alpha, alpha cmpop A`, where `cmpop` is one of : `>, >=, ==, !=, <=, <`. The result of comparison is an 8-bit single channel mask whose elements are set to 255 (if the particular element or pair of elements satisfy the condition) or 0.

- Bitwise logical operations: `A logicop B, A logicop s, s logicop A, ~A`, where `logicop` is one of : `&, |, ^`.

- Element-wise minimum and maximum: `min(A, B), min(A, alpha), max(A, B), max(A, alpha)`

- Element-wise absolute value: `abs(A)`

- Cross-product, dot-product: `A.cross(B) A.dot(B)`

- Any function of matrix or matrices and scalars that returns a matrix or a scalar, such as `norm, mean, sum, countNonZero, trace, determinant, repeat`, and others.

- Matrix initializers ( `Mat::eye(), Mat::zeros(), Mat::ones()` ), matrix comma-separated initializers, matrix constructors and operators that extract sub-matrices (see `Mat` description).

- `Mat_<destination_type>()` constructors to cast the result to the proper type.

---

**Note:** Comma-separated initializers and probably some other operations may require additional explicit `Mat()` or `Mat_<T>()` constructor calls to resolve a possible ambiguity.

---

Here are examples of matrix expressions:

```
// compute pseudo-inverse of A, equivalent to A.inv(DECOMP_SVD)
SVD svd(A);
Mat pinvA = svd.vt.t()*Mat::diag(1./svd.w)*svd.u.t();

// compute the new vector of parameters in the Levenberg-Marquardt algorithm
x -= (A.t()*A + lambda*Mat::eye(A.cols,A.cols,A.type())).inv(DECOMP_CHOLESKY)*(A.t()*err);

// sharpen image using "unsharp mask" algorithm
Mat blurred; double sigma = 1, threshold = 5, amount = 1;
GaussianBlur(img, blurred, Size(), sigma, sigma);
Mat lowContrastMask = abs(img - blurred) < threshold;
Mat sharpened = img*(1+amount) + blurred*(-amount);
img.copyTo(sharpened, lowContrastMask);
```

Below is the formal description of the `Mat` methods.

## Mat::Mat

Various Mat constructors

**C++:** `Mat::`**`Mat`**`()`

**C++:** `Mat::`**`Mat`**`(int **rows**, int **cols**, int **type**)`

**C++:** `Mat::`**`Mat`**`(Size **size**, int **type**)`

**C++:** `Mat::`**`Mat`**`(int **rows**, int **cols**, int **type**, const Scalar& **s**)`

**C++:** `Mat::`**`Mat`**`(Size **size**, int **type**, const Scalar& **s**)`

**C++:** `Mat::`**`Mat`**`(const Mat& **m**)`

**C++:** `Mat::`**`Mat`**`(int **rows**, int **cols**, int **type**, void* **data**, size_t **step**=AUTO_STEP)`

**C++:** `Mat::`**`Mat`**`(Size **size**, int **type**, void* **data**, size_t **step**=AUTO_STEP)`

**C++:** `Mat::`**`Mat`**`(const Mat& **m**, const Range& **rowRange**, const Range& **colRange**=Range::all() )`

**C++:** `Mat::`**`Mat`**`(const Mat& **m**, const Rect& **roi**)`

**C++:** `Mat::`**`Mat`**`(const CvMat* **m**, bool **copyData**=false)`

**C++:** `Mat::`**`Mat`**`(const IplImage* **img**, bool **copyData**=false)`

**C++:** template<typename T, int n> explicit `Mat::`**`Mat`**`(const Vec<T, n>& **vec**, bool **copyData**=true)`

**C++:** template<typename T, int m, int n> explicit `Mat::`**`Mat`**`(const Matx<T, m, n>& **vec**, bool **copy-Data**=true)`

**C++:** template<typename T> explicit `Mat::`**`Mat`**`(const vector<T>& **vec**, bool **copyData**=false)`

**C++:** `Mat::`**`Mat`**`(int **ndims**, const int* **sizes**, int **type**)`

**C++:** `Mat::`**`Mat`**`(int **ndims**, const int* **sizes**, int **type**, const Scalar& **s**)`

**C++:** `Mat::`**`Mat`**`(int **ndims**, const int* **sizes**, int **type**, void* **data**, const size_t* **steps**=0)`

**C++:** `Mat::`**`Mat`**`(const Mat& **m**, const Range* **ranges**)`

---

**Parameters**

**ndims** – Array dimensionality.

**rows** – Number of rows in a 2D array.

**cols** – Number of columns in a 2D array.

**roi** – Region of interest.

**size** – 2D array size: `Size(cols, rows)` . In the `Size()` constructor, the number of rows and the number of columns go in the reverse order.

**sizes** – Array of integers specifying an n-dimensional array shape.

**type** – Array type. Use `CV_8UC1`, `...`, `CV_64FC4` to create 1-4 channel matrices, or `CV_8UC(n)`, `...`, `CV_64FC(n)` to create multi-channel (up to `CV_MAX_CN` channels) matrices.

**s** – An optional value to initialize each matrix element with. To set all the matrix elements to the particular value after the construction, use the assignment operator `Mat::operator=(const Scalar& value)` .

**data** – Pointer to the user data. Matrix constructors that take `data` and `step` parameters do not allocate matrix data. Instead, they just initialize the matrix header that points to the specified data, which means that no data is copied. This operation is very efficient and can be used to process external data using OpenCV functions. The external data is not automatically deallocated, so you should take care of it.

**step** – Number of bytes each matrix row occupies. The value should include the padding bytes at the end of each row, if any. If the parameter is missing (set to `AUTO_STEP` ), no padding is assumed and the actual step is calculated as `cols*elemSize()` . See `Mat::elemSize()` .

**steps** – Array of `ndims-1` steps in case of a multi-dimensional array (the last step is always set to the element size). If not specified, the matrix is assumed to be continuous.

**m** – Array that (as a whole or partly) is assigned to the constructed matrix. No data is copied by these constructors. Instead, the header pointing to `m` data or its sub-array is constructed and associated with it. The reference counter, if any, is incremented. So, when you modify the matrix formed using such a constructor, you also modify the corresponding elements of `m` . If you want to have an independent copy of the sub-array, use `Mat::clone()` .

**img** – Pointer to the old-style `IplImage` image structure. By default, the data is shared between the original image and the new matrix. But when `copyData` is set, the full copy of the image data is created.

**vec** – STL vector whose elements form the matrix. The matrix has a single column and the number of rows equal to the number of vector elements. Type of the matrix matches the type of vector elements. The constructor can handle arbitrary types, for which there is a properly declared `DataType` . This means that the vector elements must be primitive numbers or uni-type numerical tuples of numbers. Mixed-type structures are not supported. The corresponding constructor is explicit. Since STL vectors are not automatically converted to `Mat` instances, you should write `Mat(vec)` explicitly. Unless you copy the data into the matrix ( `copyData=true` ), no new elements will be added to the vector because it can potentially yield vector data reallocation, and, thus, the matrix data pointer will be invalid.

**copyData** – Flag to specify whether the underlying data of the STL vector or the old-style `CvMat` or `IplImage` should be copied to (`true`) or shared with (`false`) the newly constructed matrix. When the data is copied, the allocated buffer is managed using `Mat` refer-

ence counting mechanism. While the data is shared, the reference counter is NULL, and you should not deallocate the data until the matrix is not destructed.

**rowRange** – Range of the `m` rows to take. As usual, the range start is inclusive and the range end is exclusive. Use `Range::all()` to take all the rows.

**colRange** – Range of the `m` columns to take. Use `Range::all()` to take all the columns.

**ranges** – Array of selected ranges of `m` along each dimensionality.

These are various constructors that form a matrix. As noted in the *Automatic Allocation of the Output Data*, often the default constructor is enough, and the proper matrix will be allocated by an OpenCV function. The constructed matrix can further be assigned to another matrix or matrix expression or can be allocated with `Mat::create()`. In the former case, the old content is de-referenced.

## Mat::~Mat

The Mat destructor.

**C++:** `Mat::~Mat()`

The matrix destructor calls `Mat::release()`.

## Mat::operator =

Provides matrix assignment operators.

**C++:** Mat& Mat::**operator=**(const Mat& **m**)

**C++:** Mat& Mat::**operator=**(const MatExpr& **expr**)

**C++:** Mat& Mat::**operator=**(const Scalar& **s**)

> **Parameters**

>> **m** – Assigned, right-hand-side matrix. Matrix assignment is an O(1) operation. This means that no data is copied but the data is shared and the reference counter, if any, is incremented. Before assigning new data, the old data is de-referenced via `Mat::release()`.

>> **expr** – Assigned matrix expression object. As opposite to the first form of the assignment operation, the second form can reuse already allocated matrix if it has the right size and type to fit the matrix expression result. It is automatically handled by the real function that the matrix expressions is expanded to. For example, `C=A+B` is expanded to `add(A, B, C)`, and `add()` takes care of automatic `C` reallocation.

>> **s** – Scalar assigned to each matrix element. The matrix size or type is not changed.

These are available assignment operators. Since they all are very different, make sure to read the operator parameters description.

## Mat::row

Creates a matrix header for the specified matrix row.

**C++:** Mat Mat::**row**(int **y**) const

> **Parameters**

>> **y** – A 0-based row index.

The method makes a new header for the specified matrix row and returns it. This is an O(1) operation, regardless of the matrix size. The underlying data of the new matrix is shared with the original matrix. Here is the example of one of the classical basic matrix processing operations, `axpy`, used by LU and many other algorithms:

```
inline void matrix_axpy(Mat& A, int i, int j, double alpha)
{
    A.row(i) += A.row(j)*alpha;
}
```

**Note:** In the current implementation, the following code does not work as expected:

```
Mat A;
...
A.row(i) = A.row(j); // will not work
```

This happens because `A.row(i)` forms a temporary header that is further assigned to another header. Remember that each of these operations is O(1), that is, no data is copied. Thus, the above assignment is not true if you may have expected the j-th row to be copied to the i-th row. To achieve that, you should either turn this simple assignment into an expression or use the `Mat::copyTo()` method:

```
Mat A;
...
// works, but looks a bit obscure.
A.row(i) = A.row(j) + 0;

// this is a bit longer, but the recommended method.
A.row(j).copyTo(A.row(i));
```

## Mat::col

Creates a matrix header for the specified matrix column.

**C++:** Mat Mat::**col**(int **x**) const

> **Parameters**
>
> > **x** – A 0-based column index.

The method makes a new header for the specified matrix column and returns it. This is an O(1) operation, regardless of the matrix size. The underlying data of the new matrix is shared with the original matrix. See also the `Mat::row()` description.

## Mat::rowRange

Creates a matrix header for the specified row span.

**C++:** Mat Mat::**rowRange**(int **startrow**, int **endrow**) const

**C++:** Mat Mat::**rowRange**(const Range& **r**) const

> **Parameters**
>
> > **startrow** – An inclusive 0-based start index of the row span.
> >
> > **endrow** – An exclusive 0-based ending index of the row span.
> >
> > **r** – Range structure containing both the start and the end indices.

The method makes a new header for the specified row span of the matrix. Similarly to `Mat::row()` and `Mat::col()`, this is an O(1) operation.

## Mat::colRange

Creates a matrix header for the specified row span.

**C++:** Mat Mat::**colRange**(int **startcol**, int **endcol**) const

**C++:** Mat Mat::**colRange**(const Range& **r**) const

> **Parameters**
>
> > **startcol** – An inclusive 0-based start index of the column span.
> >
> > **endcol** – An exclusive 0-based ending index of the column span.
> >
> > **r** – Range structure containing both the start and the end indices.

The method makes a new header for the specified column span of the matrix. Similarly to `Mat::row()` and `Mat::col()`, this is an O(1) operation.

## Mat::diag

Extracts a diagonal from a matrix, or creates a diagonal matrix.

**C++:** Mat Mat::**diag**(int **d**=0 ) const

**C++:** **static** Mat Mat::**diag**(const Mat& **d**)

> **Parameters**
>
> > **d** – Single-column matrix that forms a diagonal matrix or index of the diagonal, with the following values:
> >
> > – **d=0** is the main diagonal.
> >
> > – **d>0** is a diagonal from the lower half. For example, d=1 means the diagonal is set immediately below the main one.
> >
> > – **d<0** is a diagonal from the upper half. For example, d=1 means the diagonal is set immediately above the main one.

The method makes a new header for the specified matrix diagonal. The new matrix is represented as a single-column matrix. Similarly to `Mat::row()` and `Mat::col()`, this is an O(1) operation.

## Mat::clone

Creates a full copy of the array and the underlying data.

**C++:** Mat Mat::**clone**() const

The method creates a full copy of the array. The original `step[]` is not taken into account. So, the array copy is a continuous array occupying `total()*elemSize()` bytes.

## Mat::copyTo

Copies the matrix to another one.

**C++:** void Mat::**copyTo**(OutputArray **m**) const

**C++:** void Mat::**copyTo**(OutputArray **m**, InputArray **mask**) const

>   **Parameters**

>>      **m** – Destination matrix. If it does not have a proper size or type before the operation, it is reallocated.

>>      **mask** – Operation mask. Its non-zero elements indicate which matrix elements need to be copied.

The method copies the matrix data to another matrix. Before copying the data, the method invokes

```
m.create(this->size(), this->type);
```

so that the destination matrix is reallocated if needed. While `m.copyTo(m);` works flawlessly, the function does not handle the case of a partial overlap between the source and the destination matrices.

When the operation mask is specified, and the `Mat::create` call shown above reallocated the matrix, the newly allocated matrix is initialized with all zeros before copying the data.

## Mat::convertTo

Converts an array to another data type with optional scaling.

**C++:** void Mat::**convertTo**(OutputArray **m**, int **rtype**, double **alpha**=1, double **beta**=0 ) const

>   **Parameters**

>>      **m** – output matrix; if it does not have a proper size or type before the operation, it is reallocated.

>>      **rtype** – desired output matrix type or, rather, the depth since the number of channels are the same as the input has; if `rtype` is negative, the output matrix will have the same type as the input.

>>      **alpha** – optional scale factor.

>>      **beta** – optional delta added to the scaled values.

The method converts source pixel values to the target data type. `saturate_cast<>` is applied at the end to avoid possible overflows:

$$m(x, y) = saturate\_cast < rType > (\alpha(*this)(x, y) + \beta)$$

## Mat::assignTo

Provides a functional form of `convertTo`.

**C++:** void Mat::**assignTo**(Mat& **m**, int **type**=-1 ) const

>   **Parameters**

>>      **m** – Destination array.

>>      **type** – Desired destination array depth (or -1 if it should be the same as the source type).

This is an internally used method called by the *Matrix Expressions* engine.

## Mat::setTo

Sets all or some of the array elements to the specified value.

**C++:** Mat& Mat::**setTo**(InputArray **value**, InputArray **mask**=noArray() )

> **Parameters**
>
>> **value** – Assigned scalar converted to the actual array type.
>>
>> **mask** – Operation mask of the same size as *this. This is an advanced variant of the Mat::operator=(const Scalar& s) operator.

## Mat::reshape

Changes the shape and/or the number of channels of a 2D matrix without copying the data.

**C++:** Mat Mat::**reshape**(int **cn**, int **rows**=0) const

> **Parameters**
>
>> **cn** – New number of channels. If the parameter is 0, the number of channels remains the same.
>>
>> **rows** – New number of rows. If the parameter is 0, the number of rows remains the same.

The method makes a new matrix header for *this elements. The new matrix may have a different size and/or different number of channels. Any combination is possible if:

- No extra elements are included into the new matrix and no elements are excluded. Consequently, the product rows*cols*channels() must stay the same after the transformation.

- No data is copied. That is, this is an O(1) operation. Consequently, if you change the number of rows, or the operation changes the indices of elements row in some other way, the matrix must be continuous. See Mat::isContinuous() .

For example, if there is a set of 3D points stored as an STL vector, and you want to represent the points as a 3xN matrix, do the following:

```
std::vector<Point3f> vec;
...

Mat pointMat = Mat(vec). // convert vector to Mat, O(1) operation
                 reshape(1). // make Nx3 1-channel matrix out of Nx1 3-channel.
                             // Also, an O(1) operation
                    t(); // finally, transpose the Nx3 matrix.
                         // This involves copying all the elements
```

## Mat::t

Transposes a matrix.

**C++:** MatExpr Mat::**t**() const

The method performs matrix transposition by means of matrix expressions. It does not perform the actual transposition but returns a temporary matrix transposition object that can be further used as a part of more complex matrix expressions or can be assigned to a matrix:

```
Mat A1 = A + Mat::eye(A.size(), A.type)*lambda;
Mat C = A1.t()*A1; // compute (A + lambda*I)^t * (A + lamda*I)
```

## Mat::inv

Inverses a matrix.

**C++:** MatExpr Mat::**inv**(int **method**=DECOMP_LU) const

>  **Parameters**

>>  **method** – Matrix inversion method. Possible values are the following:

>>  – **DECOMP_LU** is the LU decomposition. The matrix must be non-singular.

>>  – **DECOMP_CHOLESKY** is the Cholesky $LL^T$ decomposition for symmetrical positively defined matrices only. This type is about twice faster than LU on big matrices.

>>  – **DECOMP_SVD** is the SVD decomposition. If the matrix is singular or even non-square, the pseudo inversion is computed.

The method performs a matrix inversion by means of matrix expressions. This means that a temporary matrix inversion object is returned by the method and can be used further as a part of more complex matrix expressions or can be assigned to a matrix.

## Mat::mul

Performs an element-wise multiplication or division of the two matrices.

**C++:** MatExpr Mat::**mul**(InputArray **m**, double **scale**=1) const

>  **Parameters**

>>  **m** – Another array of the same type and the same size as *this, or a matrix expression.

>>  **scale** – Optional scale factor.

The method returns a temporary object encoding per-element array multiplication, with optional scale. Note that this is not a matrix multiplication that corresponds to a simpler "*" operator.

Example:

```
Mat C = A.mul(5/B); // equivalent to divide(A, B, C, 5)
```

## Mat::cross

Computes a cross-product of two 3-element vectors.

**C++:** Mat Mat::**cross**(InputArray **m**) const

>  **Parameters**

>>  **m** – Another cross-product operand.

The method computes a cross-product of two 3-element vectors. The vectors must be 3-element floating-point vectors of the same shape and size. The result is another 3-element vector of the same shape and type as operands.

## Mat::dot

Computes a dot-product of two vectors.

**C++:** double Mat::**dot**(InputArray **m**) const

>  **Parameters**

**m** – another dot-product operand.

The method computes a dot-product of two matrices. If the matrices are not single-column or single-row vectors, the top-to-bottom left-to-right scan ordering is used to treat them as 1D vectors. The vectors must have the same size and type. If the matrices have more than one channel, the dot products from all the channels are summed together.

## Mat::zeros

Returns a zero array of the specified size and type.

**C++:** **static** MatExpr Mat::**zeros**(int **rows**, int **cols**, int **type**)

**C++:** **static** MatExpr Mat::**zeros**(Size **size**, int **type**)

**C++:** **static** MatExpr Mat::**zeros**(int **ndims**, const int* **sz**, int **type**)

> **Parameters**
>
>> **ndims** – Array dimensionality.
>>
>> **rows** – Number of rows.
>>
>> **cols** – Number of columns.
>>
>> **size** – Alternative to the matrix size specification `Size(cols, rows)` .
>>
>> **sz** – Array of integers specifying the array shape.
>>
>> **type** – Created matrix type.

The method returns a Matlab-style zero array initializer. It can be used to quickly form a constant array as a function parameter, part of a matrix expression, or as a matrix initializer.

```
Mat A;
A = Mat::zeros(3, 3, CV_32F);
```

In the example above, a new matrix is allocated only if A is not a 3x3 floating-point matrix. Otherwise, the existing matrix A is filled with zeros.

## Mat::ones

Returns an array of all 1's of the specified size and type.

**C++:** **static** MatExpr Mat::**ones**(int **rows**, int **cols**, int **type**)

**C++:** **static** MatExpr Mat::**ones**(Size **size**, int **type**)

**C++:** **static** MatExpr Mat::**ones**(int **ndims**, const int* **sz**, int **type**)

> **Parameters**
>
>> **ndims** – Array dimensionality.
>>
>> **rows** – Number of rows.
>>
>> **cols** – Number of columns.
>>
>> **size** – Alternative to the matrix size specification `Size(cols, rows)` .
>>
>> **sz** – Array of integers specifying the array shape.
>>
>> **type** – Created matrix type.

The method returns a Matlab-style 1's array initializer, similarly to `Mat::zeros()`. Note that using this method you can initialize an array with an arbitrary value, using the following Matlab idiom:

```
Mat A = Mat::ones(100, 100, CV_8U)*3; // make 100x100 matrix filled with 3.
```

The above operation does not form a 100x100 matrix of 1's and then multiply it by 3. Instead, it just remembers the scale factor (3 in this case) and use it when actually invoking the matrix initializer.

## Mat::eye

Returns an identity matrix of the specified size and type.

**C++:** **static** MatExpr `Mat::eye`(int **rows**, int **cols**, int **type**)

**C++:** **static** MatExpr `Mat::eye`(Size **size**, int **type**)

> **Parameters**
>
> > **rows** – Number of rows.
> >
> > **cols** – Number of columns.
> >
> > **size** – Alternative matrix size specification as `Size(cols, rows)`.
> >
> > **type** – Created matrix type.

The method returns a Matlab-style identity matrix initializer, similarly to `Mat::zeros()`. Similarly to `Mat::ones()`, you can use a scale operation to create a scaled identity matrix efficiently:

```
// make a 4x4 diagonal matrix with 0.1's on the diagonal.
Mat A = Mat::eye(4, 4, CV_32F)*0.1;
```

## Mat::create

Allocates new array data if needed.

**C++:** void `Mat::create`(int **rows**, int **cols**, int **type**)

**C++:** void `Mat::create`(Size **size**, int **type**)

**C++:** void `Mat::create`(int **ndims**, const int* **sizes**, int **type**)

> **Parameters**
>
> > **ndims** – New array dimensionality.
> >
> > **rows** – New number of rows.
> >
> > **cols** – New number of columns.
> >
> > **size** – Alternative new matrix size specification: `Size(cols, rows)`
> >
> > **sizes** – Array of integers specifying a new array shape.
> >
> > **type** – New matrix type.

This is one of the key `Mat` methods. Most new-style OpenCV functions and methods that produce arrays call this method for each output array. The method uses the following algorithm:

1. If the current array shape and the type match the new ones, return immediately. Otherwise, de-reference the previous data by calling `Mat::release()`.

2. Initialize the new header.

3. Allocate the new data of `total()*elemSize()` bytes.

4. Allocate the new, associated with the data, reference counter and set it to 1.

Such a scheme makes the memory management robust and efficient at the same time and helps avoid extra typing for you. This means that usually there is no need to explicitly allocate output arrays. That is, instead of writing:

```
Mat color;
...
Mat gray(color.rows, color.cols, color.depth());
cvtColor(color, gray, CV_BGR2GRAY);
```

you can simply write:

```
Mat color;
...
Mat gray;
cvtColor(color, gray, CV_BGR2GRAY);
```

because `cvtColor`, as well as the most of OpenCV functions, calls `Mat::create()` for the output array internally.

## Mat::addref

Increments the reference counter.

**C++:** void Mat::**addref**()

The method increments the reference counter associated with the matrix data. If the matrix header points to an external data set (see `Mat::Mat()`), the reference counter is NULL, and the method has no effect in this case. Normally, to avoid memory leaks, the method should not be called explicitly. It is called implicitly by the matrix assignment operator. The reference counter increment is an atomic operation on the platforms that support it. Thus, it is safe to operate on the same matrices asynchronously in different threads.

## Mat::release

Decrements the reference counter and deallocates the matrix if needed.

**C++:** void Mat::**release**()

The method decrements the reference counter associated with the matrix data. When the reference counter reaches 0, the matrix data is deallocated and the data and the reference counter pointers are set to NULL's. If the matrix header points to an external data set (see `Mat::Mat()`), the reference counter is NULL, and the method has no effect in this case.

This method can be called manually to force the matrix data deallocation. But since this method is automatically called in the destructor, or by any other method that changes the data pointer, it is usually not needed. The reference counter decrement and check for 0 is an atomic operation on the platforms that support it. Thus, it is safe to operate on the same matrices asynchronously in different threads.

## Mat::resize

Changes the number of matrix rows.

**C++:** void Mat::**resize**(size_t **sz**)

**C++:** void Mat::**resize**(size_t **sz**, const Scalar& **s**)

   **Parameters**

> **sz** – New number of rows.
>
> **s** – Value assigned to the newly added elements.

The methods change the number of matrix rows. If the matrix is reallocated, the first `min(Mat::rows, sz)` rows are preserved. The methods emulate the corresponding methods of the STL vector class.

## Mat::reserve

Reserves space for the certain number of rows.

**C++:** void `Mat::`**`reserve`**(size_t **sz**)

> **Parameters**
>
> > **sz** – Number of rows.

The method reserves space for `sz` rows. If the matrix already has enough space to store `sz` rows, nothing happens. If the matrix is reallocated, the first `Mat::rows` rows are preserved. The method emulates the corresponding method of the STL vector class.

## Mat::push_back

Adds elements to the bottom of the matrix.

**C++:** template<typename T> void `Mat::`**`push_back`**(const T& **elem**)

**C++:** void `Mat::`**`push_back`**(const Mat& **m**)

> **Parameters**
>
> > **elem** – Added element(s).

The methods add one or more elements to the bottom of the matrix. They emulate the corresponding method of the STL vector class. When `elem` is `Mat` , its type and the number of columns must be the same as in the container matrix.

## Mat::pop_back

Removes elements from the bottom of the matrix.

**C++:** template<typename T> void `Mat::`**`pop_back`**(size_t **nelems**=1)

> **Parameters**
>
> > **nelems** – Number of removed rows. If it is greater than the total number of rows, an exception is thrown.

The method removes one or more rows from the bottom of the matrix.

## Mat::locateROI

Locates the matrix header within a parent matrix.

**C++:** void `Mat::`**`locateROI`**(Size& **wholeSize**, Point& **ofs**) const

> **Parameters**
>
> > **wholeSize** – Output parameter that contains the size of the whole matrix containing `*this` as a part.

> > > **ofs** – Output parameter that contains an offset of `*this` inside the whole matrix.

After you extracted a submatrix from a matrix using `Mat::row()`, `Mat::col()`, `Mat::rowRange()`, `Mat::colRange()`, and others, the resultant submatrix points just to the part of the original big matrix. However, each submatrix contains information (represented by `datastart` and `dataend` fields) that helps reconstruct the original matrix size and the position of the extracted submatrix within the original matrix. The method `locateROI` does exactly that.

## Mat::adjustROI

Adjusts a submatrix size and position within the parent matrix.

**C++:** Mat& Mat::**adjustROI**(int **dtop**, int **dbottom**, int **dleft**, int **dright**)

> **Parameters**

> > **dtop** – Shift of the top submatrix boundary upwards.

> > **dbottom** – Shift of the bottom submatrix boundary downwards.

> > **dleft** – Shift of the left submatrix boundary to the left.

> > **dright** – Shift of the right submatrix boundary to the right.

The method is complimentary to `Mat::locateROI()`. The typical use of these functions is to determine the submatrix position within the parent matrix and then shift the position somehow. Typically, it can be required for filtering operations when pixels outside of the ROI should be taken into account. When all the method parameters are positive, the ROI needs to grow in all directions by the specified amount, for example:

```
A.adjustROI(2, 2, 2, 2);
```

In this example, the matrix size is increased by 4 elements in each direction. The matrix is shifted by 2 elements to the left and 2 elements up, which brings in all the necessary pixels for the filtering with the 5x5 kernel.

`adjustROI` forces the adjusted ROI to be inside of the parent matrix that is boundaries of the adjusted ROI are constrained by boundaries of the parent matrix. For example, if the submatrix A is located in the first row of a parent matrix and you called `A.adjustROI(2, 2, 2, 2)` then A will not be increased in the upward direction.

The function is used internally by the OpenCV filtering functions, like `filter2D()`, morphological operations, and so on.

**See Also:**

`copyMakeBorder()`

## Mat::operator()

Extracts a rectangular submatrix.

**C++:** Mat Mat::**operator()**(Range **rowRange**, Range **colRange**) const

**C++:** Mat Mat::**operator()**(const Rect& **roi**) const

**C++:** Mat Mat::**operator()**(const Range* **ranges**) const

> **Parameters**

> > **rowRange** – Start and end row of the extracted submatrix. The upper boundary is not included. To select all the rows, use `Range::all()`.

> > **colRange** – Start and end column of the extracted submatrix. The upper boundary is not included. To select all the columns, use `Range::all()`.

> > **roi** – Extracted submatrix specified as a rectangle.

> > **ranges** – Array of selected ranges along each array dimension.

The operators make a new header for the specified sub-array of `*this`. They are the most generalized forms of `Mat::row()`, `Mat::col()`, `Mat::rowRange()`, and `Mat::colRange()`. For example, A(Range(0, 10), Range::all()) is equivalent to `A.rowRange(0, 10)`. Similarly to all of the above, the operators are O(1) operations, that is, no matrix data is copied.

## Mat::operator CvMat

Creates the `CvMat` header for the matrix.

**C++:** Mat::**operator** CvMat() const

The operator creates the `CvMat` header for the matrix without copying the underlying data. The reference counter is not taken into account by this operation. Thus, you should make sure than the original matrix is not deallocated while the `CvMat` header is used. The operator is useful for intermixing the new and the old OpenCV API's, for example:

```
Mat img(Size(320, 240), CV_8UC3);
...

CvMat cvimg = img;
mycvOldFunc( &cvimg, ...);
```

where `mycvOldFunc` is a function written to work with OpenCV 1.x data structures.

## Mat::operator IplImage

Creates the `IplImage` header for the matrix.

**C++:** Mat::**operator** IplImage() const

The operator creates the `IplImage` header for the matrix without copying the underlying data. You should make sure than the original matrix is not deallocated while the `IplImage` header is used. Similarly to `Mat::operator CvMat`, the operator is useful for intermixing the new and the old OpenCV API's.

## Mat::total

Returns the total number of array elements.

**C++:** size_t Mat::**total**() const

The method returns the number of array elements (a number of pixels if the array represents an image).

## Mat::isContinuous

Reports whether the matrix is continuous or not.

**C++:** bool Mat::**isContinuous**() const

The method returns `true` if the matrix elements are stored continuously without gaps at the end of each row. Otherwise, it returns `false`. Obviously, `1x1` or `1xN` matrices are always continuous. Matrices created with `Mat::create()` are always continuous. But if you extract a part of the matrix using `Mat::col()`, `Mat::diag()`, and so on, or constructed a matrix header for externally allocated data, such matrices may no longer have this property.

---

The continuity flag is stored as a bit in the `Mat::flags` field and is computed automatically when you construct a matrix header. Thus, the continuity check is a very fast operation, though theoretically it could be done as follows:

```cpp
// alternative implementation of Mat::isContinuous()
bool myCheckMatContinuity(const Mat& m)
{
    //return (m.flags & Mat::CONTINUOUS_FLAG) != 0;
    return m.rows == 1 || m.step == m.cols*m.elemSize();
}
```

The method is used in quite a few of OpenCV functions. The point is that element-wise operations (such as arithmetic and logical operations, math functions, alpha blending, color space transformations, and others) do not depend on the image geometry. Thus, if all the input and output arrays are continuous, the functions can process them as very long single-row vectors. The example below illustrates how an alpha-blending function can be implemented.

```cpp
template<typename T>
void alphaBlendRGBA(const Mat& src1, const Mat& src2, Mat& dst)
{
    const float alpha_scale = (float)std::numeric_limits<T>::max(),
                inv_scale = 1.f/alpha_scale;

    CV_Assert( src1.type() == src2.type() &&
               src1.type() == CV_MAKETYPE(DataType<T>::depth, 4) &&
               src1.size() == src2.size());
    Size size = src1.size();
    dst.create(size, src1.type());

    // here is the idiom: check the arrays for continuity and,
    // if this is the case,
    // treat the arrays as 1D vectors
    if( src1.isContinuous() && src2.isContinuous() && dst.isContinuous() )
    {
        size.width *= size.height;
        size.height = 1;
    }
    size.width *= 4;

    for( int i = 0; i < size.height; i++ )
    {
        // when the arrays are continuous,
        // the outer loop is executed only once
        const T* ptr1 = src1.ptr<T>(i);
        const T* ptr2 = src2.ptr<T>(i);
        T* dptr = dst.ptr<T>(i);

        for( int j = 0; j < size.width; j += 4 )
        {
            float alpha = ptr1[j+3]*inv_scale, beta = ptr2[j+3]*inv_scale;
            dptr[j] = saturate_cast<T>(ptr1[j]*alpha + ptr2[j]*beta);
            dptr[j+1] = saturate_cast<T>(ptr1[j+1]*alpha + ptr2[j+1]*beta);
            dptr[j+2] = saturate_cast<T>(ptr1[j+2]*alpha + ptr2[j+2]*beta);
            dptr[j+3] = saturate_cast<T>((1 - (1-alpha)*(1-beta))*alpha_scale);
        }
    }
}
```

This approach, while being very simple, can boost the performance of a simple element-operation by 10-20 percents, especially if the image is rather small and the operation is quite simple.

Another OpenCV idiom in this function, a call of `Mat::create()` for the destination array, that allocates the destination array unless it already has the proper size and type. And while the newly allocated arrays are always continuous, you still need to check the destination array because `Mat::create()` does not always allocate a new matrix.

## Mat::elemSize

Returns the matrix element size in bytes.

**C++:** size_t Mat::**elemSize**() const

The method returns the matrix element size in bytes. For example, if the matrix type is `CV_16SC3`, the method returns `3*sizeof(short)` or 6.

## Mat::elemSize1

Returns the size of each matrix element channel in bytes.

**C++:** size_t Mat::**elemSize1**() const

The method returns the matrix element channel size in bytes, that is, it ignores the number of channels. For example, if the matrix type is `CV_16SC3`, the method returns `sizeof(short)` or 2.

## Mat::type

Returns the type of a matrix element.

**C++:** int Mat::**type**() const

The method returns a matrix element type. This is an identifier compatible with the `CvMat` type system, like `CV_16SC3` or 16-bit signed 3-channel array, and so on.

## Mat::depth

Returns the depth of a matrix element.

**C++:** int Mat::**depth**() const

The method returns the identifier of the matrix element depth (the type of each individual channel). For example, for a 16-bit signed 3-channel array, the method returns `CV_16S`. A complete list of matrix types contains the following values:

- `CV_8U` - 8-bit unsigned integers ( `0..255` )
- `CV_8S` - 8-bit signed integers ( `-128..127` )
- `CV_16U` - 16-bit unsigned integers ( `0..65535` )
- `CV_16S` - 16-bit signed integers ( `-32768..32767` )
- `CV_32S` - 32-bit signed integers ( `-2147483648..2147483647` )
- `CV_32F` - 32-bit floating-point numbers ( `-FLT_MAX..FLT_MAX, INF, NAN` )
- `CV_64F` - 64-bit floating-point numbers ( `-DBL_MAX..DBL_MAX, INF, NAN` )

## Mat::channels

Returns the number of matrix channels.

**C++:** int Mat::**channels**() const

The method returns the number of matrix channels.

## Mat::step1

Returns a normalized step.

**C++:** size_t Mat::**step1**(int **i**=0 ) const

The method returns a matrix step divided by `Mat::elemSize1()` . It can be useful to quickly access an arbitrary matrix element.

## Mat::size

Returns a matrix size.

**C++:** Size Mat::**size**() const

The method returns a matrix size: `Size(cols, rows)` . When the matrix is more than 2-dimensional, the returned size is (-1, -1).

## Mat::empty

Returns `true` if the array has no elements.

**C++:** bool Mat::**empty**() const

The method returns `true` if `Mat::total()` is 0 or if `Mat::data` is NULL. Because of `pop_back()` and `resize()` methods `M.total() == 0` does not imply that `M.data == NULL` .

## Mat::ptr

Returns a pointer to the specified matrix row.

**C++:** uchar* Mat::**ptr**(int **i0**=0)

**C++:** const uchar* Mat::**ptr**(int **i0**=0) const

**C++:** template<typename _Tp> _Tp* Mat::**ptr**(int **i0**=0)

**C++:** template<typename _Tp> const _Tp* Mat::**ptr**(int **i0**=0) const

> **Parameters**

>> **i0** – A 0-based row index.

The methods return `uchar*` or typed pointer to the specified matrix row. See the sample in `Mat::isContinuous()` to know how to use these methods.

## Mat::at

Returns a reference to the specified array element.

**C++:** template<typename T> T& Mat::**at**(int **i**) const

**C++:** template<typename T> const T& Mat::**at**(int **i**) const

**C++:** template<typename T> T& Mat::**at**(int **i**, int **j**)

**C++:** template<typename T> const T& Mat::**at**(int **i**, int **j**) const

**C++:** template<typename T> T& Mat::**at**(Point **pt**)

**C++:** template<typename T> const T& Mat::**at**(Point **pt**) const

**C++:** template<typename T> T& Mat::**at**(int **i**, int **j**, int **k**)

**C++:** template<typename T> const T& Mat::**at**(int **i**, int **j**, int **k**) const

**C++:** template<typename T> T& Mat::**at**(const int* **idx**)

**C++:** template<typename T> const T& Mat::**at**(const int* **idx**) const

> **Parameters**
>> **i** – Index along the dimension 0
>>
>> **j** – Index along the dimension 1
>>
>> **k** – Index along the dimension 2
>>
>> **pt** – Element position specified as `Point(j,i)` .
>>
>> **idx** – Array of `Mat::dims` indices.

The template methods return a reference to the specified array element. For the sake of higher performance, the index range checks are only performed in the Debug configuration.

Note that the variants with a single index (i) can be used to access elements of single-row or single-column 2-dimensional arrays. That is, if, for example, A is a `1 x N` floating-point matrix and B is an `M x 1` integer matrix, you can simply write `A.at<float>(k+4)` and `B.at<int>(2*i+1)` instead of `A.at<float>(0,k+4)` and `B.at<int>(2*i+1,0)` , respectively.

The example below initializes a Hilbert matrix:

```
Mat H(100, 100, CV_64F);
for(int i = 0; i < H.rows; i++)
    for(int j = 0; j < H.cols; j++)
        H.at<double>(i,j)=1./(i+j+1);
```

## Mat::begin

Returns the matrix iterator and sets it to the first matrix element.

**C++:** template<typename _Tp> MatIterator_<_Tp> Mat::**begin**()

**C++:** template<typename _Tp> MatConstIterator_<_Tp> Mat::**begin**() const

The methods return the matrix read-only or read-write iterators. The use of matrix iterators is very similar to the use of bi-directional STL iterators. In the example below, the alpha blending function is rewritten using the matrix iterators:

```
template<typename T>
void alphaBlendRGBA(const Mat& src1, const Mat& src2, Mat& dst)
{
    typedef Vec<T, 4> VT;

    const float alpha_scale = (float)std::numeric_limits<T>::max(),
                inv_scale = 1.f/alpha_scale;

    CV_Assert( src1.type() == src2.type() &&
               src1.type() == DataType<VT>::type &&
               src1.size() == src2.size());
    Size size = src1.size();
    dst.create(size, src1.type());

    MatConstIterator_<VT> it1 = src1.begin<VT>(), it1_end = src1.end<VT>();
    MatConstIterator_<VT> it2 = src2.begin<VT>();
    MatIterator_<VT> dst_it = dst.begin<VT>();

    for( ; it1 != it1_end; ++it1, ++it2, ++dst_it )
    {
        VT pix1 = *it1, pix2 = *it2;
        float alpha = pix1[3]*inv_scale, beta = pix2[3]*inv_scale;
        *dst_it = VT(saturate_cast<T>(pix1[0]*alpha + pix2[0]*beta),
                     saturate_cast<T>(pix1[1]*alpha + pix2[1]*beta),
                     saturate_cast<T>(pix1[2]*alpha + pix2[2]*beta),
                     saturate_cast<T>((1 - (1-alpha)*(1-beta))*alpha_scale));
    }
}
```

## Mat::end

Returns the matrix iterator and sets it to the after-last matrix element.

**C++:** template<typename _Tp> MatIterator_<_Tp> Mat::**end**()

**C++:** template<typename _Tp> MatConstIterator_<_Tp> Mat::**end**() const

The methods return the matrix read-only or read-write iterators, set to the point following the last matrix element.

## Mat_

**class Mat_**

Template matrix class derived from Mat .

```
template<typename _Tp> class Mat_ : public Mat
{
public:
    // ... some specific methods
    //         and
    // no new extra fields
};
```

The class Mat_<_Tp> is a "thin" template wrapper on top of the Mat class. It does not have any extra data fields. Nor this class nor Mat has any virtual methods. Thus, references or pointers to these two classes can be freely but carefully converted one to another. For example:

```
// create a 100x100 8-bit matrix
Mat M(100,100,CV_8U);
// this will be compiled fine. no any data conversion will be done.
Mat_<float>& M1 = (Mat_<float>&)M;
// the program is likely to crash at the statement below
M1(99,99) = 1.f;
```

While `Mat` is sufficient in most cases, `Mat_` can be more convenient if you use a lot of element access operations and if you know matrix type at the compilation time. Note that `Mat::at<_Tp>(int y, int x)` and `Mat_<_Tp>::operator ()(int y, int x)` do absolutely the same and run at the same speed, but the latter is certainly shorter:

```
Mat_<double> M(20,20);
for(int i = 0; i < M.rows; i++)
    for(int j = 0; j < M.cols; j++)
        M(i,j) = 1./(i+j+1);
Mat E, V;
eigen(M,E,V);
cout << E.at<double>(0,0)/E.at<double>(M.rows-1,0);
```

To use `Mat_` for multi-channel images/matrices, pass `Vec` as a `Mat_` parameter:

```
// allocate a 320x240 color image and fill it with green (in RGB space)
Mat_<Vec3b> img(240, 320, Vec3b(0,255,0));
// now draw a diagonal white line
for(int i = 0; i < 100; i++)
    img(i,i)=Vec3b(255,255,255);
// and now scramble the 2nd (red) channel of each pixel
for(int i = 0; i < img.rows; i++)
    for(int j = 0; j < img.cols; j++)
        img(i,j)[2] ^= (uchar)(i ^ j);
```

## InputArray

**class InputArray**

This is the proxy class for passing read-only input arrays into OpenCV functions. It is defined as

```
typedef const _InputArray& InputArray;
```

where `_InputArray` is a class that can be constructed from `Mat`, `Mat_<T>`, `Matx<T, m, n>`, `std::vector<T>`, `std::vector<std::vector<T> >` or `std::vector<Mat>`. It can also be constructed from a matrix expression.

Since this is mostly implementation-level class, and its interface may change in future versions, we do not describe it in details. There are a few key things, though, that should be kept in mind:

- When you see in the reference manual or in OpenCV source code a function that takes `InputArray`, it means that you can actually pass `Mat`, `Matx`, `vector<T>` etc. (see above the complete list).

- Optional input arguments: If some of the input arrays may be empty, pass `cv::noArray()` (or simply `cv::Mat()` as you probably did before).

- The class is designed solely for passing parameters. That is, normally you *should not* declare class members, local and global variables of this type.

- If you want to design your own function or a class method that can operate of arrays of multiple types, you can use `InputArray` (or `OutputArray`) for the respective parameters. Inside a function you should use `_InputArray::getMat()` method to construct a matrix header for the array (without copying data). `_InputArray::kind()` can be used to distinguish `Mat` from `vector<>` etc., but normally it is not needed.

Here is how you can use a function that takes `InputArray`

```
std::vector<Point2f> vec;
// points or a circle
for( int i = 0; i < 30; i++ )
    vec.push_back(Point2f((float)(100 + 30*cos(i*CV_PI*2/5)),
                          (float)(100 - 30*sin(i*CV_PI*2/5))));
cv::transform(vec, vec, cv::Matx23f(0.707, -0.707, 10, 0.707, 0.707, 20));
```

That is, we form an STL vector containing points, and apply in-place affine transformation to the vector using the 2x3 matrix created inline as `Matx<float, 2, 3>` instance.

Here is how such a function can be implemented (for simplicity, we implement a very specific case of it, according to the assertion statement inside)

```
void myAffineTransform(InputArray _src, OutputArray _dst, InputArray _m)
{
    // get Mat headers for input arrays. This is O(1) operation,
    // unless _src and/or _m are matrix expressions.
    Mat src = _src.getMat(), m = _m.getMat();
    CV_Assert( src.type() == CV_32FC2 && m.type() == CV_32F && m.size() == Size(3, 2) );

    // [re]create the output array so that it has the proper size and type.
    // In case of Mat it calls Mat::create, in case of STL vector it calls vector::resize.
    _dst.create(src.size(), src.type());
    Mat dst = _dst.getMat();

    for( int i = 0; i < src.rows; i++ )
        for( int j = 0; j < src.cols; j++ )
        {
            Point2f pt = src.at<Point2f>(i, j);
            dst.at<Point2f>(i, j) = Point2f(m.at<float>(0, 0)*pt.x +
                                            m.at<float>(0, 1)*pt.y +
                                            m.at<float>(0, 2),
                                            m.at<float>(1, 0)*pt.x +
                                            m.at<float>(1, 1)*pt.y +
                                            m.at<float>(1, 2));
        }
}
```

There is another related type, `InputArrayOfArrays`, which is currently defined as a synonym for `InputArray`:

```
typedef InputArray InputArrayOfArrays;
```

It denotes function arguments that are either vectors of vectors or vectors of matrices. A separate synonym is needed to generate Python/Java etc. wrappers properly. At the function implementation level their use is similar, but `_InputArray::getMat(idx)` should be used to get header for the idx-th component of the outer vector and `_InputArray::size().area()` should be used to find the number of components (vectors/matrices) of the outer vector.

## OutputArray

**class `OutputArray` : public `InputArray`**

This type is very similar to `InputArray` except that it is used for input/output and output function parameters. Just like with `InputArray`, OpenCV users should not care about `OutputArray`, they just pass `Mat`, `vector<T>` etc. to the functions. The same limitation as for `InputArray`: **Do not explicitly create OutputArray instances** applies here too.

If you want to make your function polymorphic (i.e. accept different arrays as output parameters), it is also not very difficult. Take the sample above as the reference. Note that `_OutputArray::create()` needs to be called before `_OutputArray::getMat()`. This way you guarantee that the output array is properly allocated.

Optional output parameters. If you do not need certain output array to be computed and returned to you, pass `cv::noArray()`, just like you would in the case of optional input array. At the implementation level, use `_OutputArray::needed()` to check if certain output array needs to be computed or not.

There are several synonyms for `OutputArray` that are used to assist automatic Python/Java/... wrapper generators:

```cpp
typedef OutputArray OutputArrayOfArrays;
typedef OutputArray InputOutputArray;
typedef OutputArray InputOutputArrayOfArrays;
```

## NAryMatIterator

class **NAryMatIterator**

n-ary multi-dimensional array iterator.

```cpp
class CV_EXPORTS NAryMatIterator
{
public:
    //! the default constructor
    NAryMatIterator();
    //! the full constructor taking arbitrary number of n-dim matrices
    NAryMatIterator(const Mat** arrays, Mat* planes, int narrays=-1);
    //! the separate iterator initialization method
    void init(const Mat** arrays, Mat* planes, int narrays=-1);

    //! proceeds to the next plane of every iterated matrix
    NAryMatIterator& operator ++();
    //! proceeds to the next plane of every iterated matrix (postfix increment operator)
    NAryMatIterator operator ++(int);

    ...
    int nplanes; // the total number of planes
};
```

Use the class to implement unary, binary, and, generally, n-ary element-wise operations on multi-dimensional arrays. Some of the arguments of an n-ary function may be continuous arrays, some may be not. It is possible to use conventional `MatIterator` 's for each array but incrementing all of the iterators after each small operations may be a big overhead. In this case consider using `NAryMatIterator` to iterate through several matrices simultaneously as long as they have the same geometry (dimensionality and all the dimension sizes are the same). On each iteration `it.planes[0]`, `it.planes[1]` , ... will be the slices of the corresponding matrices.

The example below illustrates how you can compute a normalized and threshold 3D color histogram:

```cpp
void computeNormalizedColorHist(const Mat& image, Mat& hist, int N, double minProb)
{
    const int histSize[] = {N, N, N};

    // make sure that the histogram has a proper size and type
    hist.create(3, histSize, CV_32F);

    // and clear it
    hist = Scalar(0);
```

```
    // the loop below assumes that the image
    // is a 8-bit 3-channel. check it.
    CV_Assert(image.type() == CV_8UC3);
    MatConstIterator_<Vec3b> it = image.begin<Vec3b>(),
                             it_end = image.end<Vec3b>();
    for( ; it != it_end; ++it )
    {
        const Vec3b& pix = *it;
        hist.at<float>(pix[0]*N/256, pix[1]*N/256, pix[2]*N/256) += 1.f;
    }

    minProb *= image.rows*image.cols;
    Mat plane;
    NAryMatIterator it(&hist, &plane, 1);
    double s = 0;
    // iterate through the matrix. on each iteration
    // it.planes[*] (of type Mat) will be set to the current plane.
    for(int p = 0; p < it.nplanes; p++, ++it)
    {
        threshold(it.planes[0], it.planes[0], minProb, 0, THRESH_TOZERO);
        s += sum(it.planes[0])[0];
    }

    s = 1./s;
    it = NAryMatIterator(&hist, &plane, 1);
    for(int p = 0; p < it.nplanes; p++, ++it)
        it.planes[0] *= s;
}
```

## SparseMat

**class SparseMat**

Sparse n-dimensional array.

```
class SparseMat
{
public:
    typedef SparseMatIterator iterator;
    typedef SparseMatConstIterator const_iterator;

    // internal structure - sparse matrix header
    struct Hdr
    {
        ...
    };

    // sparse matrix node - element of a hash table
    struct Node
    {
        size_t hashval;
        size_t next;
        int idx[CV_MAX_DIM];
    };

    ////////// constructors and destructor //////////
    // default constructor
```

```
    SparseMat();
    // creates matrix of the specified size and type
    SparseMat(int dims, const int* _sizes, int _type);
    // copy constructor
    SparseMat(const SparseMat& m);
    // converts dense array to the sparse form,
    // if try1d is true and matrix is a single-column matrix (Nx1),
    // then the sparse matrix will be 1-dimensional.
    SparseMat(const Mat& m, bool try1d=false);
    // converts an old-style sparse matrix to the new style.
    // all the data is copied so that "m" can be safely
    // deleted after the conversion
    SparseMat(const CvSparseMat* m);
    // destructor
    ~SparseMat();

    ///////// assignment operations ///////////

    // this is an O(1) operation; no data is copied
    SparseMat& operator = (const SparseMat& m);
    // (equivalent to the corresponding constructor with try1d=false)
    SparseMat& operator = (const Mat& m);

    // creates a full copy of the matrix
    SparseMat clone() const;

    // copy all the data to the destination matrix.
    // the destination will be reallocated if needed.
    void copyTo( SparseMat& m ) const;
    // converts 1D or 2D sparse matrix to dense 2D matrix.
    // If the sparse matrix is 1D, the result will
    // be a single-column matrix.
    void copyTo( Mat& m ) const;
    // converts arbitrary sparse matrix to dense matrix.
    // multiplies all the matrix elements by the specified scalar
    void convertTo( SparseMat& m, int rtype, double alpha=1 ) const;
    // converts sparse matrix to dense matrix with optional type conversion and scaling.
    // When rtype=-1, the destination element type will be the same
    // as the sparse matrix element type.
    // Otherwise, rtype will specify the depth and
    // the number of channels will remain the same as in the sparse matrix
    void convertTo( Mat& m, int rtype, double alpha=1, double beta=0 ) const;

    // not used now
    void assignTo( SparseMat& m, int type=-1 ) const;

    // reallocates sparse matrix. If it was already of the proper size and type,
    // it is simply cleared with clear(), otherwise,
    // the old matrix is released (using release()) and the new one is allocated.
    void create(int dims, const int* _sizes, int _type);
    // sets all the matrix elements to 0, which means clearing the hash table.
    void clear();
    // manually increases reference counter to the header.
    void addref();
    // decreses the header reference counter when it reaches 0.
    // the header and all the underlying data are deallocated.
    void release();
```

```
// converts sparse matrix to the old-style representation.
// all the elements are copied.
operator CvSparseMat*() const;
// size of each element in bytes
// (the matrix nodes will be bigger because of
//   element indices and other SparseMat::Node elements).
size_t elemSize() const;
// elemSize()/channels()
size_t elemSize1() const;

// the same is in Mat
int type() const;
int depth() const;
int channels() const;

// returns the array of sizes and 0 if the matrix is not allocated
const int* size() const;
// returns i-th size (or 0)
int size(int i) const;
// returns the matrix dimensionality
int dims() const;
// returns the number of non-zero elements
size_t nzcount() const;

// compute element hash value from the element indices:
// 1D case
size_t hash(int i0) const;
// 2D case
size_t hash(int i0, int i1) const;
// 3D case
size_t hash(int i0, int i1, int i2) const;
// n-D case
size_t hash(const int* idx) const;

// low-level element-access functions,
// special variants for 1D, 2D, 3D cases, and the generic one for n-D case.
//
// return pointer to the matrix element.
//   if the element is there (it is non-zero), the pointer to it is returned
//   if it is not there and createMissing=false, NULL pointer is returned
//   if it is not there and createMissing=true, the new element
//     is created and initialized with 0. Pointer to it is returned.
//   If the optional hashval pointer is not NULL, the element hash value is
//   not computed but *hashval is taken instead.
uchar* ptr(int i0, bool createMissing, size_t* hashval=0);
uchar* ptr(int i0, int i1, bool createMissing, size_t* hashval=0);
uchar* ptr(int i0, int i1, int i2, bool createMissing, size_t* hashval=0);
uchar* ptr(const int* idx, bool createMissing, size_t* hashval=0);

// higher-level element access functions:
// ref<_Tp>(i0,...[,hashval]) - equivalent to *(_Tp*)ptr(i0,...true[,hashval]).
//    always return valid reference to the element.
//    If it does not exist, it is created.
// find<_Tp>(i0,...[,hashval]) - equivalent to (_const Tp*)ptr(i0,...false[,hashval]).
//    return pointer to the element or NULL pointer if the element is not there.
// value<_Tp>(i0,...[,hashval]) - equivalent to
//    { const _Tp* p = find<_Tp>(i0,...[,hashval]); return p ? *p : _Tp(); }
//    that is, 0 is returned when the element is not there.
```

```
    // note that _Tp must match the actual matrix type -
    // the functions do not do any on-fly type conversion

    // 1D case
    template<typename _Tp> _Tp& ref(int i0, size_t* hashval=0);
    template<typename _Tp> _Tp value(int i0, size_t* hashval=0) const;
    template<typename _Tp> const _Tp* find(int i0, size_t* hashval=0) const;

    // 2D case
    template<typename _Tp> _Tp& ref(int i0, int i1, size_t* hashval=0);
    template<typename _Tp> _Tp value(int i0, int i1, size_t* hashval=0) const;
    template<typename _Tp> const _Tp* find(int i0, int i1, size_t* hashval=0) const;

    // 3D case
    template<typename _Tp> _Tp& ref(int i0, int i1, int i2, size_t* hashval=0);
    template<typename _Tp> _Tp value(int i0, int i1, int i2, size_t* hashval=0) const;
    template<typename _Tp> const _Tp* find(int i0, int i1, int i2, size_t* hashval=0) const;

    // n-D case
    template<typename _Tp> _Tp& ref(const int* idx, size_t* hashval=0);
    template<typename _Tp> _Tp value(const int* idx, size_t* hashval=0) const;
    template<typename _Tp> const _Tp* find(const int* idx, size_t* hashval=0) const;

    // erase the specified matrix element.
    // when there is no such an element, the methods do nothing
    void erase(int i0, int i1, size_t* hashval=0);
    void erase(int i0, int i1, int i2, size_t* hashval=0);
    void erase(const int* idx, size_t* hashval=0);

    // return the matrix iterators,
    //   pointing to the first sparse matrix element,
    SparseMatIterator begin();
    SparseMatConstIterator begin() const;
    //   ... or to the point after the last sparse matrix element
    SparseMatIterator end();
    SparseMatConstIterator end() const;

    // and the template forms of the above methods.
    // _Tp must match the actual matrix type.
    template<typename _Tp> SparseMatIterator_<_Tp> begin();
    template<typename _Tp> SparseMatConstIterator_<_Tp> begin() const;
    template<typename _Tp> SparseMatIterator_<_Tp> end();
    template<typename _Tp> SparseMatConstIterator_<_Tp> end() const;

    // return value stored in the sparse martix node
    template<typename _Tp> _Tp& value(Node* n);
    template<typename _Tp> const _Tp& value(const Node* n) const;

    ////////////// some internally used methods ///////////////
    ...

    // pointer to the sparse matrix header
    Hdr* hdr;
};
```

The class `SparseMat` represents multi-dimensional sparse numerical arrays. Such a sparse array can store elements of any type that `Mat` can store. *Sparse* means that only non-zero elements are stored (though, as a result of operations on a sparse matrix, some of its stored elements can actually become 0. It is up to you to detect such elements and delete

them using `SparseMat::erase` ). The non-zero elements are stored in a hash table that grows when it is filled so that the search time is O(1) in average (regardless of whether element is there or not). Elements can be accessed using the following methods:

- Query operations ( `SparseMat::ptr` and the higher-level `SparseMat::ref`, `SparseMat::value` and `SparseMat::find` ), for example:

```
const int dims = 5;
int size[] = {10, 10, 10, 10, 10};
SparseMat sparse_mat(dims, size, CV_32F);
for(int i = 0; i < 1000; i++)
{
    int idx[dims];
    for(int k = 0; k < dims; k++)
        idx[k] = rand()
    sparse_mat.ref<float>(idx) += 1.f;
}
```

- Sparse matrix iterators. They are similar to `MatIterator` but different from `NAryMatIterator`. That is, the iteration loop is familiar to STL users:

```
// prints elements of a sparse floating-point matrix
// and the sum of elements.
SparseMatConstIterator_<float>
    it = sparse_mat.begin<float>(),
    it_end = sparse_mat.end<float>();
double s = 0;
int dims = sparse_mat.dims();
for(; it != it_end; ++it)
{
    // print element indices and the element value
    const SparseMat::Node* n = it.node();
    printf("(");
    for(int i = 0; i < dims; i++)
        printf("%d%s", n->idx[i], i < dims-1 ? ", " : ")");
    printf(": %g\n", it.value<float>());
    s += *it;
}
printf("Element sum is %g\n", s);
```

If you run this loop, you will notice that elements are not enumerated in a logical order (lexicographical, and so on). They come in the same order as they are stored in the hash table (semi-randomly). You may collect pointers to the nodes and sort them to get the proper ordering. Note, however, that pointers to the nodes may become invalid when you add more elements to the matrix. This may happen due to possible buffer reallocation.

- Combination of the above 2 methods when you need to process 2 or more sparse matrices simultaneously. For example, this is how you can compute unnormalized cross-correlation of the 2 floating-point sparse matrices:

```
double cross_corr(const SparseMat& a, const SparseMat& b)
{
    const SparseMat *_a = &a, *_b = &b;
    // if b contains less elements than a,
    // it is faster to iterate through b
    if(_a->nzcount() > _b->nzcount())
        std::swap(_a, _b);
    SparseMatConstIterator_<float> it = _a->begin<float>(),
                                   it_end = _a->end<float>();
    double ccorr = 0;
    for(; it != it_end; ++it)
```

```
    {
        // take the next element from the first matrix
        float avalue = *it;
        const Node* anode = it.node();
        // and try to find an element with the same index in the second matrix.
        // since the hash value depends only on the element index,
        // reuse the hash value stored in the node
        float bvalue = _b->value<float>(anode->idx,&anode->hashval);
        ccorr += avalue*bvalue;
    }
    return ccorr;
}
```

# SparseMat_

class **SparseMat_**

Template sparse n-dimensional array class derived from SparseMat

```
template<typename _Tp> class SparseMat_ : public SparseMat
{
public:
    typedef SparseMatIterator_<_Tp> iterator;
    typedef SparseMatConstIterator_<_Tp> const_iterator;

    // constructors;
    // the created matrix will have data type = DataType<_Tp>::type
    SparseMat_();
    SparseMat_(int dims, const int* _sizes);
    SparseMat_(const SparseMat& m);
    SparseMat_(const SparseMat_& m);
    SparseMat_(const Mat& m);
    SparseMat_(const CvSparseMat* m);
    // assignment operators; data type conversion is done when necessary
    SparseMat_& operator = (const SparseMat& m);
    SparseMat_& operator = (const SparseMat_& m);
    SparseMat_& operator = (const Mat& m);

    // equivalent to the correspoding parent class methods
    SparseMat_ clone() const;
    void create(int dims, const int* _sizes);
    operator CvSparseMat*() const;

    // overriden methods that do extra checks for the data type
    int type() const;
    int depth() const;
    int channels() const;

    // more convenient element access operations.
    // ref() is retained (but <_Tp> specification is not needed anymore);
    // operator () is equivalent to SparseMat::value<_Tp>
    _Tp& ref(int i0, size_t* hashval=0);
    _Tp operator()(int i0, size_t* hashval=0) const;
    _Tp& ref(int i0, int i1, size_t* hashval=0);
    _Tp operator()(int i0, int i1, size_t* hashval=0) const;
    _Tp& ref(int i0, int i1, int i2, size_t* hashval=0);
    _Tp operator()(int i0, int i1, int i2, size_t* hashval=0) const;
```

```
    _Tp& ref(const int* idx, size_t* hashval=0);
    _Tp operator()(const int* idx, size_t* hashval=0) const;

    // iterators
    SparseMatIterator_<_Tp> begin();
    SparseMatConstIterator_<_Tp> begin() const;
    SparseMatIterator_<_Tp> end();
    SparseMatConstIterator_<_Tp> end() const;
};
```

`SparseMat_` is a thin wrapper on top of `SparseMat` created in the same way as `Mat_` . It simplifies notation of some operations.

```
int sz[] = {10, 20, 30};
SparseMat_<double> M(3, sz);
...
M.ref(1, 2, 3) = M(4, 5, 6) + M(7, 8, 9);
```

## Algorithm

**class** `Algorithm`

This is a base class for all more or less complex algorithms in OpenCV, especially for classes of algorithms, for which there can be multiple implementations. The examples are stereo correspondence (for which there are algorithms like block matching, semi-global block matching, graph-cut etc.), background subtraction (which can be done using mixture-of-gaussians models, codebook-based algorithm etc.), optical flow (block matching, Lucas-Kanade, Horn-Schunck etc.).

The class provides the following features for all derived classes:

- so called "virtual constructor". That is, each Algorithm derivative is registered at program start and you can get the list of registered algorithms and create instance of a particular algorithm by its name (see `Algorithm::create`). If you plan to add your own algorithms, it is good practice to add a unique prefix to your algorithms to distinguish them from other algorithms.

- setting/retrieving algorithm parameters by name. If you used video capturing functionality from OpenCV highgui module, you are probably familar with `cvSetCaptureProperty()`, `cvGetCaptureProperty()`, `VideoCapture::set()` and `VideoCapture::get()`. `Algorithm` provides similar method where instead of integer id's you specify the parameter names as text strings. See `Algorithm::set` and `Algorithm::get` for details.

- reading and writing parameters from/to XML or YAML files. Every Algorithm derivative can store all its parameters and then read them back. There is no need to re-implement it each time.

Here is example of SIFT use in your application via Algorithm interface:

```
#include "opencv2/opencv.hpp"
#include "opencv2/nonfree/nonfree.hpp"

...

initModule_nonfree(); // to load SURF/SIFT etc.

Ptr<Feature2D> sift = Algorithm::create<Feature2D>("Feature2D.SIFT");

FileStorage fs("sift_params.xml", FileStorage::READ);
if( fs.isOpened() ) // if we have file with parameters, read them
{
```

```
    sift->read(fs["sift_params"]);
    fs.release();
}
else // else modify the parameters and store them; user can later edit the file to use different parameters
{
    sift->set("contrastThreshold", 0.01f); // lower the contrast threshold, compared to the default value

    {
    WriteStructContext ws(fs, "sift_params", CV_NODE_MAP);
    sift->write(fs);
    }
}

Mat image = imread("myimage.png", 0), descriptors;
vector<KeyPoint> keypoints;
(*sift)(image, noArray(), keypoints, descriptors);
```

## Algorithm::get

Returns the algorithm parameter

**C++:** template<typename _Tp> typename ParamType<_Tp>::member_type Algorithm::**get**(const string&
**name**) const

> **Parameters**
>
> > **name** – The parameter name.

The method returns value of the particular parameter. Since the compiler can not deduce the type of the returned
parameter, you should specify it explicitly in angle brackets. Here are the allowed forms of get:

- myalgo.get<int>("param_name")

- myalgo.get<double>("param_name")

- myalgo.get<bool>("param_name")

- myalgo.get<string>("param_name")

- myalgo.get<Mat>("param_name")

- myalgo.get<vector<Mat> >("param_name")

- myalgo.get<Algorithm>("param_name") (it returns Ptr<Algorithm>).

In some cases the actual type of the parameter can be cast to the specified type, e.g. integer parameter can be cast to
double, `bool` can be cast to `int`. But "dangerous" transformations (string<->number, double->int, 1x1 Mat<->number,
...) are not performed and the method will throw an exception. In the case of `Mat` or `vector<Mat>` parameters the
method does not clone the matrix data, so do not modify the matrices. Use `Algorithm::set` instead - slower, but
more safe.

## Algorithm::set

Sets the algorithm parameter

**C++:** void Algorithm::**set**(const string& **name**, int **value**)

**C++:** void Algorithm::**set**(const string& **name**, double **value**)

**C++:** void Algorithm::**set**(const string& **name**, bool **value**)

**C++:** void Algorithm::**set**(const string& **name**, const string& **value**)

**C++:** void Algorithm::**set**(const string& **name**, const Mat& **value**)

**C++:** void Algorithm::**set**(const string& **name**, const vector<Mat>& **value**)

**C++:** void Algorithm::**set**(const string& **name**, const Ptr<Algorithm>& **value**)

> **Parameters**

>> **name** – The parameter name.

>> **value** – The parameter value.

The method sets value of the particular parameter. Some of the algorithm parameters may be declared as read-only. If you try to set such a parameter, you will get exception with the corresponding error message.

## Algorithm::write

Stores algorithm parameters in a file storage

**C++:** void Algorithm::**write**(FileStorage& **fs**) const

> **Parameters**

>> **fs** – File storage.

The method stores all the algorithm parameters (in alphabetic order) to the file storage. The method is virtual. If you define your own Algorithm derivative, your can override the method and store some extra information. However, it's rarely needed. Here are some examples:

- SIFT feature detector (from nonfree module). The class only stores algorithm parameters and no keypoints or their descriptors. Therefore, it's enough to store the algorithm parameters, which is what `Algorithm::write()` does. Therefore, there is no dedicated `SIFT::write()`.

- Background subtractor (from video module). It has the algorithm parameters and also it has the current background model. However, the background model is not stored. First, it's rather big. Then, if you have stored the background model, it would likely become irrelevant on the next run (because of shifted camera, changed background, different lighting etc.). Therefore, `BackgroundSubtractorMOG` and `BackgroundSubtractorMOG2` also rely on the standard `Algorithm::write()` to store just the algorithm parameters.

- Expectation Maximization (from ml module). The algorithm finds mixture of gaussians that approximates user data best of all. In this case the model may be re-used on the next run to test new data against the trained statistical model. So EM needs to store the model. However, since the model is described by a few parameters that are available as read-only algorithm parameters (i.e. they are available via `EM::get()`), EM also relies on `Algorithm::write()` to store both EM parameters and the model (represented by read-only algorithm parameters).

## Algorithm::read

Reads algorithm parameters from a file storage

**C++:** void Algorithm::**read**(const FileNode& **fn**)

> **Parameters**

>> **fn** – File node of the file storage.

The method reads all the algorithm parameters from the specified node of a file storage. Similarly to `Algorithm::write()`, if you implement an algorithm that needs to read some extra data and/or re-compute some internal data, you may override the method.

## Algorithm::getList

Returns the list of registered algorithms

**C++:** void Algorithm::**getList**(vector<string>& **algorithms**)

> Parameters

>> **algorithms** – The output vector of algorithm names.

This static method returns the list of registered algorithms in alphabetical order. Here is how to use it

```
vector<string> algorithms;
Algorithm::getList(algorithms);
cout << "Algorithms: " << algorithms.size() << endl;
for (size_t i=0; i < algorithms.size(); i++)
    cout << algorithms[i] << endl;
```

## Algorithm::create

Creates algorithm instance by name

**C++:** template<typename _Tp> Ptr<_Tp> Algorithm::**create**(const string& **name**)

> Parameters

>> **name** – The algorithm name, one of the names returned by `Algorithm::getList()`.

This static method creates a new instance of the specified algorithm. If there is no such algorithm, the method will silently return null pointer (that can be checked by `Ptr::empty()` method). Also, you should specify the particular `Algorithm` subclass as `_Tp` (or simply `Algorithm` if you do not know it at that point).

```
Ptr<BackgroundSubtractor> bgfg = Algorithm::create<BackgroundSubtractor>("BackgroundSubtractor.MOG2");
```

**Note:** This is important note about seemingly mysterious behavior of `Algorithm::create()` when it returns NULL while it should not. The reason is simple - `Algorithm::create()` resides in OpenCV's core module and the algorithms are implemented in other modules. If you create algorithms dynamically, C++ linker may decide to throw away the modules where the actual algorithms are implemented, since you do not call any functions from the modules. To avoid this problem, you need to call `initModule_<modulename>();` somewhere in the beginning of the program before `Algorithm::create()`. For example, call `initModule_nonfree()` in order to use SURF/SIFT, call `initModule_ml()` to use expectation maximization etc.

## Creating Own Algorithms

The above methods are usually enough for users. If you want to make your own algorithm, derived from `Algorithm`, you should basically follow a few conventions and add a little semi-standard piece of code to your class:

- Make a class and specify `Algorithm` as its base class.

- The algorithm parameters should be the class members. See `Algorithm::get()` for the list of possible types of the parameters.

- Add public virtual method `AlgorithmInfo* info() const;` to your class.

- Add constructor function, `AlgorithmInfo` instance and implement the `info()` method. The simplest way is to take http://code.opencv.org/projects/opencv/repository/revisions/master/entry/modules/ml/src/ml_init.cpp as the reference and modify it according to the list of your parameters.

- Add some public function (e.g. `initModule_<mymodule>()`) that calls info() of your algorithm and put it into the same source file as `info()` implementation. This is to force C++ linker to include this object file into the target application. See `Algorithm::create()` for details.

## 2.2 Basic C Structures and Operations

The section describes the main data structures, used by the OpenCV 1.x API, and the basic functions to create and process the data structures.

### CvPoint

**C:** CvPoint **cvPoint**(int **x**, int **y**)
    constructs `CvPoint` structure.

**C:** CvPoint **cvPointFrom32f**(CvPoint2D32f **point**)
    converts `CvPoint2D32f` to `CvPoint`.

**struct `CvPoint`**
    2D point with integer coordinates (usually zero-based).

> **param x** x-coordinate of the point.
>
> **param y** y-coordinate of the point.
>
> **param point** the point to convert.

**See Also:**

Point_

### CvPoint2D32f

**C:** CvPoint2D32f **cvPoint2D32f**(double **x**, double **y**)
    constructs `CvPoint2D32f` structure.

**C:** CvPoint2D32f **cvPointTo32f**(CvPoint **point**)
    converts `CvPoint` to `CvPoint2D32f`.

**struct `CvPoint2D32f`**
    2D point with floating-point coordinates.

> **param x** floating-point x-coordinate of the point.
>
> **param y** floating-point y-coordinate of the point.
>
> **param point** the point to convert.

**See Also:**

Point_

### CvPoint3D32f

**struct `CvPoint3D32f`**
    3D point with floating-point coordinates

**C:** CvPoint3D32f **cvPoint3D32f**(double **x**, double **y**, double **z**)
constructs CvPoint3D32f structure.

> Parameters

>> **x** – floating-point x-coordinate of the point.

>> **y** – floating-point y-coordinate of the point.

>> **z** – floating-point z-coordinate of the point.

**See Also:**

Point3_

## CvPoint2D64f

struct **CvPoint2D64f**
2D point with double-precision floating-point coordinates.

**C:** CvPoint2D64f **cvPoint2D64f**(double **x**, double **y**)
constructs CvPoint2D64f structure.

> Parameters

>> **x** – double-precision floating-point x-coordinate of the point.

>> **y** – double-precision floating-point y-coordinate of the point.

**See Also:**

Point_

## CvPoint3D64f

struct **CvPoint3D64f**
3D point with double-precision floating-point coordinates.

**C:** CvPoint3D64f **cvPoint3D64f**(double **x**, double **y**, double **z**)
constructs CvPoint3D64f structure.

> Parameters

>> **x** – double-precision floating-point x-coordinate of the point.

>> **y** – double-precision floating-point y-coordinate of the point.

>> **z** – double-precision floating-point z-coordinate of the point.

**See Also:**

Point3_

## CvSize

struct **CvSize**
Size of a rectangle or an image.

**C:** CvSize **cvSize**(int **width**, int **height**)
constructs CvSize structure.

> Parameters

>            **width** – width of the rectangle.

>            **height** – height of the rectangle.

**See Also:**

Size_

## CvSize2D32f

**struct CvSize2D32f**
>     Sub-pixel accurate size of a rectangle.

**C:** CvSize2D32f **cvSize2D32f**(double **width**, double **height**)
>     constructs `CvSize2D32f` structure.

>            **Parameters**

>                    **width** – floating-point width of the rectangle.

>                    **height** – floating-point height of the rectangle.

**See Also:**

Size_

## CvRect

**struct CvRect**
>     Stores coordinates of a rectangle.

**C:** CvRect **cvRect**(int **x**, int **y**, int **width**, int **height**)
>     constructs `CvRect` structure.

>            **Parameters**

>                    **x** – x-coordinate of the top-left corner.

>                    **y** – y-coordinate of the top-left corner (sometimes bottom-left corner).

>                    **width** – width of the rectangle.

>                    **height** – height of the rectangle.

**See Also:**

Rect_

## CvBox2D

**struct CvBox2D**
>     Stores coordinates of a rotated rectangle.

>     CvPoint2D32f **center**
>         Center of the box

>     CvSize2D32f **size**
>         Box width and height

>     float **angle**
>         Angle between the horizontal axis and the first side (i.e. length) in degrees

**See Also:**

RotatedRect

## CvScalar

**struct CvScalar**

    A container for 1-,2-,3- or 4-tuples of doubles.

    double[4] **val**

**See Also:**

Scalar_

## CvTermCriteria

**struct CvTermCriteria**

    Termination criteria for iterative algorithms.

    int **type**

        type of the termination criteria, one of:

- CV_TERMCRIT_ITER - stop the algorithm after max_iter iterations at maximum.
- CV_TERMCRIT_EPS - stop the algorithm after the achieved algorithm-dependent accuracy becomes lower than epsilon.
- CV_TERMCRIT_ITER+CV_TERMCRIT_EPS - stop the algorithm after max_iter iterations or when the achieved accuracy is lower than epsilon, whichever comes the earliest.

    int **max_iter**

        Maximum number of iterations

    double **epsilon**

        Required accuracy

**See Also:**

TermCriteria

## CvMat

**struct CvMat**

    A multi-channel dense matrix.

    int **type**

        CvMat signature (CV_MAT_MAGIC_VAL) plus type of the elements. Type of the matrix elements can be retrieved using CV_MAT_TYPE macro:

```
int type = CV_MAT_TYPE(matrix->type);
```

        For description of possible matrix elements, see Mat.

    int **step**

        Full row length in bytes

    int* **refcount**

        Underlying data reference counter

union **data**
    Pointers to the actual matrix data:

  - ptr - pointer to 8-bit unsigned elements

  - s - pointer to 16-bit signed elements

  - i - pointer to 32-bit signed elements

  - fl - pointer to 32-bit floating-point elements

  - db - pointer to 64-bit floating-point elements

int **rows**
    Number of rows

int **cols**
    Number of columns

Matrix elements are stored row by row. Element (i, j) (i - 0-based row index, j - 0-based column index) of a matrix can be retrieved or modified using `CV_MAT_ELEM` macro:

```
uchar pixval = CV_MAT_ELEM(grayimg, uchar, i, j)
CV_MAT_ELEM(cameraMatrix, float, 0, 2) = image.width*0.5f;
```

To access multiple-channel matrices, you can use `CV_MAT_ELEM(matrix, type, i, j*nchannels + channel_idx)`.

CvMat is now obsolete; consider using `Mat` instead.


## CvMatND

**struct CvMatND**
    Multi-dimensional dense multi-channel array.

  int **type**
      A `CvMatND` signature (`CV_MATND_MAGIC_VAL`) plus the type of elements. Type of the matrix elements can be retrieved using `CV_MAT_TYPE` macro:

      ```
      int type = CV_MAT_TYPE(ndmatrix->type);
      ```

  int **dims**
      The number of array dimensions

  int* **refcount**
      Underlying data reference counter

  union **data**
      Pointers to the actual matrix data

    - ptr - pointer to 8-bit unsigned elements

    - s - pointer to 16-bit signed elements

    - i - pointer to 32-bit signed elements

    - fl - pointer to 32-bit floating-point elements

    - db - pointer to 64-bit floating-point elements

  array **dim**
      Arrays of pairs (array size along the i-th dimension, distance between neighbor elements along i-th dimension):

```
    for(int i = 0; i < ndmatrix->dims; i++)
        printf("size[i] = %d, step[i] = %d\n", ndmatrix->dim[i].size, ndmatrix->dim[i].step);
```

CvMatND is now obsolete; consider using Mat instead.

## CvSparseMat

**struct CvSparseMat**
>   Multi-dimensional sparse multi-channel array.

>   int **type**
>>      A CvSparseMat signature (CV_SPARSE_MAT_MAGIC_VAL) plus the type of sparse matrix elements. Similarly to CvMat and CvMatND, use CV_MAT_TYPE() to retrieve type of the elements.

>   int **dims**
>>      Number of dimensions

>   int* **refcount**
>>      Underlying reference counter. Not used.

>   CvSet* **heap**
>>      A pool of hash table nodes

>   void** **hashtable**
>>      The hash table. Each entry is a list of nodes.

>   int **hashsize**
>>      Size of the hash table

>   int[] **size**
>>      Array of dimension sizes

## IplImage

**struct IplImage**
>   IPL image header

>   int **nSize**
>>      sizeof(IplImage)

>   int **ID**
>>      Version, always equals 0

>   int **nChannels**
>>      Number of channels. Most OpenCV functions support 1-4 channels.

>   int **alphaChannel**
>>      Ignored by OpenCV

>   int **depth**
>>      Channel depth in bits + the optional sign bit ( IPL_DEPTH_SIGN ). The supported depths are:

>>      • IPL_DEPTH_8U - unsigned 8-bit integer. Equivalent to CV_8U in matrix types.

>>      • IPL_DEPTH_8S - signed 8-bit integer. Equivalent to CV_8S in matrix types.

>>      • IPL_DEPTH_16U - unsigned 16-bit integer. Equivalent to CV_16U in matrix types.

>>      • IPL_DEPTH_16S - signed 8-bit integer. Equivalent to CV_16S in matrix types.

>>      • IPL_DEPTH_32S - signed 32-bit integer. Equivalent to CV_32S in matrix types.

- IPL_DEPTH_32F - single-precision floating-point number. Equivalent to CV_32F in matrix types.

- IPL_DEPTH_64F - double-precision floating-point number. Equivalent to CV_64F in matrix types.

char[] **colorModel**
> Ignored by OpenCV.

char[] **channelSeq**
> Ignored by OpenCV

int **dataOrder**
> 0 = IPL_DATA_ORDER_PIXEL - interleaved color channels, 1 - separate color channels. CreateImage() only creates images with interleaved channels. For example, the usual layout of a color image is: $b_{00}g_{00}r_{00}b_{10}g_{10}r_{10}...$

int **origin**
> 0 - top-left origin, 1 - bottom-left origin (Windows bitmap style)

int **align**
> Alignment of image rows (4 or 8). OpenCV ignores this and uses widthStep instead.

int **width**
> Image width in pixels

int **height**
> Image height in pixels

IplROI* **roi**
> Region Of Interest (ROI). If not NULL, only this image region will be processed.

IplImage* **maskROI**
> Must be NULL in OpenCV

void* **imageId**
> Must be NULL in OpenCV

void* **tileInfo**
> Must be NULL in OpenCV

int **imageSize**
> Image data size in bytes. For interleaved data, this equals image->height · image->widthStep

char* **imageData**
> A pointer to the aligned image data. Do not assign imageData directly. Use SetData().

int **widthStep**
> The size of an aligned image row, in bytes.

int[] **BorderMode**
> Border completion mode, ignored by OpenCV

int[] **BorderConst**
> Constant border value, ignored by OpenCV

char* **imageDataOrigin**
> A pointer to the origin of the image data (not necessarily aligned). This is used for image deallocation.

The IplImage is taken from the Intel Image Processing Library, in which the format is native. OpenCV only supports a subset of possible IplImage formats, as outlined in the parameter list above.

In addition to the above restrictions, OpenCV handles ROIs differently. OpenCV functions require that the image size or ROI size of all source and destination images match exactly. On the other hand, the Intel Image Processing Library processes the area of intersection between the source and destination images (or ROIs), allowing them to vary independently.

## CvArr

**struct** `CvArr`

This is the "metatype" used *only* as a function parameter. It denotes that the function accepts arrays of multiple types, such as IplImage*, CvMat* or even CvSeq* sometimes. The particular array type is determined at runtime by analyzing the first 4 bytes of the header. In C++ interface the role of `CvArr` is played by `InputArray` and `OutputArray`.

## ClearND

Clears a specific array element.

**C:** void **cvClearND**(CvArr* **arr**, const int* **idx**)

**Python:** cv.**ClearND**(arr, idx) → None

> **Parameters**
>
> > **arr** – Input array
> >
> > **idx** – Array of the element indices

The function clears (sets to zero) a specific element of a dense array or deletes the element of a sparse array. If the sparse array element does not exists, the function does nothing.

## CloneImage

Makes a full copy of an image, including the header, data, and ROI.

**C:** IplImage* **cvCloneImage**(const IplImage* **image**)

**Python:** cv.**CloneImage**(image) → image

> **Parameters**
>
> > **image** – The original image

## CloneMat

Creates a full matrix copy.

**C:** CvMat* **cvCloneMat**(const CvMat* **mat**)

**Python:** cv.**CloneMat**(mat) → mat

> **Parameters**
>
> > **mat** – Matrix to be copied

Creates a full copy of a matrix and returns a pointer to the copy. Note that the matrix copy is compacted, that is, it will not have gaps between rows.

## CloneMatND

Creates full copy of a multi-dimensional array and returns a pointer to the copy.

**C:** CvMatND* **cvCloneMatND**(const CvMatND* **mat**)

**Python:** cv.**CloneMatND**(mat) → matND

> **Parameters**
>
>> **mat** – Input array

## CloneSparseMat

Creates full copy of sparse array.

**C:** CvSparseMat* **cvCloneSparseMat**(const CvSparseMat* **mat**)

> **Parameters**
>
>> **mat** – Input array

The function creates a copy of the input array and returns pointer to the copy.

## ConvertScale

Converts one array to another with optional linear transformation.

**C:** void **cvConvertScale**(const CvArr* **src**, CvArr* **dst**, double **scale**=1, double **shift**=0)

**Python:** cv.**ConvertScale**(src, dst, scale=1.0, shift=0.0) → None

**Python:** cv.**Convert**(src, dst) → None

```
#define cvCvtScale cvConvertScale
#define cvScale  cvConvertScale
#define cvConvert(src, dst )  cvConvertScale((src), (dst), 1, 0 )
```

> **Parameters**
>
>> **src** – Source array
>>
>> **dst** – Destination array
>>
>> **scale** – Scale factor
>>
>> **shift** – Value added to the scaled source array elements

The function has several different purposes, and thus has several different names. It copies one array to another with optional scaling, which is performed first, and/or optional type conversion, performed after:

$$\texttt{dst}(I) = \texttt{scale} \texttt{src}(I) + (\texttt{shift}_0, \texttt{shift}_1, ...)$$

All the channels of multi-channel arrays are processed independently.

The type of conversion is done with rounding and saturation, that is if the result of scaling + conversion can not be represented exactly by a value of the destination array element type, it is set to the nearest representable value on the real axis.

## Copy

Copies one array to another.

**C:** void **cvCopy**(const CvArr* **src**, CvArr* **dst**, const CvArr* **mask**=NULL)

**Python:** cv.**Copy**(src, dst, mask=None) → None

> **Parameters**
>
> > **src** – The source array
> >
> > **dst** – The destination array
> >
> > **mask** – Operation mask, 8-bit single channel array; specifies elements of the destination array to be changed

The function copies selected elements from an input array to an output array:

$$\text{dst}(I) = \text{src}(I) \quad \text{if} \quad \text{mask}(I) \neq 0.$$

If any of the passed arrays is of `IplImage` type, then its ROI and COI fields are used. Both arrays must have the same type, the same number of dimensions, and the same size. The function can also copy sparse arrays (mask is not supported in this case).

## CreateData

Allocates array data

**C:** void **cvCreateData**(CvArr* **arr**)

**Python:** cv.**CreateData**(arr) → None

> **Parameters**
>
> > **arr** – Array header

The function allocates image, matrix or multi-dimensional dense array data. Note that in the case of matrix types OpenCV allocation functions are used. In the case of IplImage they are used unless `CV_TURN_ON_IPL_COMPATIBILITY()` has been called before. In the latter case IPL functions are used to allocate the data.

## CreateImage

Creates an image header and allocates the image data.

**C:** IplImage* **cvCreateImage**(CvSize **size**, int **depth**, int **channels**)

**Python:** cv.**CreateImage**(size, depth, channels) → image

> **Parameters**
>
> > **size** – Image width and height
> >
> > **depth** – Bit depth of image elements. See `IplImage` for valid depths.
> >
> > **channels** – Number of channels per pixel. See `IplImage` for details. This function only creates images with interleaved channels.

This function call is equivalent to the following code:

```
header = cvCreateImageHeader(size, depth, channels);
cvCreateData(header);
```

## CreateImageHeader

Creates an image header but does not allocate the image data.

**C:** IplImage* **cvCreateImageHeader**(CvSize **size**, int **depth**, int **channels**)

**Python:** cv.**CreateImageHeader**(size, depth, channels) → image

> **Parameters**
>> **size** – Image width and height
>>
>> **depth** – Image depth (see `CreateImage()`)
>>
>> **channels** – Number of channels (see `CreateImage()`)

## CreateMat

Creates a matrix header and allocates the matrix data.

**C:** CvMat* **cvCreateMat**(int **rows**, int **cols**, int **type**)

**Python:** cv.**CreateMat**(rows, cols, type) → mat

> **Parameters**
>> **rows** – Number of rows in the matrix
>>
>> **cols** – Number of columns in the matrix
>>
>> **type** – The type of the matrix elements in the form `CV_<bit depth><S|U|F>C<number of channels>`, where S=signed, U=unsigned, F=float. For example, CV _ 8UC1 means the elements are 8-bit unsigned and the there is 1 channel, and CV _ 32SC2 means the elements are 32-bit signed and there are 2 channels.

The function call is equivalent to the following code:

```
CvMat* mat = cvCreateMatHeader(rows, cols, type);
cvCreateData(mat);
```

## CreateMatHeader

Creates a matrix header but does not allocate the matrix data.

**C:** CvMat* **cvCreateMatHeader**(int **rows**, int **cols**, int **type**)

**Python:** cv.**CreateMatHeader**(rows, cols, type) → mat

> **Parameters**
>> **rows** – Number of rows in the matrix
>>
>> **cols** – Number of columns in the matrix
>>
>> **type** – Type of the matrix elements, see `CreateMat()`

The function allocates a new matrix header and returns a pointer to it. The matrix data can then be allocated using `CreateData()` or set explicitly to user-allocated data via `SetData()`.

## CreateMatND

Creates the header and allocates the data for a multi-dimensional dense array.

**C:** CvMatND* **cvCreateMatND**(int **dims**, const int* **sizes**, int **type**)

**Python:** cv.**CreateMatND**(dims, type) → matND

      **Parameters**

            **dims** – Number of array dimensions. This must not exceed CV_MAX_DIM (32 by default, but can be changed at build time).

            **sizes** – Array of dimension sizes.

            **type** – Type of array elements, see CreateMat() .

This function call is equivalent to the following code:

```
CvMatND* mat = cvCreateMatNDHeader(dims, sizes, type);
cvCreateData(mat);
```

## CreateMatNDHeader

Creates a new matrix header but does not allocate the matrix data.

**C:** CvMatND* **cvCreateMatNDHeader**(int **dims**, const int* **sizes**, int **type**)

**Python:** cv.**CreateMatNDHeader**(dims, type) → matND

      **Parameters**

            **dims** – Number of array dimensions

            **sizes** – Array of dimension sizes

            **type** – Type of array elements, see CreateMat()

The function allocates a header for a multi-dimensional dense array. The array data can further be allocated using CreateData() or set explicitly to user-allocated data via SetData().

## CreateSparseMat

Creates sparse array.

**C:** CvSparseMat* **cvCreateSparseMat**(int **dims**, const int* **sizes**, int **type**)

      **Parameters**

            **dims** – Number of array dimensions. In contrast to the dense matrix, the number of dimensions is practically unlimited (up to $2^{16}$ ).

            **sizes** – Array of dimension sizes

            **type** – Type of array elements. The same as for CvMat

The function allocates a multi-dimensional sparse array. Initially the array contain no elements, that is PtrND() and other related functions will return 0 for every index.

## CrossProduct

Calculates the cross product of two 3D vectors.

**C:** void **cvCrossProduct**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**)

**Python:** cv.**CrossProduct**(src1, src2, dst) → None

> **Parameters**
>
> > **src1** – The first source vector
> >
> > **src2** – The second source vector
> >
> > **dst** – The destination vector

The function calculates the cross product of two 3D vectors:

$$dst = src1 \times src2$$

or:

$$dst_1 = src1_2 src2_3 - src1_3 src2_2$$
$$dst_2 = src1_3 src2_1 - src1_1 src2_3$$
$$dst_3 = src1_1 src2_2 - src1_2 src2_1$$

## DotProduct

Calculates the dot product of two arrays in Euclidean metrics.

**C:** double **cvDotProduct**(const CvArr* **src1**, const CvArr* **src2**)

**Python:** cv.**DotProduct**(src1, src2) → float

> **Parameters**
>
> > **src1** – The first source array
> >
> > **src2** – The second source array

The function calculates and returns the Euclidean dot product of two arrays.

$$src1 \bullet src2 = \sum_I (src1(I) src2(I))$$

In the case of multiple channel arrays, the results for all channels are accumulated. In particular, `cvDotProduct(a,a)` where a is a complex vector, will return $\|a\|^2$. The function can process multi-dimensional arrays, row by row, layer by layer, and so on.

## Get?D

**C:** CvScalar **cvGet1D**(const CvArr* **arr**, int **idx0**)

**C:** CvScalar **cvGet2D**(const CvArr* **arr**, int **idx0**, int **idx1**)

**C:** CvScalar **cvGet3D**(const CvArr* **arr**, int **idx0**, int **idx1**, int **idx2**)

**C:** CvScalar **cvGetND**(const CvArr* **arr**, const int* **idx**)

**Python:** cv.**Get1D**(arr, idx) → scalar

**Python:** cv.**Get2D**(arr, idx0, idx1) → scalar

**Python:** cv.**Get3D**(arr, idx0, idx1, idx2) → scalar

**Python:** cv.**GetND**(arr, indices) → scalar

Return a specific array element.

> **Parameters**

>> **arr** – Input array

>> **idx0** – The first zero-based component of the element index

>> **idx1** – The second zero-based component of the element index

>> **idx2** – The third zero-based component of the element index

>> **idx** – Array of the element indices

The functions return a specific array element. In the case of a sparse array the functions return 0 if the requested node does not exist (no new node is created by the functions).

## GetCol(s)

Returns one of more array columns.

**C:** CvMat* **cvGetCol**(const CvArr* **arr**, CvMat* **submat**, int **col**)

**C:** CvMat* **cvGetCols**(const CvArr* **arr**, CvMat* **submat**, int **start_col**, int **end_col**)

**Python:** cv.**GetCol**(arr, col) → submat

**Python:** cv.**GetCols**(arr, startCol, endCol) → submat

> **Parameters**

>> **arr** – Input array

>> **submat** – Pointer to the resulting sub-array header

>> **col** – Zero-based index of the selected column

>> **start_col** – Zero-based index of the starting column (inclusive) of the span

>> **end_col** – Zero-based index of the ending column (exclusive) of the span

The functions return the header, corresponding to a specified column span of the input array. That is, no data is copied. Therefore, any modifications of the submatrix will affect the original array. If you need to copy the columns, use CloneMat(). cvGetCol(arr, submat, col) is a shortcut for cvGetCols(arr, submat, col, col+1).

## GetDiag

Returns one of array diagonals.

**C:** CvMat* **cvGetDiag**(const CvArr* **arr**, CvMat* **submat**, int **diag**=0)

**Python:** cv.**GetDiag**(arr, diag=0) → submat

> **Parameters**

>> **arr** – Input array

>> **submat** – Pointer to the resulting sub-array header

>> **diag** – Index of the array diagonal. Zero value corresponds to the main diagonal, -1 corresponds to the diagonal above the main, 1 corresponds to the diagonal below the main, and so forth.

The function returns the header, corresponding to a specified diagonal of the input array.

## GetDims

Return number of array dimensions

**C:** int **cvGetDims**(const CvArr* **arr**, int* **sizes**=NULL)

**Python:** cv.**GetDims**(arr) -> (dim1, dim2, ...)

> **Parameters**
>
> > **arr** – Input array
> >
> > **sizes** – Optional output vector of the array dimension sizes. For 2d arrays the number of rows (height) goes first, number of columns (width) next.

The function returns the array dimensionality and the array of dimension sizes. In the case of `IplImage` or *CvMat* it always returns 2 regardless of number of image/matrix rows. For example, the following code calculates total number of array elements:

```
int sizes[CV_MAX_DIM];
int i, total = 1;
int dims = cvGetDims(arr, size);
for(i = 0; i < dims; i++ )
    total *= sizes[i];
```

## GetDimSize

Returns array size along the specified dimension.

**C:** int **cvGetDimSize**(const CvArr* **arr**, int **index**)

> **Parameters**
>
> > **arr** – Input array
> >
> > **index** – Zero-based dimension index (for matrices 0 means number of rows, 1 means number of columns; for images 0 means height, 1 means width)

## GetElemType

Returns type of array elements.

**C:** int **cvGetElemType**(const CvArr* **arr**)

**Python:** cv.**GetElemType**(arr) → int

> **Parameters**
>
> > **arr** – Input array

The function returns type of the array elements. In the case of `IplImage` the type is converted to `CvMat`-like representation. For example, if the image has been created as:

```
IplImage* img = cvCreateImage(cvSize(640, 480), IPL_DEPTH_8U, 3);
```

The code `cvGetElemType(img)` will return `CV_8UC3`.

## GetImage

Returns image header for arbitrary array.

**C:** IplImage* **cvGetImage**(const CvArr* **arr**, IplImage* **image_header**)

**Python:** cv.**GetImage**(arr) → iplimage

> **Parameters**
>
> > **arr** – Input array
> >
> > **image_header** – Pointer to `IplImage` structure used as a temporary buffer

The function returns the image header for the input array that can be a matrix (`CvMat`) or image (`IplImage`). In the case of an image the function simply returns the input pointer. In the case of `CvMat` it initializes an `image_header` structure with the parameters of the input matrix. Note that if we transform `IplImage` to `CvMat` using `GetMat()` and then transform `CvMat` back to IplImage using this function, we will get different headers if the ROI is set in the original image.

## GetImageCOI

Returns the index of the channel of interest.

**C:** int **cvGetImageCOI**(const IplImage* **image**)

**Python:** cv.**GetImageCOI**(image) → int

> **Parameters**
>
> > **image** – A pointer to the image header

Returns the channel of interest of in an IplImage. Returned values correspond to the `coi` in `SetImageCOI()`.

## GetImageROI

Returns the image ROI.

**C:** CvRect **cvGetImageROI**(const IplImage* **image**)

**Python:** cv.**GetImageROI**(image) → CvRect

> **Parameters**
>
> > **image** – A pointer to the image header

If there is no ROI set, `cvRect(0,0,image->width,image->height)` is returned.

## GetMat

Returns matrix header for arbitrary array.

**C:** CvMat* **cvGetMat**(const CvArr* **arr**, CvMat* **header**, int* **coi**=NULL, int **allowND**=0)

**Python:** cv.**GetMat**(arr, allowND=0) → mat

> **Parameters**
>
> > **arr** – Input array
> >
> > **header** – Pointer to `CvMat` structure used as a temporary buffer

coi – Optional output parameter for storing COI

**allowND** – If non-zero, the function accepts multi-dimensional dense arrays (CvMatND*) and returns 2D matrix (if CvMatND has two dimensions) or 1D matrix (when CvMatND has 1 dimension or more than 2 dimensions). The CvMatND array must be continuous.

The function returns a matrix header for the input array that can be a matrix - CvMat, an image - IplImage, or a multi-dimensional dense array - CvMatND (the third option is allowed only if allowND != 0) . In the case of matrix the function simply returns the input pointer. In the case of IplImage* or CvMatND it initializes the header structure with parameters of the current image ROI and returns &header. Because COI is not supported by CvMat, it is returned separately.

The function provides an easy way to handle both types of arrays - IplImage and CvMat using the same code. Input array must have non-zero data pointer, otherwise the function will report an error.

**See Also:**

GetImage(), cvarrToMat().

---

**Note:** If the input array is IplImage with planar data layout and COI set, the function returns the pointer to the selected plane and COI == 0. This feature allows user to process IplImage structures with planar data layout, even though OpenCV does not support such images.

---

## GetNextSparseNode

Returns the next sparse matrix element

**C:** CvSparseNode* **cvGetNextSparseNode**(CvSparseMatIterator* **mat_iterator**)

> **Parameters**
>
> > **mat_iterator** – Sparse array iterator

The function moves iterator to the next sparse matrix element and returns pointer to it. In the current version there is no any particular order of the elements, because they are stored in the hash table. The sample below demonstrates how to iterate through the sparse matrix:

```
// print all the non-zero sparse matrix elements and compute their sum
double sum = 0;
int i, dims = cvGetDims(sparsemat);
CvSparseMatIterator it;
CvSparseNode* node = cvInitSparseMatIterator(sparsemat, &it);

for(; node != 0; node = cvGetNextSparseNode(&it))
{
    /* get pointer to the element indices */
    int* idx = CV_NODE_IDX(array, node);
    /* get value of the element (assume that the type is CV_32FC1) */
    float val = *(float*)CV_NODE_VAL(array, node);
    printf("M");
    for(i = 0; i < dims; i++ )
        printf("[%d]", idx[i]);
    printf("=%g\n", val);

    sum += val;
}

printf("nTotal sum = %g\n", sum);
```

## GetRawData

Retrieves low-level information about the array.

**C:** void **cvGetRawData**(const CvArr* **arr**, uchar** **data**, int* **step**=NULL, CvSize* **roi_size**=NULL )

> **Parameters**
>
>> **arr** – Array header
>>
>> **data** – Output pointer to the whole image origin or ROI origin if ROI is set
>>
>> **step** – Output full row length in bytes
>>
>> **roi_size** – Output ROI size

The function fills output variables with low-level information about the array data. All output parameters are optional, so some of the pointers may be set to NULL. If the array is IplImage with ROI set, the parameters of ROI are returned.

The following example shows how to get access to array elements. It computes absolute values of the array elements

```
float* data;
int step;
CvSize size;

cvGetRawData(array, (uchar**)&data, &step, &size);
step /= sizeof(data[0]);

for(int y = 0; y < size.height; y++, data += step )
    for(int x = 0; x < size.width; x++ )
        data[x] = (float)fabs(data[x]);
```

## GetReal?D

Return a specific element of single-channel 1D, 2D, 3D or nD array.

**C:** double **cvGetReal1D**(const CvArr* **arr**, int **idx0**)

**C:** double **cvGetReal2D**(const CvArr* **arr**, int **idx0**, int **idx1**)

**C:** double **cvGetReal3D**(const CvArr* **arr**, int **idx0**, int **idx1**, int **idx2**)

**C:** double **cvGetRealND**(const CvArr* **arr**, const int* **idx**)

**Python:** cv.**GetReal1D**(arr, idx0) → float

**Python:** cv.**GetReal2D**(arr, idx0, idx1) → float

**Python:** cv.**GetReal3D**(arr, idx0, idx1, idx2) → float

**Python:** cv.**GetRealND**(arr, idx) → float

> **Parameters**
>
>> **arr** – Input array. Must have a single channel.
>>
>> **idx0** – The first zero-based component of the element index
>>
>> **idx1** – The second zero-based component of the element index
>>
>> **idx2** – The third zero-based component of the element index
>>
>> **idx** – Array of the element indices

Returns a specific element of a single-channel array. If the array has multiple channels, a runtime error is raised. Note that `Get?D` functions can be used safely for both single-channel and multiple-channel arrays though they are a bit slower.

In the case of a sparse array the functions return 0 if the requested node does not exist (no new node is created by the functions).

## GetRow(s)

Returns array row or row span.

**C:** CvMat* **cvGetRow**(const CvArr* **arr**, CvMat* **submat**, int **row**)

**C:** CvMat* **cvGetRows**(const CvArr* **arr**, CvMat* **submat**, int **start_row**, int **end_row**, int **delta_row**=1 )

**Python:** cv.**GetRow**(arr, row) → submat

**Python:** cv.**GetRows**(arr, startRow, endRow, deltaRow=1) → submat

> **Parameters**

>> **arr** – Input array

>> **submat** – Pointer to the resulting sub-array header

>> **row** – Zero-based index of the selected row

>> **start_row** – Zero-based index of the starting row (inclusive) of the span

>> **end_row** – Zero-based index of the ending row (exclusive) of the span

>> **delta_row** – Index step in the row span. That is, the function extracts every `delta_row` -th row from `start_row` and up to (but not including) `end_row` .

The functions return the header, corresponding to a specified row/row span of the input array. `cvGetRow(arr, submat, row)` is a shortcut for `cvGetRows(arr, submat, row, row+1)`.

## GetSize

Returns size of matrix or image ROI.

**C:** CvSize **cvGetSize**(const CvArr* **arr**)

**Python:** cv.**GetSize**(arr)-> (width, height)

> **Parameters**

>> **arr** – array header

The function returns number of rows (CvSize::height) and number of columns (CvSize::width) of the input matrix or image. In the case of image the size of ROI is returned.

## GetSubRect

Returns matrix header corresponding to the rectangular sub-array of input image or matrix.

**C:** CvMat* **cvGetSubRect**(const CvArr* **arr**, CvMat* **submat**, CvRect **rect**)

**Python:** cv.**GetSubRect**(arr, rect) → submat

> **Parameters**

> **arr** – Input array
>
> **submat** – Pointer to the resultant sub-array header
>
> **rect** – Zero-based coordinates of the rectangle of interest

The function returns header, corresponding to a specified rectangle of the input array. In other words, it allows the user to treat a rectangular part of input array as a stand-alone array. ROI is taken into account by the function so the sub-array of ROI is actually extracted.

## DecRefData

Decrements an array data reference counter.

**C:** void **cvDecRefData**(CvArr* **arr**)

> **Parameters**
>
> > **arr** – Pointer to an array header

The function decrements the data reference counter in a `CvMat` or `CvMatND` if the reference counter pointer is not NULL. If the counter reaches zero, the data is deallocated. In the current implementation the reference counter is not NULL only if the data was allocated using the `CreateData()` function. The counter will be NULL in other cases such as: external data was assigned to the header using `SetData()`, header is part of a larger matrix or image, or the header was converted from an image or n-dimensional matrix header.

## IncRefData

Increments array data reference counter.

**C:** int **cvIncRefData**(CvArr* **arr**)

> **Parameters**
>
> > **arr** – Array header

The function increments `CvMat` or `CvMatND` data reference counter and returns the new counter value if the reference counter pointer is not NULL, otherwise it returns zero.

## InitImageHeader

Initializes an image header that was previously allocated.

**C:** IplImage* **cvInitImageHeader**(IplImage* **image**, CvSize **size**, int **depth**, int **channels**, int **origin**=0, int **align**=4)

> **Parameters**
>
> > **image** – Image header to initialize
> >
> > **size** – Image width and height
> >
> > **depth** – Image depth (see `CreateImage()`)
> >
> > **channels** – Number of channels (see `CreateImage()`)
> >
> > **origin** – Top-left `IPL_ORIGIN_TL` or bottom-left `IPL_ORIGIN_BL`
> >
> > **align** – Alignment for image rows, typically 4 or 8 bytes

The returned `IplImage*` points to the initialized header.

---

## InitMatHeader

Initializes a pre-allocated matrix header.

**C:** CvMat* **cvInitMatHeader**(CvMat* **mat**, int **rows**, int **cols**, int **type**, void* **data**=NULL, int **step**=CV_AUTOSTEP)

> **Parameters**
>
>> **mat** – A pointer to the matrix header to be initialized
>>
>> **rows** – Number of rows in the matrix
>>
>> **cols** – Number of columns in the matrix
>>
>> **type** – Type of the matrix elements, see `CreateMat()` .
>>
>> **data** – Optional: data pointer assigned to the matrix header
>>
>> **step** – Optional: full row width in bytes of the assigned data. By default, the minimal possible step is used which assumes there are no gaps between subsequent rows of the matrix.

This function is often used to process raw data with OpenCV matrix functions. For example, the following code computes the matrix product of two matrices, stored as ordinary arrays:

```
double a[] = { 1, 2, 3, 4,
               5, 6, 7, 8,
               9, 10, 11, 12 };

double b[] = { 1, 5, 9,
               2, 6, 10,
               3, 7, 11,
               4, 8, 12 };

double c[9];
CvMat Ma, Mb, Mc ;

cvInitMatHeader(&Ma, 3, 4, CV_64FC1, a);
cvInitMatHeader(&Mb, 4, 3, CV_64FC1, b);
cvInitMatHeader(&Mc, 3, 3, CV_64FC1, c);

cvMatMulAdd(&Ma, &Mb, 0, &Mc);
// the c array now contains the product of a (3x4) and b (4x3)
```

## InitMatNDHeader

Initializes a pre-allocated multi-dimensional array header.

**C:** CvMatND* **cvInitMatNDHeader**(CvMatND* **mat**, int **dims**, const int* **sizes**, int **type**, void* **data**=NULL)

> **Parameters**
>
>> **mat** – A pointer to the array header to be initialized
>>
>> **dims** – The number of array dimensions
>>
>> **sizes** – An array of dimension sizes
>>
>> **type** – Type of array elements, see `CreateMat()`
>>
>> **data** – Optional data pointer assigned to the matrix header

## InitSparseMatIterator

Initializes sparse array elements iterator.

**C:** CvSparseNode* **cvInitSparseMatIterator**(const CvSparseMat* **mat**, CvSparseMatIterator* **mat_iterator**)

> **Parameters**
>
> > **mat** – Input array
> >
> > **mat_iterator** – Initialized iterator

The function initializes iterator of sparse array elements and returns pointer to the first element, or NULL if the array is empty.

## Mat

Initializes matrix header (lightweight variant).

**C:** CvMat **cvMat**(int **rows**, int **cols**, int **type**, void* **data**=NULL)

> **Parameters**
>
> > **rows** – Number of rows in the matrix
> >
> > **cols** – Number of columns in the matrix
> >
> > **type** – Type of the matrix elements - see `CreateMat()`
> >
> > **data** – Optional data pointer assigned to the matrix header

Initializes a matrix header and assigns data to it. The matrix is filled *row*-wise (the first `cols` elements of data form the first row of the matrix, etc.)

This function is a fast inline substitution for `InitMatHeader()`. Namely, it is equivalent to:

```
CvMat mat;
cvInitMatHeader(&mat, rows, cols, type, data, CV_AUTOSTEP);
```

## Ptr?D

Return pointer to a particular array element.

**C:** uchar* **cvPtr1D**(const CvArr* **arr**, int **idx0**, int* **type**=NULL)

**C:** uchar* **cvPtr2D**(const CvArr* **arr**, int **idx0**, int **idx1**, int* **type**=NULL)

**C:** uchar* **cvPtr3D**(const CvArr* **arr**, int **idx0**, int **idx1**, int **idx2**, int* **type**=NULL)

**C:** uchar* **cvPtrND**(const CvArr* **arr**, const int* **idx**, int* **type**=NULL, int **create_node**=1, unsigned int* **precalc_hashval**=NULL )

> **Parameters**
>
> > **arr** – Input array
> >
> > **idx0** – The first zero-based component of the element index
> >
> > **idx1** – The second zero-based component of the element index
> >
> > **idx2** – The third zero-based component of the element index
> >
> > **idx** – Array of the element indices

**type** – Optional output parameter: type of matrix elements

**create_node** – Optional input parameter for sparse matrices. Non-zero value of the parameter means that the requested element is created if it does not exist already.

**precalc_hashval** – Optional input parameter for sparse matrices. If the pointer is not NULL, the function does not recalculate the node hash value, but takes it from the specified location. It is useful for speeding up pair-wise operations (TODO: provide an example)

The functions return a pointer to a specific array element. Number of array dimension should match to the number of indices passed to the function except for `cvPtr1D` function that can be used for sequential access to 1D, 2D or nD dense arrays.

The functions can be used for sparse arrays as well - if the requested node does not exist they create it and set it to zero.

All these as well as other functions accessing array elements ( `GetND()` , `GetRealND()` , `Set()` , `SetND()` , `SetRealND()` ) raise an error in case if the element index is out of range.

## ReleaseData

Releases array data.

**C:** void **cvReleaseData**(CvArr* **arr**)

> **Parameters**

>> **arr** – Array header

The function releases the array data. In the case of `CvMat` or `CvMatND` it simply calls cvDecRefData(), that is the function can not deallocate external data. See also the note to `CreateData()` .

## ReleaseImage

Deallocates the image header and the image data.

**C:** void **cvReleaseImage**(IplImage** **image**)

> **Parameters**

>> **image** – Double pointer to the image header

This call is a shortened form of

```
if(*image )
{
    cvReleaseData(*image);
    cvReleaseImageHeader(image);
}
```

## ReleaseImageHeader

Deallocates an image header.

**C:** void **cvReleaseImageHeader**(IplImage** **image**)

> **Parameters**

>> **image** – Double pointer to the image header

This call is an analogue of

```
if(image )
{
    iplDeallocate(*image, IPL_IMAGE_HEADER | IPL_IMAGE_ROI);
    *image = 0;
}
```

but it does not use IPL functions by default (see the `CV_TURN_ON_IPL_COMPATIBILITY` macro).

## ReleaseMat

Deallocates a matrix.

**C:** void **cvReleaseMat**(CvMat** **mat**)

        **Parameters**

                **mat** – Double pointer to the matrix

The function decrements the matrix data reference counter and deallocates matrix header. If the data reference counter is 0, it also deallocates the data.

```
if(*mat )
    cvDecRefData(*mat);
cvFree((void**)mat);
```

## ReleaseMatND

Deallocates a multi-dimensional array.

**C:** void **cvReleaseMatND**(CvMatND** **mat**)

        **Parameters**

                **mat** – Double pointer to the array

The function decrements the array data reference counter and releases the array header. If the reference counter reaches 0, it also deallocates the data.

```
if(*mat )
    cvDecRefData(*mat);
cvFree((void**)mat);
```

## ReleaseSparseMat

Deallocates sparse array.

**C:** void **cvReleaseSparseMat**(CvSparseMat** **mat**)

        **Parameters**

                **mat** – Double pointer to the array

The function releases the sparse array and clears the array pointer upon exit.

## ResetImageROI

Resets the image ROI to include the entire image and releases the ROI structure.

**C:** void **cvResetImageROI**(IplImage* **image**)

**Python:** cv.**ResetImageROI**(image) → None

> **Parameters**
>
> > **image** – A pointer to the image header

This produces a similar result to the following, but in addition it releases the ROI structure.

```
cvSetImageROI(image, cvRect(0, 0, image->width, image->height ));
cvSetImageCOI(image, 0);
```

## Reshape

Changes shape of matrix/image without copying data.

**C:** CvMat* **cvReshape**(const CvArr* **arr**, CvMat* **header**, int **new_cn**, int **new_rows**=0 )

**Python:** cv.**Reshape**(arr, newCn, newRows=0) → mat

> **Parameters**
>
> > **arr** – Input array
> >
> > **header** – Output header to be filled
> >
> > **new_cn** – New number of channels. 'new_cn = 0' means that the number of channels remains unchanged.
> >
> > **new_rows** – New number of rows. 'new_rows = 0' means that the number of rows remains unchanged unless it needs to be changed according to new_cn value.

The function initializes the CvMat header so that it points to the same data as the original array but has a different shape - different number of channels, different number of rows, or both.

The following example code creates one image buffer and two image headers, the first is for a 320x240x3 image and the second is for a 960x240x1 image:

```
IplImage* color_img = cvCreateImage(cvSize(320,240), IPL_DEPTH_8U, 3);
CvMat gray_mat_hdr;
IplImage gray_img_hdr, *gray_img;
cvReshape(color_img, &gray_mat_hdr, 1);
gray_img = cvGetImage(&gray_mat_hdr, &gray_img_hdr);
```

And the next example converts a 3x3 matrix to a single 1x9 vector:

```
CvMat* mat = cvCreateMat(3, 3, CV_32F);
CvMat row_header, *row;
row = cvReshape(mat, &row_header, 0, 1);
```

## ReshapeMatND

Changes the shape of a multi-dimensional array without copying the data.

**C:** CvArr* **cvReshapeMatND**(const CvArr* **arr**, int **sizeof_header**, CvArr* **header**, int **new_cn**, int **new_dims**, int* **new_sizes**)

---

**Python:** cv.**ReshapeMatND**(arr, newCn, newDims) → mat

>> **Parameters**

>>> **arr** – Input array

>>> **sizeof_header** – Size of output header to distinguish between IplImage, CvMat and Cv-MatND output headers

>>> **header** – Output header to be filled

>>> **new_cn** – New number of channels. `new_cn = 0` means that the number of channels remains unchanged.

>>> **new_dims** – New number of dimensions. `new_dims = 0` means that the number of dimensions remains the same.

>>> **new_sizes** – Array of new dimension sizes. Only `new_dims-1` values are used, because the total number of elements must remain the same. Thus, if `new_dims = 1`, `new_sizes` array is not used.

The function is an advanced version of Reshape() that can work with multi-dimensional arrays as well (though it can work with ordinary images and matrices) and change the number of dimensions.

Below are the two samples from the Reshape() description rewritten using ReshapeMatND():

```
IplImage* color_img = cvCreateImage(cvSize(320,240), IPL_DEPTH_8U, 3);
IplImage gray_img_hdr, *gray_img;
gray_img = (IplImage*)cvReshapeND(color_img, &gray_img_hdr, 1, 0, 0);


...

/* second example is modified to convert 2x2x2 array to 8x1 vector */
int size[] = { 2, 2, 2 };
CvMatND* mat = cvCreateMatND(3, size, CV_32F);
CvMat row_header, *row;
row = (CvMat*)cvReshapeND(mat, &row_header, 0, 1, 0);
```

## Set

Sets every element of an array to a given value.

**C:** void **cvSet**(CvArr* **arr**, CvScalar **value**, const CvArr* **mask**=NULL)

**Python:** cv.**Set**(arr, value, mask=None) → None

>> **Parameters**

>>> **arr** – The destination array

>>> **value** – Fill value

>>> **mask** – Operation mask, 8-bit single channel array; specifies elements of the destination array to be changed

The function copies the scalar `value` to every selected element of the destination array:

$$\mathtt{arr}(I) = \mathtt{value} \quad \text{if} \quad \mathtt{mask}(I) \neq 0$$

If array `arr` is of IplImage type, then is ROI used, but COI must not be set.

## Set?D

Change the particular array element.

**C:** void **cvSet1D**(CvArr* **arr**, int **idx0**, CvScalar **value**)

**C:** void **cvSet2D**(CvArr* **arr**, int **idx0**, int **idx1**, CvScalar **value**)

**C:** void **cvSet3D**(CvArr* **arr**, int **idx0**, int **idx1**, int **idx2**, CvScalar **value**)

**C:** void **cvSetND**(CvArr* **arr**, const int* **idx**, CvScalar **value**)

**Python:** cv.**Set1D**(arr, idx, value) → None

**Python:** cv.**Set2D**(arr, idx0, idx1, value) → None

**Python:** cv.**Set3D**(arr, idx0, idx1, idx2, value) → None

**Python:** cv.**SetND**(arr, indices, value) → None

> **Parameters**
>> **arr** – Input array
>>
>> **idx0** – The first zero-based component of the element index
>>
>> **idx1** – The second zero-based component of the element index
>>
>> **idx2** – The third zero-based component of the element index
>>
>> **idx** – Array of the element indices
>>
>> **value** – The assigned value

The functions assign the new value to a particular array element. In the case of a sparse array the functions create the node if it does not exist yet.

## SetData

Assigns user data to the array header.

**C:** void **cvSetData**(CvArr* **arr**, void* **data**, int **step**)

**Python:** cv.**SetData**(arr, data, step) → None

> **Parameters**
>> **arr** – Array header
>>
>> **data** – User data
>>
>> **step** – Full row length in bytes

The function assigns user data to the array header. Header should be initialized before using `cvCreateMatHeader()`, `cvCreateImageHeader()`, `cvCreateMatNDHeader()`, `cvInitMatHeader()`, `cvInitImageHeader()` or `cvInitMatNDHeader()`.

## SetImageCOI

Sets the channel of interest in an IplImage.

**C:** void **cvSetImageCOI**(IplImage* **image**, int **coi**)

**Python:** cv.**SetImageCOI**(image, coi) → None

**Parameters**

> **image** – A pointer to the image header
>
> **coi** – The channel of interest. 0 - all channels are selected, 1 - first channel is selected, etc. Note that the channel indices become 1-based.

If the ROI is set to `NULL` and the coi is *not* 0, the ROI is allocated. Most OpenCV functions do *not* support the COI setting, so to process an individual image/matrix channel one may copy (via `Copy()` or `Split()`) the channel to a separate image/matrix, process it and then copy the result back (via `Copy()` or `Merge()`) if needed.

## SetImageROI

Sets an image Region Of Interest (ROI) for a given rectangle.

**C:** void **cvSetImageROI**(IplImage* **image**, CvRect **rect**)

**Python:** cv.**SetImageROI**(image, rect) → None

> **Parameters**
>
> > **image** – A pointer to the image header
> >
> > **rect** – The ROI rectangle

If the original image ROI was `NULL` and the `rect` is not the whole image, the ROI structure is allocated.

Most OpenCV functions support the use of ROI and treat the image rectangle as a separate image. For example, all of the pixel coordinates are counted from the top-left (or bottom-left) corner of the ROI, not the original image.

## SetReal?D

Change a specific array element.

**C:** void **cvSetReal1D**(CvArr* **arr**, int **idx0**, double **value**)

**C:** void **cvSetReal2D**(CvArr* **arr**, int **idx0**, int **idx1**, double **value**)

**C:** void **cvSetReal3D**(CvArr* **arr**, int **idx0**, int **idx1**, int **idx2**, double **value**)

**C:** void **cvSetRealND**(CvArr* **arr**, const int* **idx**, double **value**)

**Python:** cv.**SetReal1D**(arr, idx, value) → None

**Python:** cv.**SetReal2D**(arr, idx0, idx1, value) → None

**Python:** cv.**SetReal3D**(arr, idx0, idx1, idx2, value) → None

**Python:** cv.**SetRealND**(arr, indices, value) → None

> **Parameters**
>
> > **arr** – Input array
> >
> > **idx0** – The first zero-based component of the element index
> >
> > **idx1** – The second zero-based component of the element index
> >
> > **idx2** – The third zero-based component of the element index
> >
> > **idx** – Array of the element indices
> >
> > **value** – The assigned value

The functions assign a new value to a specific element of a single-channel array. If the array has multiple channels, a runtime error is raised. Note that the `Set*D` function can be used safely for both single-channel and multiple-channel arrays, though they are a bit slower.

In the case of a sparse array the functions create the node if it does not yet exist.

## SetZero

Clears the array.

**C:** void **cvSetZero**(CvArr* **arr**)

**Python:** cv.**SetZero**(arr) → None

> **Parameters**
>
> > **arr** – Array to be cleared

The function clears the array. In the case of dense arrays (CvMat, CvMatND or IplImage), cvZero(array) is equivalent to cvSet(array,cvScalarAll(0),0). In the case of sparse arrays all the elements are removed.

## mGet

Returns the particular element of single-channel floating-point matrix.

**C:** double **cvmGet**(const CvMat* **mat**, int **row**, int **col**)

**Python:** cv.**mGet**(mat, row, col) → float

> **Parameters**
>
> > **mat** – Input matrix
> >
> > **row** – The zero-based index of row
> >
> > **col** – The zero-based index of column

The function is a fast replacement for GetReal2D() in the case of single-channel floating-point matrices. It is faster because it is inline, it does fewer checks for array type and array element type, and it checks for the row and column ranges only in debug mode.

## mSet

Sets a specific element of a single-channel floating-point matrix.

**C:** void **cvmSet**(CvMat* **mat**, int **row**, int **col**, double **value**)

**Python:** cv.**mSet**(mat, row, col, value) → None

> **Parameters**
>
> > **mat** – The matrix
> >
> > **row** – The zero-based index of row
> >
> > **col** – The zero-based index of column
> >
> > **value** – The new value of the matrix element

The function is a fast replacement for SetReal2D() in the case of single-channel floating-point matrices. It is faster because it is inline, it does fewer checks for array type and array element type, and it checks for the row and column ranges only in debug mode.

## SetIPLAllocators

Makes OpenCV use IPL functions for allocating IplImage and IplROI structures.

**C:** void **cvSetIPLAllocators**(Cv_iplCreateImageHeader **create_header**, Cv_iplAllocateImageData **allocate_data**, Cv_iplDeallocate **deallocate**, Cv_iplCreateROI **create_roi**, Cv_iplCloneImage **clone_image**)

> **Parameters**
>
>> **create_header** – pointer to a function, creating IPL image header.
>>
>> **allocate_data** – pointer to a function, allocating IPL image data.
>>
>> **deallocate** – pointer to a function, deallocating IPL image.
>>
>> **create_roi** – pointer to a function, creating IPL image ROI (i.e. Region of Interest).
>>
>> **clone_image** – pointer to a function, cloning an IPL image.

Normally, the function is not called directly. Instead, a simple macro `CV_TURN_ON_IPL_COMPATIBILITY()` is used that calls `cvSetIPLAllocators` and passes there pointers to IPL allocation functions.

```
...
CV_TURN_ON_IPL_COMPATIBILITY()
...
```

## RNG

Initializes a random number generator state.

**C:** CvRNG **cvRNG**(int64 **seed**=-1)

**Python:** cv.**RNG**(seed=-1LL) → CvRNG

> **Parameters**
>
>> **seed** – 64-bit value used to initiate a random sequence

The function initializes a random number generator and returns the state. The pointer to the state can be then passed to the `RandInt()`, `RandReal()` and `RandArr()` functions. In the current implementation a multiply-with-carry generator is used.

**See Also:**

the C++ class `RNG` replaced `CvRNG`.

## RandArr

Fills an array with random numbers and updates the RNG state.

**C:** void **cvRandArr**(CvRNG* **rng**, CvArr* **arr**, int **dist_type**, CvScalar **param1**, CvScalar **param2**)

**Python:** cv.**RandArr**(rng, arr, distType, param1, param2) → None

> **Parameters**
>
>> **rng** – CvRNG state initialized by `RNG()`
>>
>> **arr** – The destination array
>>
>> **dist_type** – Distribution type
>>
>>> **– CV_RAND_UNI** uniform distribution

– **CV_RAND_NORMAL** normal or Gaussian distribution

**param1** – The first parameter of the distribution. In the case of a uniform distribution it is the inclusive lower boundary of the random numbers range. In the case of a normal distribution it is the mean value of the random numbers.

**param2** – The second parameter of the distribution. In the case of a uniform distribution it is the exclusive upper boundary of the random numbers range. In the case of a normal distribution it is the standard deviation of the random numbers.

The function fills the destination array with uniformly or normally distributed random numbers.

**See Also:**

`randu()`, `randn()`, `RNG::fill()`.

## RandInt

Returns a 32-bit unsigned integer and updates RNG.

**C:** unsigned int **cvRandInt**(CvRNG* **rng**)

**Python:** cv.**RandInt**(rng) → unsigned

> **Parameters**
>
> > **rng** – CvRNG state initialized by `RNG()`.

The function returns a uniformly-distributed random 32-bit unsigned integer and updates the RNG state. It is similar to the rand() function from the C runtime library, except that OpenCV functions always generates a 32-bit random number, regardless of the platform.

## RandReal

Returns a floating-point random number and updates RNG.

**C:** double **cvRandReal**(CvRNG* **rng**)

**Python:** cv.**RandReal**(rng) → float

> **Parameters**
>
> > **rng** – RNG state initialized by `RNG()`

The function returns a uniformly-distributed random floating-point number between 0 and 1 (1 is not included).

## fromarray

Create a CvMat from an object that supports the array interface.

**Python:** cv.**fromarray**(array, allowND=False) → mat

> **Parameters**
>
> > **object** – Any object that supports the array interface
> >
> > **allowND** – If true, will return a CvMatND

If the object supports the array interface , return a `CvMat` or `CvMatND`, depending on `allowND` flag:

- If `allowND = False`, then the object's array must be either 2D or 3D. If it is 2D, then the returned CvMat has a single channel. If it is 3D, then the returned CvMat will have N channels, where N is the last dimension of the array. In this case, N cannot be greater than OpenCV's channel limit, `CV_CN_MAX`.

- If``allowND = True``, then `fromarray` returns a single-channel `CvMatND` with the same shape as the original array.

For example, NumPy arrays support the array interface, so can be converted to OpenCV objects:

**Note:** In the new Python wrappers (**cv2** module) the function is not needed, since cv2 can process Numpy arrays (and this is the only supported array type).

## 2.3 Dynamic Structures

The section describes OpenCV 1.x API for creating growable sequences and other dynamic data structures allocated in `CvMemStorage`. If you use the new C++, Python, Java etc interface, you will unlikely need this functionality. Use `std::vector` or other high-level data structures.

### CvMemStorage

**struct `CvMemStorage`**

A storage for various OpenCV dynamic data structures, such as `CvSeq`, `CvSet` etc.

CvMemBlock* **`bottom`**
the first memory block in the double-linked list of blocks

CvMemBlock* **`top`**
the current partially allocated memory block in the list of blocks

CvMemStorage* **`parent`**
the parent storage (if any) from which the new memory blocks are borrowed.

int **`free_space`**
number of free bytes in the `top` block

int **`block_size`**
the total size of the memory blocks

Memory storage is a low-level structure used to store dynamically growing data structures such as sequences, contours, graphs, subdivisions, etc. It is organized as a list of memory blocks of equal size - `bottom` field is the beginning of the list of blocks and `top` is the currently used block, but not necessarily the last block of the list. All blocks between `bottom` and `top`, not including the latter, are considered fully occupied; all blocks between `top` and the last block, not including `top`, are considered free and `top` itself is partly occupied - `free_space` contains the number of free bytes left in the end of `top`.

A new memory buffer that may be allocated explicitly by `MemStorageAlloc()` function or implicitly by higher-level functions, such as `SeqPush()`, `GraphAddEdge()` etc.

The buffer is put in the end of already allocated space in the `top` memory block, if there is enough free space. After allocation, `free_space` is decreased by the size of the allocated buffer plus some padding to keep the proper alignment. When the allocated buffer does not fit into the available portion of `top`, the next storage block from the list is taken as `top` and `free_space` is reset to the whole block size prior to the allocation.

If there are no more free blocks, a new block is allocated (or borrowed from the parent, see `CreateChildMemStorage()`) and added to the end of list. Thus, the storage behaves as a stack with `bottom` in-

dicating bottom of the stack and the pair (`top`, `free_space`) indicating top of the stack. The stack top may be saved via SaveMemStoragePos(), restored via RestoreMemStoragePos(), or reset via ClearMemStorage().

## CvMemBlock

**struct CvMemBlock**

The structure `CvMemBlock` represents a single block of memory storage. The actual data in the memory blocks follows the header.

## CvMemStoragePos

**struct CvMemStoragePos**

The structure stores the position in the memory storage. It is used by SaveMemStoragePos() and RestoreMemStoragePos().

## CvSeq

**struct CvSeq**
>   Dynamically growing sequence.

>   int **flags**
>   >   sequence flags, including the sequence signature (CV_SEQ_MAGIC_VAL or CV_SET_MAGIC_VAL), type of the elements and some other information about the sequence.

>   int **header_size**
>   >   size of the sequence header. It should be sizeof(CvSeq) at minimum. See CreateSeq().

>   CvSeq* **h_prev**

>   CvSeq* **h_next**

>   CvSeq* **v_prev**

>   CvSeq* **v_next**
>   >   pointers to another sequences in a sequence tree. Sequence trees are used to store hierarchical contour structures, retrieved by FindContours()

>   int **total**
>   >   the number of sequence elements

>   int **elem_size**
>   >   size of each sequence element in bytes

>   CvMemStorage* **storage**
>   >   memory storage where the sequence resides. It can be a NULL pointer.

>   CvSeqBlock* **first**
>   >   pointer to the first data block

The structure `CvSeq` is a base for all of OpenCV dynamic data structures. There are two types of sequences - dense and sparse. The base type for dense sequences is CvSeq and such sequences are used to represent growable 1d arrays - vectors, stacks, queues, and deques. They have no gaps in the middle - if an element is removed from the middle or inserted into the middle of the sequence, the elements from the closer end are shifted. Sparse sequences have CvSet as a base class and they are discussed later in more detail. They are sequences of nodes; each may be either occupied or free as indicated by the node flag. Such sequences are used for unordered data structures such as sets of elements, graphs, hash tables and so forth.

## CvSlice

**struct CvSlice**

> A sequence slice. In C++ interface the class Range should be used instead.

> **int start_index**
>
>> inclusive start index of the sequence slice

> **int end_index**
>
>> exclusive end index of the sequence slice

There are helper functions to construct the slice and to compute its length:

**C:** CvSlice **cvSlice**(int **start**, int **end**)

> **Parameters**
>
>> **start** – Inclusive left boundary.
>>
>> **end** – Exclusive right boundary.

```
#define CV_WHOLE_SEQ_END_INDEX 0x3ffffff
#define CV_WHOLE_SEQ  cvSlice(0, CV_WHOLE_SEQ_END_INDEX)
```

**C:** int **cvSliceLength**(CvSlice **slice**, const CvSeq* **seq**)

> **Parameters**
>
>> **slice** – The slice of sequence.
>>
>> **seq** – Source sequence.

Calculates the sequence slice length.

Some of functions that operate on sequences take a `CvSlice slice` parameter that is often set to the whole sequence (CV_WHOLE_SEQ) by default. Either of the `start_index` and `end_index` may be negative or exceed the sequence length. If they are equal, the slice is considered empty (i.e., contains no elements). Because sequences are treated as circular structures, the slice may select a few elements in the end of a sequence followed by a few elements at the beginning of the sequence. For example, `cvSlice(-2, 3)` in the case of a 10-element sequence will select a 5-element slice, containing the pre-last (8th), last (9th), the very first (0th), second (1th) and third (2nd) elements. The functions normalize the slice argument in the following way:

1. `SliceLength()` is called to determine the length of the slice,

2. `start_index` of the slice is normalized similarly to the argument of `GetSeqElem()` (i.e., negative indices are allowed). The actual slice to process starts at the normalized `start_index` and lasts `SliceLength()` elements (again, assuming the sequence is a circular structure).

If a function does not accept a slice argument, but you want to process only a part of the sequence, the sub-sequence may be extracted using the `SeqSlice()` function, or stored into a continuous buffer with `CvtSeqToArray()` (optionally, followed by `MakeSeqHeaderForArray()`).

## CvSet

**struct CvSet**

The structure `CvSet` is a base for OpenCV 1.x sparse data structures. It is derived from CvSeq and includes an additional member `free_elems` - a list of free nodes. Every node of the set, whether free or not, is an element of the underlying sequence. While there are no restrictions on elements of dense sequences, the set (and derived structures) elements must start with an integer field and be able to fit CvSetElem structure, because these two fields (an integer followed by a pointer) are required for the organization of a node set with the list of free nodes. If a node is free, the `flags` field is negative (the most-significant bit, or MSB, of the field is set), and the `next_free` points to the next free

node (the first free node is referenced by the `free_elems` field of `CvSet`). And if a node is occupied, the `flags` field is positive and contains the node index that may be retrieved using the (`set_elem->flags & CV_SET_ELEM_IDX_MASK`) expressions, the rest of the node content is determined by the user. In particular, the occupied nodes are not linked as the free nodes are, so the second field can be used for such a link as well as for some different purpose. The macro `CV_IS_SET_ELEM(set_elem_ptr)` can be used to determined whether the specified node is occupied or not.

Initially the set and the free node list are empty. When a new node is requested from the set, it is taken from the list of free nodes, which is then updated. If the list appears to be empty, a new sequence block is allocated and all the nodes within the block are joined in the list of free nodes. Thus, the `total` field of the set is the total number of nodes both occupied and free. When an occupied node is released, it is added to the list of free nodes. The node released last will be occupied first.

`CvSet` is used to represent graphs (`CvGraph`), sparse multi-dimensional arrays (`CvSparseMat`), and planar subdivisions (`CvSubdiv2D`).

## CvGraph

**struct `CvGraph`**

The structure `CvGraph` is a base for graphs used in OpenCV 1.x. It inherits from `CvSet`, that is, it is considered as a set of vertices. Besides, it contains another set as a member, a set of graph edges. Graphs in OpenCV are represented using adjacency lists format.

## CvGraphScanner

**struct `CvGraphScanner`**

The structure `CvGraphScanner` is used for depth-first graph traversal. See discussion of the functions below.

## CvTreeNodeIterator

**struct `CvTreeNodeIterator`**

The structure `CvTreeNodeIterator` is used to traverse trees of sequences.

## ClearGraph

Clears a graph.

**C:** void **`cvClearGraph`**(CvGraph* **graph**)

> **Parameters**
>
> > **graph** – Graph

The function removes all vertices and edges from a graph. The function has O(1) time complexity.

## ClearMemStorage

Clears memory storage.

**C:** void **`cvClearMemStorage`**(CvMemStorage* **storage**)

> **Parameters**
>
> > **storage** – Memory storage

The function resets the top (free space boundary) of the storage to the very beginning. This function does not deallocate any memory. If the storage has a parent, the function returns all blocks to the parent.

## ClearSeq

Clears a sequence.

**C:** void **cvClearSeq**(CvSeq* **seq**)

> **Parameters**
>
> > **seq** – Sequence

The function removes all elements from a sequence. The function does not return the memory to the storage block, but this memory is reused later when new elements are added to the sequence. The function has 'O(1)' time complexity.

---

**Note:** It is impossible to deallocate a sequence, i.e. free space in the memory storage occupied by the sequence. Instead, call `ClearMemStorage()` or `ReleaseMemStorage()` from time to time somewhere in a top-level processing loop.

---

## ClearSet

Clears a set.

**C:** void **cvClearSet**(CvSet* **set_header**)

> **Parameters**
>
> > **set_header** – Cleared set

The function removes all elements from set. It has O(1) time complexity.

## CloneGraph

Clones a graph.

**C:** CvGraph* **cvCloneGraph**(const CvGraph* **graph**, CvMemStorage* **storage**)

> **Parameters**
>
> > **graph** – The graph to copy
> >
> > **storage** – Container for the copy

The function creates a full copy of the specified graph. If the graph vertices or edges have pointers to some external data, it can still be shared between the copies. The vertex and edge indices in the new graph may be different from the original because the function defragments the vertex and edge sets.

## CloneSeq

Creates a copy of a sequence.

**C:** CvSeq* **cvCloneSeq**(const CvSeq* **seq**, CvMemStorage* **storage**=NULL )

**Python:** cv.**CloneSeq**(seq, storage) → None

> **Parameters**

> **seq** – Sequence
>
> **storage** – The destination storage block to hold the new sequence header and the copied data, if any. If it is NULL, the function uses the storage block containing the input sequence.

The function makes a complete copy of the input sequence and returns it.

The call `cvCloneSeq( seq, storage )` is equivalent to `cvSeqSlice( seq, CV_WHOLE_SEQ, storage, 1 )`.

## CreateChildMemStorage

Creates child memory storage.

**C:** CvMemStorage* **cvCreateChildMemStorage**(CvMemStorage* **parent**)
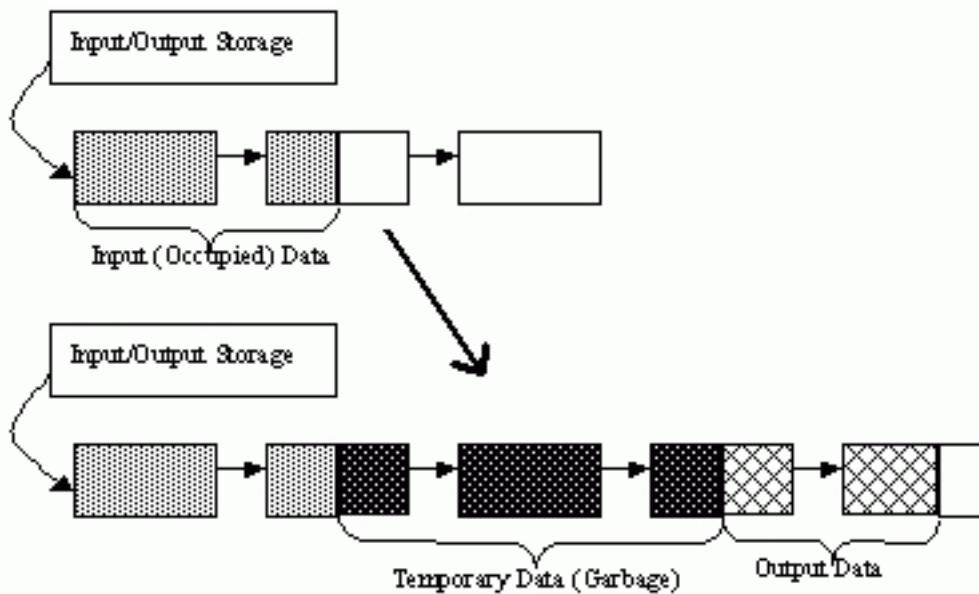
> **Parameters**
>
> > **parent** – Parent memory storage

The function creates a child memory storage that is similar to simple memory storage except for the differences in the memory allocation/deallocation mechanism. When a child storage needs a new block to add to the block list, it tries to get this block from the parent. The first unoccupied parent block available is taken and excluded from the parent block list. If no blocks are available, the parent either allocates a block or borrows one from its own parent, if any. In other words, the chain, or a more complex structure, of memory storages where every storage is a child/parent of another is possible. When a child storage is released or even cleared, it returns all blocks to the parent. In other aspects, child storage is the same as simple storage.
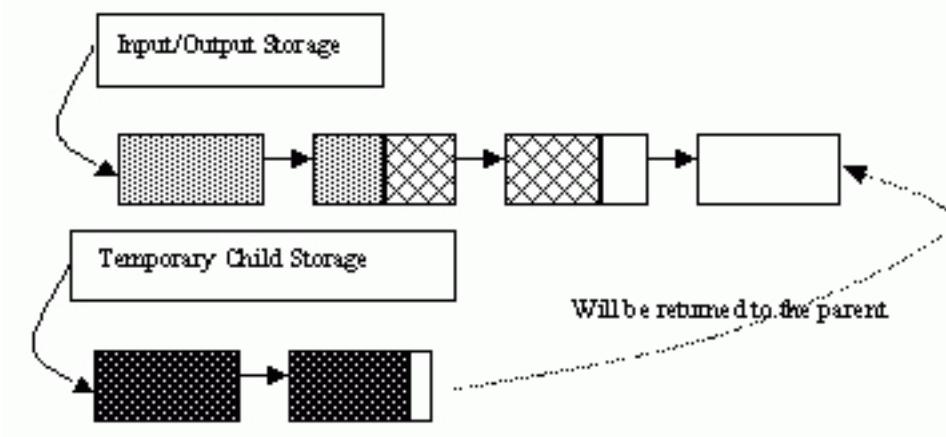
Child storage is useful in the following situation. Imagine that the user needs to process dynamic data residing in a given storage area and put the result back to that same storage area. With the simplest approach, when temporary data is resided in the same storage area as the input and output data, the storage area will look as follows after processing:

Dynamic data processing without using child storage



That is, garbage appears in the middle of the storage. However, if one creates a child memory storage at the beginning of processing, writes temporary data there, and releases the child storage at the end, no garbage will appear in the source/destination storage:

Dynamic data processing using a child storage

## CreateGraph

Creates an empty graph.

**C:** CvGraph* **cvCreateGraph**(int **graph_flags**, int **header_size**, int **vtx_size**, int **edge_size**, CvMemStorage*
**storage**)

> **Parameters**
>
> > **graph_flags** – Type of the created graph. Usually, it is either `CV_SEQ_KIND_GRAPH`
> > for generic unoriented graphs and `CV_SEQ_KIND_GRAPH | CV_GRAPH_FLAG_ORIENTED` for
> > generic oriented graphs.
> >
> > **header_size** – Graph header size; may not be less than `sizeof(CvGraph)`
> >
> > **vtx_size** – Graph vertex size; the custom vertex structure must start with `CvGraphVtx` (use
> > `CV_GRAPH_VERTEX_FIELDS()` )
> >
> > **edge_size** – Graph edge size; the custom edge structure must start with `CvGraphEdge` (use
> > `CV_GRAPH_EDGE_FIELDS()` )
> >
> > **storage** – The graph container

The function creates an empty graph and returns a pointer to it.

## CreateGraphScanner

Creates structure for depth-first graph traversal.

**C:** CvGraphScanner* **cvCreateGraphScanner**(CvGraph* **graph**, CvGraphVtx* **vtx**=NULL, int
**mask**=CV_GRAPH_ALL_ITEMS )

> **Parameters**
>
> > **graph** – Graph
> >
> > **vtx** – Initial vertex to start from. If NULL, the traversal starts from the first vertex (a vertex
> > with the minimal index in the sequence of vertices).
> >
> > **mask** – Event mask indicating which events are of interest to the user (where
> > `NextGraphItem()` function returns control to the user) It can be `CV_GRAPH_ALL_ITEMS`
> > (all events are of interest) or a combination of the following flags:
> >
> > > – **CV_GRAPH_VERTEX** stop at the graph vertices visited for the first time

- **CV_GRAPH_TREE_EDGE** stop at tree edges ( `tree edge` is the edge connecting the last visited vertex and the vertex to be visited next)

- **CV_GRAPH_BACK_EDGE** stop at back edges ( `back edge` is an edge connecting the last visited vertex with some of its ancestors in the search tree)

- **CV_GRAPH_FORWARD_EDGE** stop at forward edges ( `forward edge` is an edge connecting the last visited vertex with some of its descendants in the search tree. The forward edges are only possible during oriented graph traversal)

- **CV_GRAPH_CROSS_EDGE** stop at cross edges ( `cross edge` is an edge connecting different search trees or branches of the same tree. The `cross edges` are only possible during oriented graph traversal)

- **CV_GRAPH_ANY_EDGE** stop at any edge ( `tree`, `back`, `forward` , and `cross edges` )

- **CV_GRAPH_NEW_TREE** stop in the beginning of every new search tree. When the traversal procedure visits all vertices and edges reachable from the initial vertex (the visited vertices together with tree edges make up a tree), it searches for some unvisited vertex in the graph and resumes the traversal process from that vertex. Before starting a new tree (including the very first tree when `cvNextGraphItem` is called for the first time) it generates a `CV_GRAPH_NEW_TREE` event. For unoriented graphs, each search tree corresponds to a connected component of the graph.

- **CV_GRAPH_BACKTRACKING** stop at every already visited vertex during backtracking - returning to already visited vertexes of the traversal tree.

The function creates a structure for depth-first graph traversal/search. The initialized structure is used in the NextGraphItem() function - the incremental traversal procedure.

## CreateMemStorage

Creates memory storage.

**C:** CvMemStorage* **cvCreateMemStorage**(int **block_size**=0 )

**Python:** cv.**CreateMemStorage**(blockSize=0) → memstorage

> **Parameters**
>
> > **block_size** – Size of the storage blocks in bytes. If it is 0, the block size is set to a default value - currently it is about 64K.

The function creates an empty memory storage. See CvMemStorage description.

## CreateSeq

Creates a sequence.

**C:** CvSeq* **cvCreateSeq**(int **seq_flags**, size_t **header_size**, size_t **elem_size**, CvMemStorage* **storage**)

> **Parameters**
>
> > **seq_flags** – Flags of the created sequence. If the sequence is not passed to any function working with a specific type of sequences, the sequence value may be set to 0, otherwise the appropriate type must be selected from the list of predefined sequence types.
> >
> > **header_size** – Size of the sequence header; must be greater than or equal to `sizeof(CvSeq)` . If a specific type or its extension is indicated, this type must fit the base type header.

> **elem_size** – Size of the sequence elements in bytes. The size must be consistent with the sequence type. For example, for a sequence of points to be created, the element type `CV_SEQ_ELTYPE_POINT` should be specified and the parameter `elem_size` must be equal to `sizeof(CvPoint)`.
>
> **storage** – Sequence location

The function creates a sequence and returns the pointer to it. The function allocates the sequence header in the storage block as one continuous chunk and sets the structure fields `flags`, `elemSize`, `headerSize`, and `storage` to passed values, sets `delta_elems` to the default value (that may be reassigned using the `SetSeqBlockSize()` function), and clears other header fields, including the space following the first `sizeof(CvSeq)` bytes.

## CreateSet

Creates an empty set.

**C:** CvSet* **cvCreateSet**(int **set_flags**, int **header_size**, int **elem_size**, CvMemStorage* **storage**)

> **Parameters**
>
> > **set_flags** – Type of the created set
> >
> > **header_size** – Set header size; may not be less than `sizeof(CvSet)`
> >
> > **elem_size** – Set element size; may not be less than `CvSetElem`
> >
> > **storage** – Container for the set

The function creates an empty set with a specified header size and element size, and returns the pointer to the set. This function is just a thin layer on top of `CreateSeq()`.

## CvtSeqToArray

Copies a sequence to one continuous block of memory.

**C:** void* **cvCvtSeqToArray**(const CvSeq* **seq**, void* **elements**, CvSlice **slice**=CV_WHOLE_SEQ )

> **Parameters**
>
> > **seq** – Sequence
> >
> > **elements** – Pointer to the destination array that must be large enough. It should be a pointer to data, not a matrix header.
> >
> > **slice** – The sequence portion to copy to the array

The function copies the entire sequence or subsequence to the specified buffer and returns the pointer to the buffer.

## EndWriteSeq

Finishes the process of writing a sequence.

**C:** CvSeq* **cvEndWriteSeq**(CvSeqWriter* **writer**)

> **Parameters**
>
> > **writer** – Writer state

The function finishes the writing process and returns the pointer to the written sequence. The function also truncates the last incomplete sequence block to return the remaining part of the block to memory storage. After that, the sequence can be read and modified safely. See `StartWriteSeq()` and `StartAppendToSeq()`

---

## FindGraphEdge

Finds an edge in a graph.

**C:** CvGraphEdge* **cvFindGraphEdge**(const CvGraph* **graph**, int **start_idx**, int **end_idx**)

> **Parameters**
>
>> **graph** – Graph
>>
>> **start_idx** – Index of the starting vertex of the edge
>>
>> **end_idx** – Index of the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

```
#define cvGraphFindEdge cvFindGraphEdge
```

The function finds the graph edge connecting two specified vertices and returns a pointer to it or NULL if the edge does not exist.

## FindGraphEdgeByPtr

Finds an edge in a graph by using its pointer.

**C:** CvGraphEdge* **cvFindGraphEdgeByPtr**(const CvGraph* **graph**, const CvGraphVtx* **start_vtx**, const CvGraphVtx* **end_vtx**)

> **Parameters**
>
>> **graph** – Graph
>>
>> **start_vtx** – Pointer to the starting vertex of the edge
>>
>> **end_vtx** – Pointer to the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

```
#define cvGraphFindEdgeByPtr cvFindGraphEdgeByPtr
```

The function finds the graph edge connecting two specified vertices and returns pointer to it or NULL if the edge does not exists.

## FlushSeqWriter

Updates sequence headers from the writer.

**C:** void **cvFlushSeqWriter**(CvSeqWriter* **writer**)

> **Parameters**
>
>> **writer** – Writer state

The function is intended to enable the user to read sequence elements, whenever required, during the writing process, e.g., in order to check specific conditions. The function updates the sequence headers to make reading from the sequence possible. The writer is not closed, however, so that the writing process can be continued at any time. If an algorithm requires frequent flushes, consider using `SeqPush()` instead.

## GetGraphVtx

Finds a graph vertex by using its index.

**C:** CvGraphVtx* **cvGetGraphVtx**(CvGraph* **graph**, int **vtx_idx**)

>> **Parameters**

>>> **graph** – Graph

>>> **vtx_idx** – Index of the vertex

The function finds the graph vertex by using its index and returns the pointer to it or NULL if the vertex does not belong to the graph.

## GetSeqElem

Returns a pointer to a sequence element according to its index.

**C:** schar* **cvGetSeqElem**(const CvSeq* **seq**, int **index**)

>> **Parameters**

>>> **seq** – Sequence

>>> **index** – Index of element

```
#define CV_GET_SEQ_ELEM( TYPE, seq, index )  (TYPE*)cvGetSeqElem( (CvSeq*)(seq), (index) )
```

The function finds the element with the given index in the sequence and returns the pointer to it. If the element is not found, the function returns 0. The function supports negative indices, where -1 stands for the last sequence element, -2 stands for the one before last, etc. If the sequence is most likely to consist of a single sequence block or the desired element is likely to be located in the first block, then the macro `CV_GET_SEQ_ELEM( elemType, seq, index )` should be used, where the parameter `elemType` is the type of sequence elements ( `CvPoint` for example), the parameter `seq` is a sequence, and the parameter `index` is the index of the desired element. The macro checks first whether the desired element belongs to the first block of the sequence and returns it if it does; otherwise the macro calls the main function GetSeqElem . Negative indices always cause the `GetSeqElem()` call. The function has O(1) time complexity assuming that the number of blocks is much smaller than the number of elements.

## GetSeqReaderPos

Returns the current reader position.

**C:** int **cvGetSeqReaderPos**(CvSeqReader* **reader**)

>> **Parameters**

>>> **reader** – Reader state

The function returns the current reader position (within 0 ... `reader->seq->total` - 1).

## GetSetElem

Finds a set element by its index.

**C:** CvSetElem* **cvGetSetElem**(const CvSet* **set_header**, int **idx**)

>> **Parameters**

>>> **set_header** – Set

**idx** – Index of the set element within a sequence

The function finds a set element by its index. The function returns the pointer to it or 0 if the index is invalid or the corresponding node is free. The function supports negative indices as it uses `GetSeqElem()` to locate the node.

## GraphAddEdge

Adds an edge to a graph.

**C:** int **cvGraphAddEdge**(CvGraph* **graph**, int **start_idx**, int **end_idx**, const CvGraphEdge* **edge**=NULL, CvGraphEdge** **inserted_edge**=NULL )

      **Parameters**

            **graph** – Graph

            **start_idx** – Index of the starting vertex of the edge

            **end_idx** – Index of the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

            **edge** – Optional input parameter, initialization data for the edge

            **inserted_edge** – Optional output parameter to contain the address of the inserted edge

The function connects two specified vertices. The function returns 1 if the edge has been added successfully, 0 if the edge connecting the two vertices exists already and -1 if either of the vertices was not found, the starting and the ending vertex are the same, or there is some other critical situation. In the latter case (i.e., when the result is negative), the function also reports an error by default.

## GraphAddEdgeByPtr

Adds an edge to a graph by using its pointer.

**C:** int **cvGraphAddEdgeByPtr**(CvGraph* **graph**, CvGraphVtx* **start_vtx**, CvGraphVtx* **end_vtx**, const CvGraphEdge* **edge**=NULL, CvGraphEdge** **inserted_edge**=NULL )

      **Parameters**

            **graph** – Graph

            **start_vtx** – Pointer to the starting vertex of the edge

            **end_vtx** – Pointer to the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

            **edge** – Optional input parameter, initialization data for the edge

            **inserted_edge** – Optional output parameter to contain the address of the inserted edge within the edge set

The function connects two specified vertices. The function returns 1 if the edge has been added successfully, 0 if the edge connecting the two vertices exists already, and -1 if either of the vertices was not found, the starting and the ending vertex are the same or there is some other critical situation. In the latter case (i.e., when the result is negative), the function also reports an error by default.

## GraphAddVtx

Adds a vertex to a graph.

**C:** int **cvGraphAddVtx**(CvGraph* **graph**, const CvGraphVtx* **vtx**=NULL, CvGraphVtx** **inserted_vtx**=NULL )

> Parameters

>> **graph** – Graph

>> **vtx** – Optional input argument used to initialize the added vertex (only user-defined fields beyond `sizeof(CvGraphVtx)` are copied)

>> **inserted_vtx** – Optional output argument. If not `NULL` , the address of the new vertex is written here.

The function adds a vertex to the graph and returns the vertex index.

## GraphEdgeIdx

Returns the index of a graph edge.

**C:** int **cvGraphEdgeIdx**(CvGraph* **graph**, CvGraphEdge* **edge**)

> Parameters

>> **graph** – Graph

>> **edge** – Pointer to the graph edge

The function returns the index of a graph edge.

## GraphRemoveEdge

Removes an edge from a graph.

**C:** void **cvGraphRemoveEdge**(CvGraph* **graph**, int **start_idx**, int **end_idx**)

> Parameters

>> **graph** – Graph

>> **start_idx** – Index of the starting vertex of the edge

>> **end_idx** – Index of the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

The function removes the edge connecting two specified vertices. If the vertices are not connected [in that order], the function does nothing.

## GraphRemoveEdgeByPtr

Removes an edge from a graph by using its pointer.

**C:** void **cvGraphRemoveEdgeByPtr**(CvGraph* **graph**, CvGraphVtx* **start_vtx**, CvGraphVtx* **end_vtx**)

> Parameters

>> **graph** – Graph

>> **start_vtx** – Pointer to the starting vertex of the edge

>> **end_vtx** – Pointer to the ending vertex of the edge. For an unoriented graph, the order of the vertex parameters does not matter.

The function removes the edge connecting two specified vertices. If the vertices are not connected [in that order], the function does nothing.

## GraphRemoveVtx

Removes a vertex from a graph.

**C:** int **cvGraphRemoveVtx**(CvGraph* **graph**, int **index**)

> **Parameters**
>
> > **graph** – Graph
> >
> > **index** – Index of the removed vertex

The function removes a vertex from a graph together with all the edges incident to it. The function reports an error if the input vertex does not belong to the graph. The return value is the number of edges deleted, or -1 if the vertex does not belong to the graph.

## GraphRemoveVtxByPtr

Removes a vertex from a graph by using its pointer.

**C:** int **cvGraphRemoveVtxByPtr**(CvGraph* **graph**, CvGraphVtx* **vtx**)

> **Parameters**
>
> > **graph** – Graph
> >
> > **vtx** – Pointer to the removed vertex

The function removes a vertex from the graph by using its pointer together with all the edges incident to it. The function reports an error if the vertex does not belong to the graph. The return value is the number of edges deleted, or -1 if the vertex does not belong to the graph.

## GraphVtxDegree

Counts the number of edges incident to the vertex.

**C:** int **cvGraphVtxDegree**(const CvGraph* **graph**, int **vtx_idx**)

> **Parameters**
>
> > **graph** – Graph
> >
> > **vtx_idx** – Index of the graph vertex

The function returns the number of edges incident to the specified vertex, both incoming and outgoing. To count the edges, the following code is used:

```
CvGraphEdge* edge = vertex->first; int count = 0;
while( edge )
{
    edge = CV_NEXT_GRAPH_EDGE( edge, vertex );
    count++;
}
```

The macro CV_NEXT_GRAPH_EDGE( edge, vertex ) returns the edge incident to vertex that follows after edge .

## GraphVtxDegreeByPtr

Finds an edge in a graph.

**C:** int **cvGraphVtxDegreeByPtr**(const CvGraph* **graph**, const CvGraphVtx* **vtx**)

> **Parameters**
>
> > **graph** – Graph
> >
> > **vtx** – Pointer to the graph vertex

The function returns the number of edges incident to the specified vertex, both incoming and outcoming.

## GraphVtxIdx

Returns the index of a graph vertex.

**C:** int **cvGraphVtxIdx**(CvGraph* **graph**, CvGraphVtx* **vtx**)

> **Parameters**
>
> > **graph** – Graph
> >
> > **vtx** – Pointer to the graph vertex

The function returns the index of a graph vertex.

## InitTreeNodeIterator

Initializes the tree node iterator.

**C:** void **cvInitTreeNodeIterator**(CvTreeNodeIterator* **tree_iterator**, const void* **first**, int **max_level**)

> **Parameters**
>
> > **tree_iterator** – Tree iterator initialized by the function
> >
> > **first** – The initial node to start traversing from
> >
> > **max_level** – The maximal level of the tree ( `first` node assumed to be at the first level) to traverse up to. For example, 1 means that only nodes at the same level as `first` should be visited, 2 means that the nodes on the same level as `first` and their direct children should be visited, and so forth.

The function initializes the tree iterator. The tree is traversed in depth-first order.

## InsertNodeIntoTree

Adds a new node to a tree.

**C:** void **cvInsertNodeIntoTree**(void* **node**, void* **parent**, void* **frame**)

> **Parameters**
>
> > **node** – The inserted node
> >
> > **parent** – The parent node that is already in the tree
> >
> > **frame** – The top level node. If `parent` and `frame` are the same, the v_prev field of `node` is set to NULL rather than `parent` .

The function adds another node into tree. The function does not allocate any memory, it can only modify links of the tree nodes.

## MakeSeqHeaderForArray

Constructs a sequence header for an array.

**C:** CvSeq* **cvMakeSeqHeaderForArray**(int **seq_type**, int **header_size**, int **elem_size**, void* **elements**, int **total**, CvSeq* **seq**, CvSeqBlock* **block**)

> **Parameters**
>
>> **seq_type** – Type of the created sequence
>>
>> **header_size** – Size of the header of the sequence. Parameter sequence must point to the structure of that size or greater
>>
>> **elem_size** – Size of the sequence elements
>>
>> **elements** – Elements that will form a sequence
>>
>> **total** – Total number of elements in the sequence. The number of array elements must be equal to the value of this parameter.
>>
>> **seq** – Pointer to the local variable that is used as the sequence header
>>
>> **block** – Pointer to the local variable that is the header of the single sequence block

The function initializes a sequence header for an array. The sequence header as well as the sequence block are allocated by the user (for example, on stack). No data is copied by the function. The resultant sequence will consists of a single block and have NULL storage pointer; thus, it is possible to read its elements, but the attempts to add elements to the sequence will raise an error in most cases.

## MemStorageAlloc

Allocates a memory buffer in a storage block.

**C:** void* **cvMemStorageAlloc**(CvMemStorage* **storage**, size_t **size**)

> **Parameters**
>
>> **storage** – Memory storage
>>
>> **size** – Buffer size

The function allocates a memory buffer in a storage block. The buffer size must not exceed the storage block size, otherwise a runtime error is raised. The buffer address is aligned by `CV_STRUCT_ALIGN=sizeof(double)` (for the moment) bytes.

## MemStorageAllocString

Allocates a text string in a storage block.

**C:** CvString **cvMemStorageAllocString**(CvMemStorage* **storage**, const char* **ptr**, int **len**=-1)

> **Parameters**
>
>> **storage** – Memory storage
>>
>> **ptr** – The string

> **len** – Length of the string (not counting the ending NUL ) . If the parameter is negative, the
> function computes the length.

```
typedef struct CvString
{
    int len;
    char* ptr;
}
CvString;
```

The function creates copy of the string in memory storage. It returns the structure that contains user-passed or computed length of the string and pointer to the copied string.

## NextGraphItem

Executes one or more steps of the graph traversal procedure.

**C:** int **cvNextGraphItem**(CvGraphScanner* **scanner**)

> **Parameters**
>
> > **scanner** – Graph traversal state. It is updated by this function.

The function traverses through the graph until an event of interest to the user (that is, an event, specified in the mask in the CreateGraphScanner() call) is met or the traversal is completed. In the first case, it returns one of the events listed in the description of the mask parameter above and with the next call it resumes the traversal. In the latter case, it returns CV_GRAPH_OVER (-1). When the event is CV_GRAPH_VERTEX , CV_GRAPH_BACKTRACKING , or CV_GRAPH_NEW_TREE , the currently observed vertex is stored in scanner-:math:'>'vtx . And if the event is edge-related, the edge itself is stored at scanner-:math:'>'edge , the previously visited vertex - at scanner-:math:'>'vtx and the other ending vertex of the edge - at scanner-:math:'>'dst .

## NextTreeNode

Returns the currently observed node and moves the iterator toward the next node.

**C:** void* **cvNextTreeNode**(CvTreeNodeIterator* **tree_iterator**)

> **Parameters**
>
> > **tree_iterator** – Tree iterator initialized by the function

The function returns the currently observed node and then updates the iterator - moving it toward the next node. In other words, the function behavior is similar to the *p++ expression on a typical C pointer or C++ collection iterator. The function returns NULL if there are no more nodes.

## PrevTreeNode

Returns the currently observed node and moves the iterator toward the previous node.

**C:** void* **cvPrevTreeNode**(CvTreeNodeIterator* **tree_iterator**)

> **Parameters**
>
> > **tree_iterator** – Tree iterator initialized by the function

The function returns the currently observed node and then updates the iterator - moving it toward the previous node. In other words, the function behavior is similar to the *p-- expression on a typical C pointer or C++ collection iterator. The function returns NULL if there are no more nodes.

## ReleaseGraphScanner

Completes the graph traversal procedure.

**C:** void **cvReleaseGraphScanner**(CvGraphScanner** **scanner**)

> **Parameters**
>
> > **scanner** – Double pointer to graph traverser

The function completes the graph traversal procedure and releases the traverser state.

## ReleaseMemStorage

Releases memory storage.

**C:** void **cvReleaseMemStorage**(CvMemStorage** **storage**)

> **Parameters**
>
> > **storage** – Pointer to the released storage

The function deallocates all storage memory blocks or returns them to the parent, if any. Then it deallocates the storage header and clears the pointer to the storage. All child storage associated with a given parent storage block must be released before the parent storage block is released.

## RestoreMemStoragePos

Restores memory storage position.

**C:** void **cvRestoreMemStoragePos**(CvMemStorage* **storage**, CvMemStoragePos* **pos**)

> **Parameters**
>
> > **storage** – Memory storage
> >
> > **pos** – New storage top position

The function restores the position of the storage top from the parameter `pos` . This function and the function `cvClearMemStorage` are the only methods to release memory occupied in memory blocks. Note again that there is no way to free memory in the middle of an occupied portion of a storage block.

## SaveMemStoragePos

Saves memory storage position.

**C:** void **cvSaveMemStoragePos**(const CvMemStorage* **storage**, CvMemStoragePos* **pos**)

> **Parameters**
>
> > **storage** – Memory storage
> >
> > **pos** – The output position of the storage top

The function saves the current position of the storage top to the parameter `pos` . The function `cvRestoreMemStoragePos` can further retrieve this position.

## SeqElemIdx

Returns the index of a specific sequence element.

**C:** int **cvSeqElemIdx**(const CvSeq* **seq**, const void* **element**, CvSeqBlock** **block**=NULL )

> **Parameters**
>
>> **seq** – Sequence
>>
>> **element** – Pointer to the element within the sequence
>>
>> **block** – Optional argument. If the pointer is not NULL , the address of the sequence block that contains the element is stored in this location.

The function returns the index of a sequence element or a negative number if the element is not found.

## SeqInsert

Inserts an element in the middle of a sequence.

**C:** schar* **cvSeqInsert**(CvSeq* **seq**, int **before_index**, const void* **element**=NULL )

> **Parameters**
>
>> **seq** – Sequence
>>
>> **before_index** – Index before which the element is inserted. Inserting before 0 (the minimal allowed value of the parameter) is equal to SeqPushFront() and inserting before seq->total (the maximal allowed value of the parameter) is equal to SeqPush() .
>>
>> **element** – Inserted element

The function shifts the sequence elements from the inserted position to the nearest end of the sequence and copies the element content there if the pointer is not NULL. The function returns a pointer to the inserted element.

## SeqInsertSlice

Inserts an array in the middle of a sequence.

**C:** void **cvSeqInsertSlice**(CvSeq* **seq**, int **before_index**, const CvArr* **from_arr**)

> **Parameters**
>
>> **seq** – Sequence
>>
>> **before_index** – Index before which the array is inserted
>>
>> **from_arr** – The array to take elements from

The function inserts all fromArr array elements at the specified position of the sequence. The array fromArr can be a matrix or another sequence.

## SeqInvert

Reverses the order of sequence elements.

**C:** void **cvSeqInvert**(CvSeq* **seq**)

> **Parameters**
>
>> **seq** – Sequence

The function reverses the sequence in-place - the first element becomes the last one, the last element becomes the first one and so forth.

## SeqPop

Removes an element from the end of a sequence.

**C:** void **cvSeqPop**(CvSeq* **seq**, void* **element**=NULL )

      **Parameters**

            **seq** – Sequence

            **element** – Optional parameter . If the pointer is not zero, the function copies the removed element to this location.

The function removes an element from a sequence. The function reports an error if the sequence is already empty. The function has O(1) complexity.

## SeqPopFront

Removes an element from the beginning of a sequence.

**C:** void **cvSeqPopFront**(CvSeq* **seq**, void* **element**=NULL )

      **Parameters**

            **seq** – Sequence

            **element** – Optional parameter. If the pointer is not zero, the function copies the removed element to this location.

The function removes an element from the beginning of a sequence. The function reports an error if the sequence is already empty. The function has O(1) complexity.

## SeqPopMulti

Removes several elements from either end of a sequence.

**C:** void **cvSeqPopMulti**(CvSeq* **seq**, void* **elements**, int **count**, int **in_front**=0 )

      **Parameters**

            **seq** – Sequence

            **elements** – Removed elements

            **count** – Number of elements to pop

            **in_front** – The flags specifying which end of the modified sequence.

              **– CV_BACK** the elements are added to the end of the sequence

              **– CV_FRONT** the elements are added to the beginning of the sequence

The function removes several elements from either end of the sequence. If the number of the elements to be removed exceeds the total number of elements in the sequence, the function removes as many elements as possible.

## SeqPush

Adds an element to the end of a sequence.

**C:** schar* **cvSeqPush**(CvSeq* **seq**, const void* **element**=NULL )

> **Parameters**
>
> > **seq** – Sequence
> >
> > **element** – Added element

The function adds an element to the end of a sequence and returns a pointer to the allocated element. If the input element is NULL, the function simply allocates a space for one more element.

The following code demonstrates how to create a new sequence using this function:

```
CvMemStorage* storage = cvCreateMemStorage(0);
CvSeq* seq = cvCreateSeq( CV_32SC1, /* sequence of integer elements */
                          sizeof(CvSeq), /* header size - no extra fields */
                          sizeof(int), /* element size */
                          storage /* the container storage */ );
int i;
for( i = 0; i < 100; i++ )
{
    int* added = (int*)cvSeqPush( seq, &i );
    printf( "
}

...
/* release memory storage in the end */
cvReleaseMemStorage( &storage );
```

The function has O(1) complexity, but there is a faster method for writing large sequences (see StartWriteSeq() and related functions).

## SeqPushFront

Adds an element to the beginning of a sequence.

**C:** schar* **cvSeqPushFront**(CvSeq* **seq**, const void* **element**=NULL )

> **Parameters**
>
> > **seq** – Sequence
> >
> > **element** – Added element

The function is similar to SeqPush() but it adds the new element to the beginning of the sequence. The function has O(1) complexity.

## SeqPushMulti

Pushes several elements to either end of a sequence.

**C:** void **cvSeqPushMulti**(CvSeq* **seq**, const void* **elements**, int **count**, int **in_front**=0 )

> **Parameters**
>
> > **seq** – Sequence

> **elements** – Added elements
>
> **count** – Number of elements to push
>
> **in_front** – The flags specifying which end of the modified sequence.
>
> > – **CV_BACK** the elements are added to the end of the sequence
> >
> > – **CV_FRONT** the elements are added to the beginning of the sequence

The function adds several elements to either end of a sequence. The elements are added to the sequence in the same order as they are arranged in the input array but they can fall into different sequence blocks.

## SeqRemove

Removes an element from the middle of a sequence.

**C:** void **cvSeqRemove**(CvSeq* **seq**, int **index**)

> **Parameters**
>
> > **seq** – Sequence
> >
> > **index** – Index of removed element

The function removes elements with the given index. If the index is out of range the function reports an error. An attempt to remove an element from an empty sequence is a special case of this situation. The function removes an element by shifting the sequence elements between the nearest end of the sequence and the index -th position, not counting the latter.

## SeqRemoveSlice

Removes a sequence slice.

**C:** void **cvSeqRemoveSlice**(CvSeq* **seq**, CvSlice **slice**)

> **Parameters**
>
> > **seq** – Sequence
> >
> > **slice** – The part of the sequence to remove

The function removes a slice from the sequence.

## SeqSearch

Searches for an element in a sequence.

**C:** schar* **cvSeqSearch**(CvSeq* **seq**, const void* **elem**, CvCmpFunc **func**, int **is_sorted**, int* **elem_idx**, void* **userdata**=NULL )

> **Parameters**
>
> > **seq** – The sequence
> >
> > **elem** – The element to look for
> >
> > **func** – The comparison function that returns negative, zero or positive value depending on the relationships among the elements (see also SeqSort() )
> >
> > **is_sorted** – Whether the sequence is sorted or not
> >
> > **elem_idx** – Output parameter; index of the found element

> > **userdata** – The user parameter passed to the comparison function; helps to avoid global variables in some cases

```
/* a < b ? -1 : a > b ? 1 : 0 */
typedef int (CV_CDECL* CvCmpFunc)(const void* a, const void* b, void* userdata);
```

The function searches for the element in the sequence. If the sequence is sorted, a binary O(log(N)) search is used; otherwise, a simple linear search is used. If the element is not found, the function returns a NULL pointer and the index is set to the number of sequence elements if a linear search is used, or to the smallest index i, `seq(i)>elem`.

## SeqSlice

Makes a separate header for a sequence slice.

**C:** CvSeq* **cvSeqSlice**(const CvSeq* **seq**, CvSlice **slice**, CvMemStorage* **storage**=NULL, int **copy_data**=0
)

> **Parameters**

> > **seq** – Sequence

> > **slice** – The part of the sequence to be extracted

> > **storage** – The destination storage block to hold the new sequence header and the copied data, if any. If it is NULL, the function uses the storage block containing the input sequence.

> > **copy_data** – The flag that indicates whether to copy the elements of the extracted slice ( `copy_data!=0` ) or not ( `copy_data=0` )

The function creates a sequence that represents the specified slice of the input sequence. The new sequence either shares the elements with the original sequence or has its own copy of the elements. So if one needs to process a part of sequence but the processing function does not have a slice parameter, the required sub-sequence may be extracted using this function.

## SeqSort

Sorts sequence element using the specified comparison function.

**C:** void **cvSeqSort**(CvSeq* **seq**, CvCmpFunc **func**, void* **userdata**=NULL )

> **Parameters**

> > **seq** – The sequence to sort

> > **func** – The comparison function that returns a negative, zero, or positive value depending on the relationships among the elements (see the above declaration and the example below) - a similar function is used by `qsort` from C runline except that in the latter, `userdata` is not used

> > **userdata** – The user parameter passed to the comparison function; helps to avoid global variables in some cases

```
/* a < b ? -1 : a > b ? 1 : 0 */
typedef int (CV_CDECL* CvCmpFunc)(const void* a, const void* b, void* userdata);
```

The function sorts the sequence in-place using the specified criteria. Below is an example of using this function:

```
/* Sort 2d points in top-to-bottom left-to-right order */
static int cmp_func( const void* _a, const void* _b, void* userdata )
{
```

```
    CvPoint* a = (CvPoint*)_a;
    CvPoint* b = (CvPoint*)_b;
    int y_diff = a->y - b->y;
    int x_diff = a->x - b->x;
    return y_diff ? y_diff : x_diff;
}

...

CvMemStorage* storage = cvCreateMemStorage(0);
CvSeq* seq = cvCreateSeq( CV_32SC2, sizeof(CvSeq), sizeof(CvPoint), storage );
int i;

for( i = 0; i < 10; i++ )
{
    CvPoint pt;
    pt.x = rand()
    pt.y = rand()
    cvSeqPush( seq, &pt );
}

cvSeqSort( seq, cmp_func, 0 /* userdata is not used here */ );

/* print out the sorted sequence */
for( i = 0; i < seq->total; i++ )
{
    CvPoint* pt = (CvPoint*)cvSeqElem( seq, i );
    printf( "(
}

cvReleaseMemStorage( &storage );
```

## SetAdd

Occupies a node in the set.

**C:** int **cvSetAdd**(CvSet* **set_header**, CvSetElem* **elem**=NULL, CvSetElem** **inserted_elem**=NULL )

>>> **Parameters**

>>>> **set_header** – Set

>>>> **elem** – Optional input argument, an inserted element. If not NULL, the function copies the data to the allocated node (the MSB of the first integer field is cleared after copying).

>>>> **inserted_elem** – Optional output argument; the pointer to the allocated cell

The function allocates a new node, optionally copies input element data to it, and returns the pointer and the index to the node. The index value is taken from the lower bits of the `flags` field of the node. The function has O(1) complexity; however, there exists a faster function for allocating set nodes (see `SetNew()` ).

## SetNew

Adds an element to a set (fast variant).

**C:** CvSetElem* **cvSetNew**(CvSet* **set_header**)

>>> **Parameters**

**set_header** – Set

The function is an inline lightweight variant of `SetAdd()` . It occupies a new node and returns a pointer to it rather than an index.

## SetRemove

Removes an element from a set.

**C:** void **cvSetRemove**(CvSet* **set_header**, int **index**)

>   **Parameters**

>>   **set_header** – Set

>>   **index** – Index of the removed element

The function removes an element with a specified index from the set. If the node at the specified location is not occupied, the function does nothing. The function has O(1) complexity; however, `SetRemoveByPtr()` provides a quicker way to remove a set element if it is located already.

## SetRemoveByPtr

Removes a set element based on its pointer.

**C:** void **cvSetRemoveByPtr**(CvSet* **set_header**, void* **elem**)

>   **Parameters**

>>   **set_header** – Set

>>   **elem** – Removed element

The function is an inline lightweight variant of `SetRemove()` that requires an element pointer. The function does not check whether the node is occupied or not - the user should take care of that.

## SetSeqBlockSize

Sets up sequence block size.

**C:** void **cvSetSeqBlockSize**(CvSeq* **seq**, int **delta_elems**)

>   **Parameters**

>>   **seq** – Sequence

>>   **delta_elems** – Desirable sequence block size for elements

The function affects memory allocation granularity. When the free space in the sequence buffers has run out, the function allocates the space for `delta_elems` sequence elements. If this block immediately follows the one previously allocated, the two blocks are concatenated; otherwise, a new sequence block is created. Therefore, the bigger the parameter is, the lower the possible sequence fragmentation, but the more space in the storage block is wasted. When the sequence is created, the parameter `delta_elems` is set to the default value of about 1K. The function can be called any time after the sequence is created and affects future allocations. The function can modify the passed value of the parameter to meet memory storage constraints.

## SetSeqReaderPos

Moves the reader to the specified position.

**C:** void **cvSetSeqReaderPos**(CvSeqReader* **reader**, int **index**, int **is_relative**=0 )

> **Parameters**
>
> > **reader** – Reader state
> >
> > **index** – The destination position. If the positioning mode is used (see the next parameter), the actual position will be index mod reader->seq->total .
> >
> > **is_relative** – If it is not zero, then index is a relative to the current position

The function moves the read position to an absolute position or relative to the current position.

## StartAppendToSeq

Initializes the process of writing data to a sequence.

**C:** void **cvStartAppendToSeq**(CvSeq* **seq**, CvSeqWriter* **writer**)

> **Parameters**
>
> > **seq** – Pointer to the sequence
> >
> > **writer** – Writer state; initialized by the function

The function initializes the process of writing data to a sequence. Written elements are added to the end of the sequence by using the CV_WRITE_SEQ_ELEM( written_elem, writer ) macro. Note that during the writing process, other operations on the sequence may yield an incorrect result or even corrupt the sequence (see description of FlushSeqWriter() , which helps to avoid some of these problems).

## StartReadSeq

Initializes the process of sequential reading from a sequence.

**C:** void **cvStartReadSeq**(const CvSeq* **seq**, CvSeqReader* **reader**, int **reverse**=0 )

> **Parameters**
>
> > **seq** – Sequence
> >
> > **reader** – Reader state; initialized by the function
> >
> > **reverse** – Determines the direction of the sequence traversal. If reverse is 0, the reader is positioned at the first sequence element; otherwise it is positioned at the last element.

The function initializes the reader state. After that, all the sequence elements from the first one down to the last one can be read by subsequent calls of the macro CV_READ_SEQ_ELEM( read_elem, reader ) in the case of forward reading and by using CV_REV_READ_SEQ_ELEM( read_elem, reader ) in the case of reverse reading. Both macros put the sequence element to read_elem and move the reading pointer toward the next element. A circular structure of sequence blocks is used for the reading process, that is, after the last element has been read by the macro CV_READ_SEQ_ELEM , the first element is read when the macro is called again. The same applies to CV_REV_READ_SEQ_ELEM . There is no function to finish the reading process, since it neither changes the sequence nor creates any temporary buffers. The reader field ptr points to the current element of the sequence that is to be read next. The code below demonstrates how to use the sequence writer and reader.

```
CvMemStorage* storage = cvCreateMemStorage(0);
CvSeq* seq = cvCreateSeq( CV_32SC1, sizeof(CvSeq), sizeof(int), storage );
CvSeqWriter writer;
CvSeqReader reader;
int i;

cvStartAppendToSeq( seq, &writer );
for( i = 0; i < 10; i++ )
{
    int val = rand()
    CV_WRITE_SEQ_ELEM( val, writer );
    printf("
}
cvEndWriteSeq( &writer );

cvStartReadSeq( seq, &reader, 0 );
for( i = 0; i < seq->total; i++ )
{
    int val;
#if 1
    CV_READ_SEQ_ELEM( val, reader );
    printf("
#else /* alternative way, that is prefferable if sequence elements are large,
         or their size/type is unknown at compile time */
    printf("
    CV_NEXT_SEQ_ELEM( seq->elem_size, reader );
#endif
}
...

cvReleaseStorage( &storage );
```

## StartWriteSeq

Creates a new sequence and initializes a writer for it.

**C:** void **cvStartWriteSeq**(int **seq_flags**, int **header_size**, int **elem_size**, CvMemStorage* **storage**, CvSeqWriter* **writer**)

> **Parameters**
>
>> **seq_flags** – Flags of the created sequence. If the sequence is not passed to any function working with a specific type of sequences, the sequence value may be equal to 0; otherwise the appropriate type must be selected from the list of predefined sequence types.
>>
>> **header_size** – Size of the sequence header. The parameter value may not be less than `sizeof(CvSeq)`. If a certain type or extension is specified, it must fit within the base type header.
>>
>> **elem_size** – Size of the sequence elements in bytes; must be consistent with the sequence type. For example, if a sequence of points is created (element type `CV_SEQ_ELTYPE_POINT`), then the parameter `elem_size` must be equal to `sizeof(CvPoint)`.
>>
>> **storage** – Sequence location
>>
>> **writer** – Writer state; initialized by the function

The function is a combination of `CreateSeq()` and `StartAppendToSeq()`. The pointer to the created sequence is stored at `writer->seq` and is also returned by the `EndWriteSeq()` function that should be called at the end.

---

### TreeToNodeSeq

Gathers all node pointers to a single sequence.

**C:** CvSeq* **cvTreeToNodeSeq**(const void* **first**, int **header_size**, CvMemStorage* **storage**)

> **Parameters**
>
>> **first** – The initial tree node
>>
>> **header_size** – Header size of the created sequence (sizeof(CvSeq) is the most frequently used value)
>>
>> **storage** – Container for the sequence

The function puts pointers of all nodes reachable from `first` into a single sequence. The pointers are written sequentially in the depth-first order.

## 2.4 Operations on Arrays

### abs

Calculates an absolute value of each matrix element.

**C++:** MatExpr **abs**(const Mat& **m**)

**C++:** MatExpr **abs**(const MatExpr& **e**)

> **Parameters**
>
>> **m** – matrix.
>>
>> **e** – matrix expression.

abs is a meta-function that is expanded to one of `absdiff()` forms:

- `C = abs(A-B)` is equivalent to `absdiff(A, B, C)`

- `C = abs(A)` is equivalent to `absdiff(A, Scalar::all(0), C)`

- `C = Mat_<Vec<uchar,n> >(abs(A*alpha + beta))` is equivalent to `convertScaleAbs` (A, C, alpha, beta)

The output matrix has the same size and the same type as the input one except for the last case, where C is `depth=CV_8U`.

> **See Also:**
>
> *Matrix Expressions*, `absdiff()`

### absdiff

Calculates the per-element absolute difference between two arrays or between an array and a scalar.

**C++:** void **absdiff**(InputArray **src1**, InputArray **src2**, OutputArray **dst**)

**Python:** cv2.**absdiff**(src1, src2[, dst]) → dst

**C:** void **cvAbsDiff**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**)

**C:** void **cvAbsDiffS**(const CvArr* **src**, CvArr* **dst**, CvScalar **value**)

**Python:** cv.**AbsDiff**(src1, src2, dst) → None

---

**Python:** `cv.AbsDiffS`(src, dst, value) $\rightarrow$ None

> **Parameters**
>
> > **src1** – first input array or a scalar.
> >
> > **src2** – second input array or a scalar.
> >
> > **src** – single input array.
> >
> > **value** – scalar value.
> >
> > **dst** – output array that has the same size and type as input arrays.

The function `absdiff` calculates:

- Absolute difference between two arrays when they have the same size and type:

$$\texttt{dst}(I) = \texttt{saturate}(|\texttt{src1}(I) - \texttt{src2}(I)|)$$

- Absolute difference between an array and a scalar when the second array is constructed from `Scalar` or has as many elements as the number of channels in `src1`:

$$\texttt{dst}(I) = \texttt{saturate}(|\texttt{src1}(I) - \texttt{src2}|)$$

- Absolute difference between a scalar and an array when the first array is constructed from `Scalar` or has as many elements as the number of channels in `src2`:

$$\texttt{dst}(I) = \texttt{saturate}(|\texttt{src1} - \texttt{src2}(I)|)$$

where `I` is a multi-dimensional index of array elements. In case of multi-channel arrays, each channel is processed independently.

---

**Note:** Saturation is not applied when the arrays have the depth `CV_32S`. You may even get a negative value in the case of overflow.

---

**See Also:**

`abs()`

## add

Calculates the per-element sum of two arrays or an array and a scalar.

**C++:** void **add**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, InputArray **mask**=noArray(), int **dtype**=-1)

**Python:** `cv2.add`(src1, src2[, dst[, mask[, dtype ] ] ]) $\rightarrow$ dst

**C:** void **cvAdd**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, const CvArr* **mask**=NULL)

**C:** void **cvAddS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL)

**Python:** `cv.Add`(src1, src2, dst, mask=None) $\rightarrow$ None

**Python:** `cv.AddS`(src, value, dst, mask=None) $\rightarrow$ None

> **Parameters**
>
> > **src1** – first input array or a scalar.
> >
> > **src2** – second input array or a scalar.

> **src** – single input array.
>
> **value** – scalar value.
>
> **dst** – output array that has the same size and number of channels as the input array(s); the depth is defined by `dtype` or `src1`/`src2`.
>
> **mask** – optional operation mask – 8-bit single channel array, that specifies elements of the output array to be changed.
>
> **dtype** – optional depth of the output array (see the discussion below).

The function `add` calculates:

- Sum of two arrays when both input arrays have the same size and the same number of channels:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{src1}(I) + \texttt{src2}(I)) \quad \text{if mask}(I) \neq 0$$

- Sum of an array and a scalar when `src2` is constructed from `Scalar` or has the same number of elements as `src1.channels()`:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{src1}(I) + \texttt{src2}) \quad \text{if mask}(I) \neq 0$$

- Sum of a scalar and an array when `src1` is constructed from `Scalar` or has the same number of elements as `src2.channels()`:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{src1} + \texttt{src2}(I)) \quad \text{if mask}(I) \neq 0$$

where `I` is a multi-dimensional index of array elements. In case of multi-channel arrays, each channel is processed independently.

The first function in the list above can be replaced with matrix expressions:

```
dst = src1 + src2;
dst += src1; // equivalent to add(dst, src1, dst);
```

The input arrays and the output array can all have the same or different depths. For example, you can add a 16-bit unsigned array to a 8-bit signed array and store the sum as a 32-bit floating-point array. Depth of the output array is determined by the `dtype` parameter. In the second and third cases above, as well as in the first case, when `src1.depth() == src2.depth()`, `dtype` can be set to the default -1. In this case, the output array will have the same depth as the input array, be it `src1`, `src2` or both.

---

**Note:** Saturation is not applied when the output array has the depth `CV_32S`. You may even get result of an incorrect sign in the case of overflow.

---

**See Also:**

`subtract()`, `addWeighted()`, `scaleAdd()`, `Mat::convertTo()`, *Matrix Expressions*

## addWeighted

Calculates the weighted sum of two arrays.

**C++:** void **addWeighted**(InputArray **src1**, double **alpha**, InputArray **src2**, double **beta**, double **gamma**, OutputArray **dst**, int **dtype**=-1)

**Python:** cv2.**addWeighted**(src1, alpha, src2, beta, gamma[, dst[, dtype ]]) → dst

**C:** void **cvAddWeighted**(const CvArr* **src1**, double **alpha**, const CvArr* **src2**, double **beta**, double **gamma**, CvArr* **dst**)

---

**Python:** cv.**AddWeighted**(src1, alpha, src2, beta, gamma, dst) → None

>  **Parameters**

> > **src1** – first input array.

> > **alpha** – weight of the first array elements.

> > **src2** – second input array of the same size and channel number as `src1`.

> > **beta** – weight of the second array elements.

> > **dst** – output array that has the same size and number of channels as the input arrays.

> > **gamma** – scalar added to each sum.

> > **dtype** – optional depth of the output array; when both input arrays have the same depth, `dtype` can be set to `-1`, which will be equivalent to `src1.depth()`.

The function `addWeighted` calculates the weighted sum of two arrays as follows:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{src1}(I) * \texttt{alpha} + \texttt{src2}(I) * \texttt{beta} + \texttt{gamma})$$

where `I` is a multi-dimensional index of array elements. In case of multi-channel arrays, each channel is processed independently.

The function can be replaced with a matrix expression:

```
dst = src1*alpha + src2*beta + gamma;
```

---

**Note:** Saturation is not applied when the output array has the depth `CV_32S`. You may even get result of an incorrect sign in the case of overflow.

---

**See Also:**

add(), subtract(), scaleAdd(), Mat::convertTo(), *Matrix Expressions*

## bitwise_and

Calculates the per-element bit-wise conjunction of two arrays or an array and a scalar.

**C++:** void **bitwise_and**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, InputArray **mask**=noArray())

**Python:** cv2.**bitwise_and**(src1, src2[, dst[, mask]]) → dst

**C:** void **cvAnd**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, const CvArr* **mask**=NULL)

**C:** void **cvAndS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL)

**Python:** cv.**And**(src1, src2, dst, mask=None) → None

**Python:** cv.**AndS**(src, value, dst, mask=None) → None

>  **Parameters**

> > **src1** – first input array or a scalar.

> > **src2** – second input array or a scalar.

> > **src** – single input array.

> > **value** – scalar value.

> > **dst** – output array that has the same size and type as the input arrays.

       **mask** – optional operation mask, 8-bit single channel array, that specifies elements of the output array to be changed.

The function calculates the per-element bit-wise logical conjunction for:

- Two arrays when `src1` and `src2` have the same size:

$$\mathtt{dst}(I) = \mathtt{src1}(I) \wedge \mathtt{src2}(I) \quad \text{if } \mathtt{mask}(I) \neq 0$$

- An array and a scalar when `src2` is constructed from `Scalar` or has the same number of elements as `src1.channels()`:

$$\mathtt{dst}(I) = \mathtt{src1}(I) \wedge \mathtt{src2} \quad \text{if } \mathtt{mask}(I) \neq 0$$

- A scalar and an array when `src1` is constructed from `Scalar` or has the same number of elements as `src2.channels()`:

$$\mathtt{dst}(I) = \mathtt{src1} \wedge \mathtt{src2}(I) \quad \text{if } \mathtt{mask}(I) \neq 0$$

In case of floating-point arrays, their machine-specific bit representations (usually IEEE754-compliant) are used for the operation. In case of multi-channel arrays, each channel is processed independently. In the second and third cases above, the scalar is first converted to the array type.

## bitwise_not

Inverts every bit of an array.

**C++:** void **bitwise_not**(InputArray **src**, OutputArray **dst**, InputArray **mask**=noArray())

**Python:** cv2.**bitwise_not**(src[, dst[, mask]]) → dst

**C:** void **cvNot**(const CvArr* **src**, CvArr* **dst**)

**Python:** cv.**Not**(src, dst) → None

       **Parameters**

              **src** – input array.

              **dst** – output array that has the same size and type as the input array.

              **mask** – optional operation mask, 8-bit single channel array, that specifies elements of the output array to be changed.

The function calculates per-element bit-wise inversion of the input array:

$$\mathtt{dst}(I) = \neg \mathtt{src}(I)$$

In case of a floating-point input array, its machine-specific bit representation (usually IEEE754-compliant) is used for the operation. In case of multi-channel arrays, each channel is processed independently.

## bitwise_or

Calculates the per-element bit-wise disjunction of two arrays or an array and a scalar.

**C++:** void **bitwise_or**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, InputArray **mask**=noArray())

**Python:** cv2.**bitwise_or**(src1, src2[, dst[, mask]]) → dst

**C:** void **cvOr**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, const CvArr* **mask**=NULL)

**C:** void **cvOrS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL)

**Python:** cv.**Or**(src1, src2, dst, mask=None) → None

**Python:** cv.**OrS**(src, value, dst, mask=None) → None

> **Parameters**

>> **src1** – first input array or a scalar.

>> **src2** – second input array or a scalar.

>> **src** – single input array.

>> **value** – scalar value.

>> **dst** – output array that has the same size and type as the input arrays.

>> **mask** – optional operation mask, 8-bit single channel array, that specifies elements of the output array to be changed.

The function calculates the per-element bit-wise logical disjunction for:

- Two arrays when `src1` and `src2` have the same size:

$$\texttt{dst}(I) = \texttt{src1}(I) \vee \texttt{src2}(I) \quad \text{if } \texttt{mask}(I) \neq 0$$

- An array and a scalar when `src2` is constructed from `Scalar` or has the same number of elements as `src1.channels()`:

$$\texttt{dst}(I) = \texttt{src1}(I) \vee \texttt{src2} \quad \text{if } \texttt{mask}(I) \neq 0$$

- A scalar and an array when `src1` is constructed from `Scalar` or has the same number of elements as `src2.channels()`:

$$\texttt{dst}(I) = \texttt{src1} \vee \texttt{src2}(I) \quad \text{if } \texttt{mask}(I) \neq 0$$

In case of floating-point arrays, their machine-specific bit representations (usually IEEE754-compliant) are used for the operation. In case of multi-channel arrays, each channel is processed independently. In the second and third cases above, the scalar is first converted to the array type.

## bitwise_xor

Calculates the per-element bit-wise "exclusive or" operation on two arrays or an array and a scalar.

**C++:** void **bitwise_xor**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, InputArray **mask**=noArray())

**Python:** cv2.**bitwise_xor**(src1, src2[, dst[, mask]]) → dst

**C:** void **cvXor**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, const CvArr* **mask**=NULL)

**C:** void **cvXorS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL)

**Python:** cv.**Xor**(src1, src2, dst, mask=None) → None

**Python:** cv.**XorS**(src, value, dst, mask=None) → None

> **Parameters**

> **src1** – first input array or a scalar.
>
> **src2** – second input array or a scalar.
>
> **src** – single input array.
>
> **value** – scalar value.
>
> **dst** – output array that has the same size and type as the input arrays.
>
> **mask** – optional operation mask, 8-bit single channel array, that specifies elements of the output array to be changed.

The function calculates the per-element bit-wise logical "exclusive-or" operation for:

- Two arrays when `src1` and `src2` have the same size:

$$\texttt{dst}(I) = \texttt{src1}(I) \oplus \texttt{src2}(I) \quad \text{if } \texttt{mask}(I) \neq 0$$

- An array and a scalar when `src2` is constructed from `Scalar` or has the same number of elements as `src1.channels()`:

$$\texttt{dst}(I) = \texttt{src1}(I) \oplus \texttt{src2} \quad \text{if } \texttt{mask}(I) \neq 0$$

- A scalar and an array when `src1` is constructed from `Scalar` or has the same number of elements as `src2.channels()`:

$$\texttt{dst}(I) = \texttt{src1} \oplus \texttt{src2}(I) \quad \text{if } \texttt{mask}(I) \neq 0$$

In case of floating-point arrays, their machine-specific bit representations (usually IEEE754-compliant) are used for the operation. In case of multi-channel arrays, each channel is processed independently. In the 2nd and 3rd cases above, the scalar is first converted to the array type.

## calcCovarMatrix

Calculates the covariance matrix of a set of vectors.

**C++:** void **calcCovarMatrix**(const Mat* **samples**, int **nsamples**, Mat& **covar**, Mat& **mean**, int **flags**, int **ctype**=CV_64F)

**C++:** void **calcCovarMatrix**(InputArray **samples**, OutputArray **covar**, OutputArray **mean**, int **flags**, int **ctype**=CV_64F)

**Python:** cv2.**calcCovarMatrix**(samples, flags[, covar[, mean[, ctype]]]) → covar, mean

**C:** void **cvCalcCovarMatrix**(const CvArr** **vects**, int **count**, CvArr* **cov_mat**, CvArr* **avg**, int **flags**)

**Python:** cv.**CalcCovarMatrix**(vects, covMat, avg, flags) → None

> **Parameters**
>
> > **samples** – samples stored either as separate matrices or as rows/columns of a single matrix.
> >
> > **nsamples** – number of samples when they are stored separately.
> >
> > **covar** – output covariance matrix of the type `ctype` and square size.

**ctype** – type of the matrixl; it equals 'CV_64F' by default.

**mean** – input or output (depending on the flags) array as the average value of the input vectors.

**vects** – a set of vectors.

**flags** – operation flags as a combination of the following values:

– **CV_COVAR_SCRAMBLED** The output covariance matrix is calculated as:

$$\text{scale} \cdot [\text{vects}[0] - \text{mean}, \text{vects}[1] - \text{mean}, ...]^T \cdot [\text{vects}[0] - \text{mean}, \text{vects}[1] - \text{mean}, ...],$$

The covariance matrix will be `nsamples x nsamples`. Such an unusual covariance matrix is used for fast PCA of a set of very large vectors (see, for example, the EigenFaces technique for face recognition). Eigenvalues of this "scrambled" matrix match the eigenvalues of the true covariance matrix. The "true" eigenvectors can be easily calculated from the eigenvectors of the "scrambled" covariance matrix.

– **CV_COVAR_NORMAL** The output covariance matrix is calculated as:

$$\text{scale} \cdot [\text{vects}[0] - \text{mean}, \text{vects}[1] - \text{mean}, ...] \cdot [\text{vects}[0] - \text{mean}, \text{vects}[1] - \text{mean}, ...]^T,$$

`covar` will be a square matrix of the same size as the total number of elements in each input vector. One and only one of `CV_COVAR_SCRAMBLED` and `CV_COVAR_NORMAL` must be specified.

– **CV_COVAR_USE_AVG** If the flag is specified, the function does not calculate `mean` from the input vectors but, instead, uses the passed `mean` vector. This is useful if `mean` has been pre-calculated or known in advance, or if the covariance matrix is calculated by parts. In this case, `mean` is not a mean vector of the input sub-set of vectors but rather the mean vector of the whole set.

– **CV_COVAR_SCALE** If the flag is specified, the covariance matrix is scaled. In the "normal" mode, `scale` is `1./nsamples` . In the "scrambled" mode, `scale` is the reciprocal of the total number of elements in each input vector. By default (if the flag is not specified), the covariance matrix is not scaled ( `scale=1` ).

– **CV_COVAR_ROWS** [Only useful in the second variant of the function] If the flag is specified, all the input vectors are stored as rows of the `samples` matrix. `mean` should be a single-row vector in this case.

– **CV_COVAR_COLS** [Only useful in the second variant of the function] If the flag is specified, all the input vectors are stored as columns of the `samples` matrix. `mean` should be a single-column vector in this case.

The functions `calcCovarMatrix` calculate the covariance matrix and, optionally, the mean vector of the set of input vectors.

**See Also:**

`PCA`, `mulTransposed()`, `Mahalanobis()`

## cartToPolar

Calculates the magnitude and angle of 2D vectors.

**C++:** void **cartToPolar**(InputArray **x**, InputArray **y**, OutputArray **magnitude**, OutputArray **angle**, bool **angleInDegrees**=false)

**Python:** cv2.**cartToPolar**(x, y[, magnitude[, angle[, angleInDegrees ]]]) → magnitude, angle

**C:** void **cvCartToPolar**(const CvArr* **x**, const CvArr* **y**, CvArr* **magnitude**, CvArr* **angle**=NULL, int **angle_in_degrees**=0 )

**Python:** cv.**CartToPolar**(x, y, magnitude, angle=None, angleInDegrees=0) → None

> **Parameters**
>
>> **x** – array of x-coordinates; this must be a single-precision or double-precision floating-point array.
>>
>> **y** – array of y-coordinates, that must have the same size and same type as x.
>>
>> **magnitude** – output array of magnitudes of the same size and type as x.
>>
>> **angle** – output array of angles that has the same size and type as x; the angles are measured in radians (from 0 to 2*Pi) or in degrees (0 to 360 degrees).
>>
>> **angleInDegrees** – a flag, indicating whether the angles are measured in radians (which is by default), or in degrees.
>>
>> **angle_in_degrees** – a flag, indicating whether the angles are measured in radians, or in degrees (specific to C syntax).

The function cartToPolar calculates either the magnitude, angle, or both for every 2D vector (x(I),y(I)):

$$\text{magnitude}(I) = \sqrt{x(I)^2 + y(I)^2},$$
$$\text{angle}(I) = \text{atan2}(y(I), x(I))[\cdot 180/\pi]$$

The angles are calculated with accuracy about 0.3 degrees. For the point (0,0), the angle is set to 0.

**See Also:**

Sobel(), Scharr()

## checkRange

Checks every element of an input array for invalid values.

**C++:** bool **checkRange**(InputArray **a**, bool **quiet**=true, Point* **pos**=0, double **minVal**=-DBL_MAX, double **maxVal**=DBL_MAX )

**Python:** cv2.**checkRange**(a[, quiet[, minVal[, maxVal ]]]) → retval, pos

> **Parameters**
>
>> **a** – input array.
>>
>> **quiet** – a flag, indicating whether the functions quietly return false when the array elements are out of range or they throw an exception.
>>
>> **pos** – optional output parameter, where the position of the first outlier is stored; in the second function pos, when not NULL, must be a pointer to array of src.dims elements.
>>
>> **minVal** – inclusive lower boundary of valid values range.
>>
>> **maxVal** – exclusive upper boundary of valid values range.

The functions `checkRange` check that every array element is neither NaN nor infinite. When `minVal < -DBL_MAX` and `maxVal < DBL_MAX`, the functions also check that each value is between `minVal` and `maxVal`. In case of multi-channel arrays, each channel is processed independently. If some values are out of range, position of the first outlier is stored in `pos` (when `pos != NULL`). Then, the functions either return false (when `quiet=true`) or throw an exception.

## compare

Performs the per-element comparison of two arrays or an array and scalar value.

**C++:** void **compare**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, int **cmpop**)

**Python:** cv2.**compare**(src1, src2, cmpop[, dst]) → dst

**C:** void **cvCmp**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, int **cmp_op**)

**Python:** cv.**Cmp**(src1, src2, dst, cmpOp) → None

**C:** void **cvCmpS**(const CvArr* **src**, double **value**, CvArr* **dst**, int **cmp_op**)

**Python:** cv.**CmpS**(src, value, dst, cmpOp) → None

> **Parameters**

>> **src1** – first input array or a scalar (in the case of `cvCmp`, `cv.Cmp`, `cvCmpS`, `cv.CmpS` it is always an array); when it is an array, it must have a single channel.

>> **src2** – second input array or a scalar (in the case of `cvCmp` and `cv.Cmp` it is always an array; in the case of `cvCmpS`, `cv.CmpS` it is always a scalar); when it is an array, it must have a single channel.

>> **src** – single input array.

>> **value** – scalar value.

>> **dst** – output array that has the same size as the input arrays and type= `CV_8UC1` .

>> **cmpop** – a flag, that specifies correspondence between the arrays:

>> – **CMP_EQ** `src1` is equal to `src2`.

>> – **CMP_GT** `src1` is greater than `src2`.

>> – **CMP_GE** `src1` is greater than or equal to `src2`.

>> – **CMP_LT** `src1` is less than `src2`.

>> – **CMP_LE** `src1` is less than or equal to `src2`.

>> – **CMP_NE** `src1` is unequal to `src2`.

The function compares:

- Elements of two arrays when `src1` and `src2` have the same size:

$$\texttt{dst}(I) = \texttt{src1}(I) \ \texttt{cmpop} \ \texttt{src2}(I)$$

- Elements of `src1` with a scalar `src2` when `src2` is constructed from `Scalar` or has a single element:

$$\texttt{dst}(I) = \texttt{src1}(I) \ \texttt{cmpop} \ \texttt{src2}$$

- `src1` with elements of `src2` when `src1` is constructed from `Scalar` or has a single element:

$$\texttt{dst}(I) = \texttt{src1} \ \texttt{cmpop} \ \texttt{src2}(I)$$

When the comparison result is true, the corresponding element of output array is set to 255. The comparison operations can be replaced with the equivalent matrix expressions:

```
Mat dst1 = src1 >= src2;
Mat dst2 = src1 < 8;
...
```

**See Also:**

`checkRange()`, `min()`, `max()`, `threshold()`, *Matrix Expressions*

## completeSymm

Copies the lower or the upper half of a square matrix to another half.

**C++:** void **completeSymm**(InputOutputArray **mtx**, bool **lowerToUpper**=false)

**Python:** cv2.**completeSymm**(mtx[, lowerToUpper]) → None

> **Parameters**
>
> > **mtx** – input-output floating-point square matrix.
> >
> > **lowerToUpper** – operation flag; if true, the lower half is copied to the upper half. Otherwise, the upper half is copied to the lower half.

The function `completeSymm` copies the lower half of a square matrix to its another half. The matrix diagonal remains unchanged:

- $\mathtt{mtx}_{ij} = \mathtt{mtx}_{ji}$ for $i > j$ if `lowerToUpper=false`
- $\mathtt{mtx}_{ij} = \mathtt{mtx}_{ji}$ for $i < j$ if `lowerToUpper=true`

**See Also:**

`flip()`, `transpose()`

## convertScaleAbs

Scales, calculates absolute values, and converts the result to 8-bit.

**C++:** void **convertScaleAbs**(InputArray **src**, OutputArray **dst**, double **alpha**=1, double **beta**=0)

**Python:** cv2.**convertScaleAbs**(src[, dst[, alpha[, beta]]]) → dst

**C:** void **cvConvertScaleAbs**(const CvArr* **src**, CvArr* **dst**, double **scale**=1, double **shift**=0)

**Python:** cv.**ConvertScaleAbs**(src, dst, scale=1.0, shift=0.0) → None

> **Parameters**
>
> > **src** – input array.
> >
> > **dst** – output array.
> >
> > **alpha** – optional scale factor.
> >
> > **beta** – optional delta added to the scaled values.

On each element of the input array, the function `convertScaleAbs` performs three operations sequentially: scaling, taking an absolute value, conversion to an unsigned 8-bit type:

$$\mathtt{dst}(I) = \mathtt{saturate\_cast<uchar>}(|\mathtt{src}(I) * \mathtt{alpha} + \mathtt{beta}|)$$

In case of multi-channel arrays, the function processes each channel independently. When the output is not 8-bit, the operation can be emulated by calling the `Mat::convertTo` method (or by using matrix expressions) and then by calculating an absolute value of the result. For example:

```
Mat_<float> A(30,30);
randu(A, Scalar(-100), Scalar(100));
Mat_<float> B = A*5 + 3;
B = abs(B);
// Mat_<float> B = abs(A*5+3) will also do the job,
// but it will allocate a temporary matrix
```

**See Also:**

`Mat::convertTo()`, `abs()`

## countNonZero

Counts non-zero array elements.

**C++:** int **countNonZero**(InputArray **src**)

**Python:** cv2.**countNonZero**(src) → retval

**C:** int **cvCountNonZero**(const CvArr* **arr**)

**Python:** cv.**CountNonZero**(arr) → int

>    **Parameters**

>        **src** – single-channel array.

The function returns the number of non-zero elements in `src` :

$$\sum_{I:\ \text{src}(I)\neq 0} 1$$

**See Also:**

`mean()`, `meanStdDev()`, `norm()`, `minMaxLoc()`, `calcCovarMatrix()`

## cvarrToMat

Converts `CvMat`, `IplImage` , or `CvMatND` to `Mat`.

**C++:** Mat **cvarrToMat**(const CvArr* **arr**, bool **copyData**=false, bool **allowND**=true, int **coiMode**=0 )

>    **Parameters**

>        **arr** – input `CvMat`, `IplImage` , or `CvMatND`.

>        **copyData** – when false (default value), no data is copied and only the new header is created, in this case, the original array should not be deallocated while the new matrix header is used; if the parameter is true, all the data is copied and you may deallocate the original array right after the conversion.

>        **allowND** – when true (default value), `CvMatND` is converted to 2-dimensional `Mat`, if it is possible (see the discussion below); if it is not possible, or when the parameter is false, the function will report an error.

>        **coiMode** – parameter specifying how the IplImage COI (when set) is handled.

– If `coiMode=0` and COI is set, the function reports an error.

– If `coiMode=1` , the function never reports an error. Instead, it returns the header to the whole original image and you will have to check and process COI manually. See `extractImageCOI()` .

The function `cvarrToMat` converts `CvMat`, `IplImage` , or `CvMatND` header to `Mat` header, and optionally duplicates the underlying data. The constructed header is returned by the function.

When `copyData=false` , the conversion is done really fast (in O(1) time) and the newly created matrix header will have `refcount=0` , which means that no reference counting is done for the matrix data. In this case, you have to preserve the data until the new header is destructed. Otherwise, when `copyData=true` , the new buffer is allocated and managed as if you created a new matrix from scratch and copied the data there. That is, `cvarrToMat(arr, true)` is equivalent to `cvarrToMat(arr, false).clone()` (assuming that COI is not set). The function provides a uniform way of supporting `CvArr` paradigm in the code that is migrated to use new-style data structures internally. The reverse transformation, from `Mat` to `CvMat` or `IplImage` can be done by a simple assignment:

```
CvMat* A = cvCreateMat(10, 10, CV_32F);
cvSetIdentity(A);
IplImage A1; cvGetImage(A, &A1);
Mat B = cvarrToMat(A);
Mat B1 = cvarrToMat(&A1);
IplImage C = B;
CvMat C1 = B1;
// now A, A1, B, B1, C and C1 are different headers
// for the same 10x10 floating-point array.
// note that you will need to use "&"
// to pass C & C1 to OpenCV functions, for example:
printf("%g\n", cvNorm(&C1, 0, CV_L2));
```

Normally, the function is used to convert an old-style 2D array ( `CvMat` or `IplImage` ) to `Mat` . However, the function can also take `CvMatND` as an input and create `Mat()` for it, if it is possible. And, for `CvMatND A` , it is possible if and only if `A.dim[i].size*A.dim.step[i] == A.dim.step[i-1]` for all or for all but one i, `0 < i < A.dims` . That is, the matrix data should be continuous or it should be representable as a sequence of continuous matrices. By using this function in this way, you can process `CvMatND` using an arbitrary element-wise function.

The last parameter, `coiMode` , specifies how to deal with an image with COI set. By default, it is 0 and the function reports an error when an image with COI comes in. And `coiMode=1` means that no error is signalled. You have to check COI presence and handle it manually. The modern structures, such as `Mat` and `MatND` do not support COI natively. To process an individual channel of a new-style array, you need either to organize a loop over the array (for example, using matrix iterators) where the channel of interest will be processed, or extract the COI using `mixChannels()` (for new-style arrays) or `extractImageCOI()` (for old-style arrays), process this individual channel, and insert it back to the output array if needed (using `mixChannels()` or `insertImageCOI()` , respectively).

**See Also:**

`cvGetImage()`, `cvGetMat()`, `extractImageCOI()`, `insertImageCOI()`, `mixChannels()`

## dct

Performs a forward or inverse discrete Cosine transform of 1D or 2D array.

**C++:** void **dct**(InputArray **src**, OutputArray **dst**, int **flags**=0)

**Python:** cv2.**dct**(src[, dst[, flags ]]) → dst

**C:** void **cvDCT**(const CvArr* **src**, CvArr* **dst**, int **flags**)

**Python:** cv.**DCT**(src, dst, flags) → None

**Parameters**

**src** – input floating-point array.

**dst** – output array of the same size and type as `src` .

**flags** – transformation flags as a combination of the following values:

– **DCT_INVERSE** performs an inverse 1D or 2D transform instead of the default forward transform.

– **DCT_ROWS** performs a forward or inverse transform of every individual row of the input matrix. This flag enables you to transform multiple vectors simultaneously and can be used to decrease the overhead (which is sometimes several times larger than the processing itself) to perform 3D and higher-dimensional transforms and so forth.

The function `dct` performs a forward or inverse discrete Cosine transform (DCT) of a 1D or 2D floating-point array:

• Forward Cosine transform of a 1D vector of `N` elements:

$$Y = C^{(N)} \cdot X$$

where

$$C^{(N)}_{jk} = \sqrt{\alpha_j/N} \cos \left( \frac{\pi(2k+1)j}{2N} \right)$$

and

$\alpha_0 = 1$, $\alpha_j = 2$ for *j > 0*.

• Inverse Cosine transform of a 1D vector of `N` elements:

$$X = \left( C^{(N)} \right)^{-1} \cdot Y = \left( C^{(N)} \right)^T \cdot Y$$

(since $C^{(N)}$ is an orthogonal matrix, $C^{(N)} \cdot \left( C^{(N)} \right)^T = I$ )

• Forward 2D Cosine transform of `M x N` matrix:

$$Y = C^{(N)} \cdot X \cdot \left( C^{(N)} \right)^T$$

• Inverse 2D Cosine transform of `M x N` matrix:

$$X = \left( C^{(N)} \right)^T \cdot X \cdot C^{(N)}$$

The function chooses the mode of operation by looking at the flags and size of the input array:

• If `(flags & DCT_INVERSE) == 0` , the function does a forward 1D or 2D transform. Otherwise, it is an inverse 1D or 2D transform.

• If `(flags & DCT_ROWS) != 0` , the function performs a 1D transform of each row.

• If the array is a single column or a single row, the function performs a 1D transform.

• If none of the above is true, the function performs a 2D transform.

---

**Note:** Currently `dct` supports even-size arrays (2, 4, 6 ...). For data analysis and approximation, you can pad the array when necessary.

Also, the function performance depends very much, and not monotonically, on the array size (see `getOptimalDFTSize()` ). In the current implementation DCT of a vector of size `N` is calculated via DFT of a vector of size `N/2` . Thus, the optimal DCT size `N1 >= N` can be calculated as:

---

```
size_t getOptimalDCTSize(size_t N) { return 2*getOptimalDFTSize((N+1)/2); }
N1 = getOptimalDCTSize(N);
```

**See Also:**

`dft()`, `getOptimalDFTSize()`, `idct()`

## dft

Performs a forward or inverse Discrete Fourier transform of a 1D or 2D floating-point array.

**C++:** void **dft**(InputArray **src**, OutputArray **dst**, int **flags**=0, int **nonzeroRows**=0)

**Python:** cv2.**dft**(src[, dst[, flags[, nonzeroRows ]]]) → dst

**C:** void **cvDFT**(const CvArr* **src**, CvArr* **dst**, int **flags**, int **nonzero_rows**=0 )

**Python:** cv.**DFT**(src, dst, flags, nonzeroRows=0) → None

> **Parameters**
>
> > **src** – input array that could be real or complex.
> >
> > **dst** – output array whose size and type depends on the `flags` .
> >
> > **flags** – transformation flags, representing a combination of the following values:
> >
> > – **DFT_INVERSE** performs an inverse 1D or 2D transform instead of the default forward transform.
> >
> > – **DFT_SCALE** scales the result: divide it by the number of array elements. Normally, it is combined with `DFT_INVERSE`.
> >
> > – **DFT_ROWS** performs a forward or inverse transform of every individual row of the input matrix; this flag enables you to transform multiple vectors simultaneously and can be used to decrease the overhead (which is sometimes several times larger than the processing itself) to perform 3D and higher-dimensional transformations and so forth.
> >
> > – **DFT_COMPLEX_OUTPUT** performs a forward transformation of 1D or 2D real array; the result, though being a complex array, has complex-conjugate symmetry (*CCS*, see the function description below for details), and such an array can be packed into a real array of the same size as input, which is the fastest option and which is what the function does by default; however, you may wish to get a full complex array (for simpler spectrum analysis, and so on) – pass the flag to enable the function to produce a full-size complex output array.
> >
> > – **DFT_REAL_OUTPUT** performs an inverse transformation of a 1D or 2D complex array; the result is normally a complex array of the same size, however, if the input array has conjugate-complex symmetry (for example, it is a result of forward transformation with `DFT_COMPLEX_OUTPUT` flag), the output is a real array; while the function itself does not check whether the input is symmetrical or not, you can pass the flag and then the function will assume the symmetry and produce the real output array (note that when the input is packed into a real array and inverse transformation is executed, the function treats the input as a packed complex-conjugate symmetrical array, and the output will also be a real array).
> >
> > **nonzeroRows** – when the parameter is not zero, the function assumes that only the first `nonzeroRows` rows of the input array (DFT_INVERSE is not set) or only the first `nonzeroRows` of the output array (DFT_INVERSE is set) contain non-zeros, thus, the function

can handle the rest of the rows more efficiently and save some time; this technique is very useful for calculating array cross-correlation or convolution using DFT.

The function performs one of the following:

- Forward the Fourier transform of a 1D vector of N elements:

$$Y = F^{(N)} \cdot X,$$

where $F_{jk}^{(N)} = \exp(-2\pi i j k/N)$ and $i = \sqrt{-1}$

- Inverse the Fourier transform of a 1D vector of N elements:

$$X' = \left(F^{(N)}\right)^{-1} \cdot Y = \left(F^{(N)}\right)^* \cdot y$$
$$X = (1/N) \cdot X,$$

where $F^* = \left(\mathrm{Re}(F^{(N)}) - \mathrm{Im}(F^{(N)})\right)^T$

- Forward the 2D Fourier transform of a M x N matrix:

$$Y = F^{(M)} \cdot X \cdot F^{(N)}$$

- Inverse the 2D Fourier transform of a M x N matrix:

$$X' = \left(F^{(M)}\right)^* \cdot Y \cdot \left(F^{(N)}\right)^*$$
$$X = \frac{1}{M \cdot N} \cdot X'$$

In case of real (single-channel) data, the output spectrum of the forward Fourier transform or input spectrum of the inverse Fourier transform can be represented in a packed format called *CCS* (complex-conjugate-symmetrical). It was borrowed from IPL (Intel* Image Processing Library). Here is how 2D *CCS* spectrum looks:

$$
\begin{bmatrix}
\mathrm{Re}Y_{0,0} & \mathrm{Re}Y_{0,1} & \mathrm{Im}Y_{0,1} & \mathrm{Re}Y_{0,2} & \mathrm{Im}Y_{0,2} & \cdots & \mathrm{Re}Y_{0,N/2-1} & \mathrm{Im}Y_{0,N/2-1} & \mathrm{Re}Y_{0,N/2} \\
\mathrm{Re}Y_{1,0} & \mathrm{Re}Y_{1,1} & \mathrm{Im}Y_{1,1} & \mathrm{Re}Y_{1,2} & \mathrm{Im}Y_{1,2} & \cdots & \mathrm{Re}Y_{1,N/2-1} & \mathrm{Im}Y_{1,N/2-1} & \mathrm{Re}Y_{1,N/2} \\
\mathrm{Im}Y_{1,0} & \mathrm{Re}Y_{2,1} & \mathrm{Im}Y_{2,1} & \mathrm{Re}Y_{2,2} & \mathrm{Im}Y_{2,2} & \cdots & \mathrm{Re}Y_{2,N/2-1} & \mathrm{Im}Y_{2,N/2-1} & \mathrm{Im}Y_{1,N/2} \\
 & & & \cdots\cdots\cdots\cdots\cdots\cdots\cdots\cdots\cdots\cdots & & & & & \\
\mathrm{Re}Y_{M/2-1,0} & \mathrm{Re}Y_{M-3,1} & \mathrm{Im}Y_{M-3,1} & & \cdots\cdots\cdots & & \mathrm{Re}Y_{M-3,N/2-1} & \mathrm{Im}Y_{M-3,N/2-1} & \mathrm{Re}Y_{M/2-1,N/2} \\
\mathrm{Im}Y_{M/2-1,0} & \mathrm{Re}Y_{M-2,1} & \mathrm{Im}Y_{M-2,1} & & \cdots\cdots\cdots & & \mathrm{Re}Y_{M-2,N/2-1} & \mathrm{Im}Y_{M-2,N/2-1} & \mathrm{Im}Y_{M/2-1,N/2} \\
\mathrm{Re}Y_{M/2,0} & \mathrm{Re}Y_{M-1,1} & \mathrm{Im}Y_{M-1,1} & & \cdots\cdots\cdots & & \mathrm{Re}Y_{M-1,N/2-1} & \mathrm{Im}Y_{M-1,N/2-1} & \mathrm{Re}Y_{M/2,N/2}
\end{bmatrix}
$$

In case of 1D transform of a real vector, the output looks like the first row of the matrix above.

So, the function chooses an operation mode depending on the flags and size of the input array:

- If `DFT_ROWS` is set or the input array has a single row or single column, the function performs a 1D forward or inverse transform of each row of a matrix when `DFT_ROWS` is set. Otherwise, it performs a 2D transform.

- If the input array is real and `DFT_INVERSE` is not set, the function performs a forward 1D or 2D transform:

  - When `DFT_COMPLEX_OUTPUT` is set, the output is a complex matrix of the same size as input.

  - When `DFT_COMPLEX_OUTPUT` is not set, the output is a real matrix of the same size as input. In case of 2D transform, it uses the packed format as shown above. In case of a single 1D transform, it looks like the first row of the matrix above. In case of multiple 1D transforms (when using the `DCT_ROWS` flag), each row of the output matrix looks like the first row of the matrix above.

- If the input array is complex and either `DFT_INVERSE` or `DFT_REAL_OUTPUT` are not set, the output is a complex array of the same size as input. The function performs a forward or inverse 1D or 2D transform of the whole input array or each row of the input array independently, depending on the flags `DFT_INVERSE` and `DFT_ROWS`.

- When `DFT_INVERSE` is set and the input array is real, or it is complex but `DFT_REAL_OUTPUT` is set, the output is a real array of the same size as input. The function performs a 1D or 2D inverse transformation of the whole input array or each individual row, depending on the flags `DFT_INVERSE` and `DFT_ROWS`.

If `DFT_SCALE` is set, the scaling is done after the transformation.

Unlike `dct()` , the function supports arrays of arbitrary size. But only those arrays are processed efficiently, whose sizes can be factorized in a product of small prime numbers (2, 3, and 5 in the current implementation). Such an efficient DFT size can be calculated using the `getOptimalDFTSize()` method.

The sample below illustrates how to calculate a DFT-based convolution of two 2D real arrays:

```cpp
void convolveDFT(InputArray A, InputArray B, OutputArray C)
{
    // reallocate the output array if needed
    C.create(abs(A.rows - B.rows)+1, abs(A.cols - B.cols)+1, A.type());
    Size dftSize;
    // calculate the size of DFT transform
    dftSize.width = getOptimalDFTSize(A.cols + B.cols - 1);
    dftSize.height = getOptimalDFTSize(A.rows + B.rows - 1);

    // allocate temporary buffers and initialize them with 0's
    Mat tempA(dftSize, A.type(), Scalar::all(0));
    Mat tempB(dftSize, B.type(), Scalar::all(0));

    // copy A and B to the top-left corners of tempA and tempB, respectively
    Mat roiA(tempA, Rect(0,0,A.cols,A.rows));
    A.copyTo(roiA);
    Mat roiB(tempB, Rect(0,0,B.cols,B.rows));
    B.copyTo(roiB);

    // now transform the padded A & B in-place;
    // use "nonzeroRows" hint for faster processing
    dft(tempA, tempA, 0, A.rows);
    dft(tempB, tempB, 0, B.rows);

    // multiply the spectrums;
    // the function handles packed spectrum representations well
    mulSpectrums(tempA, tempB, tempA);

    // transform the product back from the frequency domain.
    // Even though all the result rows will be non-zero,
    // you need only the first C.rows of them, and thus you
    // pass nonzeroRows == C.rows
    dft(tempA, tempA, DFT_INVERSE + DFT_SCALE, C.rows);

    // now copy the result back to C.
    tempA(Rect(0, 0, C.cols, C.rows)).copyTo(C);

    // all the temporary buffers will be deallocated automatically
}
```

To optimize this sample, consider the following approaches:

- Since `nonzeroRows != 0` is passed to the forward transform calls and since A and B are copied to the top-left corners of `tempA` and `tempB`, respectively, it is not necessary to clear the whole `tempA` and `tempB`. It is only necessary to clear the `tempA.cols - A.cols` ( `tempB.cols - B.cols`) rightmost columns of the matrices.

- This DFT-based convolution does not have to be applied to the whole big arrays, especially if B is significantly smaller than A or vice versa. Instead, you can calculate convolution by parts. To do this, you need to split

the output array C into multiple tiles. For each tile, estimate which parts of A and B are required to calculate convolution in this tile. If the tiles in C are too small, the speed will decrease a lot because of repeated work. In the ultimate case, when each tile in C is a single pixel, the algorithm becomes equivalent to the naive convolution algorithm. If the tiles are too big, the temporary arrays tempA and tempB become too big and there is also a slowdown because of bad cache locality. So, there is an optimal tile size somewhere in the middle.

- If different tiles in C can be calculated in parallel and, thus, the convolution is done by parts, the loop can be threaded.

All of the above improvements have been implemented in matchTemplate() and filter2D() . Therefore, by using them, you can get the performance even better than with the above theoretically optimal implementation. Though, those two functions actually calculate cross-correlation, not convolution, so you need to "flip" the second convolution operand B vertically and horizontally using flip() .

**See Also:**

dct() , getOptimalDFTSize() , mulSpectrums(), filter2D() , matchTemplate() , flip() , cartToPolar() , magnitude() , phase()

## divide

Performs per-element division of two arrays or a scalar by an array.

**C++:** void **divide**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, double **scale**=1, int **dtype**=-1 )

**C++:** void **divide**(double **scale**, InputArray **src2**, OutputArray **dst**, int **dtype**=-1 )

**Python:** cv2.**divide**(src1, src2$\big[$, dst$\big[$, scale$\big[$, dtype $\big]\big]\big]$) $\rightarrow$ dst

**Python:** cv2.**divide**(scale, src2$\big[$, dst$\big[$, dtype $\big]\big]$) $\rightarrow$ dst

**C:** void **cvDiv**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, double **scale**=1)

**Python:** cv.**Div**(src1, src2, dst, scale=1) $\rightarrow$ None

> **Parameters**

>> **src1** – first input array.

>> **src2** – second input array of the same size and type as src1.

>> **scale** – scalar factor.

>> **dst** – output array of the same size and type as src2.

>> **dtype** – optional depth of the output array; if -1, dst will have depth src2.depth(), but in case of an array-by-array division, you can only pass -1 when src1.depth()==src2.depth().

The functions divide divide one array by another:

$$dst(I) = saturate(src1(I)*scale/src2(I))$$

or a scalar by an array when there is no src1 :

$$dst(I) = saturate(scale/src2(I))$$

When src2(I) is zero, dst(I) will also be zero. Different channels of multi-channel arrays are processed independently.

---

**Note:** Saturation is not applied when the output array has the depth CV_32S. You may even get result of an incorrect sign in the case of overflow.

---

**See Also:**

multiply(), add(), subtract(), *Matrix Expressions*

## determinant

Returns the determinant of a square floating-point matrix.

**C++:** double **determinant**(InputArray **mtx**)

**Python:** cv2.**determinant**(mtx) → retval

**C:** double **cvDet**(const CvArr* **mat**)

**Python:** cv.**Det**(mat) → float

>       **Parameters**
>
>               **mtx** – input matrix that must have CV_32FC1 or CV_64FC1 type and square size.

The function determinant calculates and returns the determinant of the specified matrix. For small matrices ( mtx.cols=mtx.rows<=3 ), the direct method is used. For larger matrices, the function uses LU factorization with partial pivoting.

For symmetric positively-determined matrices, it is also possible to use eigen() decomposition to calculate the determinant.

**See Also:**

trace(), invert(), solve(), eigen(), *Matrix Expressions*

## eigen

Calculates eigenvalues and eigenvectors of a symmetric matrix.

**C++:** bool **eigen**(InputArray **src**, OutputArray **eigenvalues**, int **lowindex**=-1, int **highindex**=-1)

**C++:** bool **eigen**(InputArray **src**, OutputArray **eigenvalues**, OutputArray **eigenvectors**, int **lowindex**=-1, int **highindex**=-1)

**Python:** cv2.**eigen**(src, computeEigenvectors[, eigenvalues[, eigenvectors]]) → retval, eigenvalues, eigenvectors

**C:** void **cvEigenVV**(CvArr* **mat**, CvArr* **evects**, CvArr* **evals**, double **eps**=0, int **lowindex**=-1, int **highindex**=-1 )

**Python:** cv.**EigenVV**(mat, evects, evals, eps, lowindex=-1, highindex=-1) → None

>       **Parameters**
>
>               **src** – input matrix that must have CV_32FC1 or CV_64FC1 type, square size and be symmetrical ($src^T == src$).
>
>               **eigenvalues** – output vector of eigenvalues of the same type as src; the eigenvalues are stored in the descending order.
>
>               **eigenvectors** – output matrix of eigenvectors; it has the same size and type as src; the eigenvectors are stored as subsequent matrix rows, in the same order as the corresponding eigenvalues.
>
>               **lowindex** – optional index of largest eigenvalue/-vector to calculate; the parameter is ignored in the current implementation.

> **highindex** – optional index of smallest eigenvalue/-vector to calculate; the parameter is ignored in the current implementation.

The functions `eigen` calculate just eigenvalues, or eigenvalues and eigenvectors of the symmetric matrix `src` :

```
src*eigenvectors.row(i).t() = eigenvalues.at<srcType>(i)*eigenvectors.row(i).t()
```

---

**Note:**  in the new and the old interfaces different ordering of eigenvalues and eigenvectors parameters is used.

---

**See Also:**

`completeSymm()` , `PCA`

## exp

Calculates the exponent of every array element.

**C++:**  void **exp**(InputArray **src**, OutputArray **dst**)

**Python:**  cv2.**exp**(src[, dst]) → dst

**C:**  void **cvExp**(const CvArr* **src**, CvArr* **dst**)

**Python:**  cv.**Exp**(src, dst) → None

>       **Parameters**

>               **src** – input array.

>               **dst** – output array of the same size and type as `src`.

The function `exp` calculates the exponent of every element of the input array:

$$\mathtt{dst}[I] = e^{src(I)}$$

The maximum relative error is about `7e-6` for single-precision input and less than `1e-10` for double-precision input. Currently, the function converts denormalized values to zeros on output. Special values (NaN, Inf) are not handled.

**See Also:**

`log()` , `cartToPolar()` , `polarToCart()` , `phase()` , `pow()` , `sqrt()` , `magnitude()`

## extractImageCOI

Extracts the selected image channel.

**C++:**  void **extractImageCOI**(const CvArr* **arr**, OutputArray **coiimg**, int **coi**=-1 )

>       **Parameters**

>               **arr** – input array; it should be a pointer to `CvMat` or `IplImage`.

>               **coiimg** – output array with a single channel and the same size and depth as `arr`.

>               **coi** – if the parameter is `>=0`, it specifies the channel to extract, if it is `<0` and `arr` is a pointer to `IplImage` with a valid COI set, the selected COI is extracted.

The function `extractImageCOI` is used to extract an image COI from an old-style array and put the result to the new-style C++ matrix. As usual, the output matrix is reallocated using `Mat::create` if needed.

To extract a channel from a new-style matrix, use `mixChannels()` or `split()` .

**See Also:**

`mixChannels()`, `split()`, `merge()`, `cvarrToMat()`, `cvSetImageCOI()`, `cvGetImageCOI()`

## insertImageCOI

Copies the selected image channel from a new-style C++ matrix to the old-style C array.

**C++:** void **insertImageCOI**(InputArray **coiimg**, CvArr* **arr**, int **coi**=-1 )

> **Parameters**
>
>> **coiimg** – input array with a single channel and the same size and depth as `arr`.
>>
>> **arr** – output array, it should be a pointer to `CvMat` or `IplImage`.
>>
>> **coi** – if the parameter is >=0, it specifies the channel to insert, if it is <0 and `arr` is a pointer to `IplImage` with a valid COI set, the selected COI is extracted.

The function `insertImageCOI` is used to extract an image COI from a new-style C++ matrix and put the result to the old-style array.

The sample below illustrates how to use the function:

```
Mat temp(240, 320, CV_8UC1, Scalar(255));
IplImage* img = cvCreateImage(cvSize(320,240), IPL_DEPTH_8U, 3);
insertImageCOI(temp, img, 1); //insert to the first channel
cvNamedWindow("window",1);
cvShowImage("window", img); //you should see green image, because channel number 1 is green (BGR)
cvWaitKey(0);
cvDestroyAllWindows();
cvReleaseImage(&img);
```

To insert a channel to a new-style matrix, use `merge()`.

**See Also:**

`mixChannels()`, `split()`, `merge()`, `cvarrToMat()`, `cvSetImageCOI()`, `cvGetImageCOI()`

## flip

Flips a 2D array around vertical, horizontal, or both axes.

**C++:** void **flip**(InputArray **src**, OutputArray **dst**, int **flipCode**)

**Python:** cv2.**flip**(src, flipCode[, dst]) → dst

**C:** void **cvFlip**(const CvArr* **src**, CvArr* **dst**=NULL, int **flip_mode**=0 )

**Python:** cv.**Flip**(src, dst=None, flipMode=0) → None

> **Parameters**
>
>> **src** – input array.
>>
>> **dst** – output array of the same size and type as `src`.
>>
>> **flipCode** – a flag to specify how to flip the array; 0 means flipping around the x-axis and positive value (for example, 1) means flipping around y-axis. Negative value (for example, -1) means flipping around both axes (see the discussion below for the formulas).

The function `flip` flips the array in one of three different ways (row and column indices are 0-based):

$$\text{dst}_{ij} = \begin{cases} \text{src}_{\text{src.rows}-i-1,j} & \text{if flipCode} = 0 \\ \text{src}_{i,\text{src.cols}-j-1} & \text{if flipCode} > 0 \\ \text{src}_{\text{src.rows}-i-1,\text{src.cols}-j-1} & \text{if flipCode} < 0 \end{cases}$$

The example scenarios of using the function are the following:

- Vertical flipping of the image (`flipCode == 0`) to switch between top-left and bottom-left image origin. This is a typical operation in video processing on Microsoft Windows* OS.

- Horizontal flipping of the image with the subsequent horizontal shift and absolute difference calculation to check for a vertical-axis symmetry (`flipCode > 0`).

- Simultaneous horizontal and vertical flipping of the image with the subsequent shift and absolute difference calculation to check for a central symmetry (`flipCode < 0`).

- Reversing the order of point arrays (`flipCode > 0` or `flipCode == 0`).

**See Also:**

`transpose()` , `repeat()` , `completeSymm()`

## gemm

Performs generalized matrix multiplication.

**C++:** void **gemm**(InputArray **src1**, InputArray **src2**, double **alpha**, InputArray **src3**, double **gamma**, OutputArray **dst**, int **flags**=0 )

**Python:** cv2.**gemm**(src1, src2, alpha, src3, gamma$\big[$, dst$\big[$, flags $\big]\big]$) $\rightarrow$ dst

**C:** void **cvGEMM**(const CvArr* **src1**, const CvArr* **src2**, double **alpha**, const CvArr* **src3**, double **beta**, CvArr* **dst**, int **tABC**=0)

**Python:** cv.**GEMM**(src1, src2, alpha, src3, beta, dst, tABC=0) $\rightarrow$ None

> Parameters
>
> > **src1** – first multiplied input matrix that should have `CV_32FC1`, `CV_64FC1`, `CV_32FC2`, or `CV_64FC2` type.
> >
> > **src2** – second multiplied input matrix of the same type as `src1`.
> >
> > **alpha** – weight of the matrix product.
> >
> > **src3** – third optional delta matrix added to the matrix product; it should have the same type as `src1` and `src2`.
> >
> > **beta** – weight of `src3`.
> >
> > **dst** – output matrix; it has the proper size and the same type as input matrices.
> >
> > **flags** – operation flags:
> >
> > – **GEMM_1_T** transposes `src1`.
> >
> > – **GEMM_2_T** transposes `src2`.
> >
> > – **GEMM_3_T** transposes `src3`.

The function performs generalized matrix multiplication similar to the `gemm` functions in BLAS level 3. For example, `gemm(src1, src2, alpha, src3, beta, dst, GEMM_1_T + GEMM_3_T)` corresponds to

$$\text{dst} = \text{alpha} \cdot \text{src1}^T \cdot \text{src2} + \text{beta} \cdot \text{src3}^T$$

The function can be replaced with a matrix expression. For example, the above call can be replaced with:

```
dst = alpha*src1.t()*src2 + beta*src3.t();
```

**See Also:**

`mulTransposed()` , `transform()` , *Matrix Expressions*

## getConvertElem

Returns a conversion function for a single pixel.

**C++:** ConvertData **getConvertElem**(int **fromType**, int **toType**)

**C++:** ConvertScaleData **getConvertScaleElem**(int **fromType**, int **toType**)

> **Parameters**
>
> > **fromType** – input pixel type.
> >
> > **toType** – output pixel type.
> >
> > **from** – callback parameter: pointer to the input pixel.
> >
> > **to** – callback parameter: pointer to the output pixel
> >
> > **cn** – callback parameter: the number of channels; it can be arbitrary, 1, 100, 100000, etc.
> >
> > **alpha** – `ConvertScaleData` callback optional parameter: the scale factor.
> >
> > **beta** – `ConvertScaleData` callback optional parameter: the delta or offset.

The functions `getConvertElem` and `getConvertScaleElem` return pointers to the functions for converting individual pixels from one type to another. While the main function purpose is to convert single pixels (actually, for converting sparse matrices from one type to another), you can use them to convert the whole row of a dense matrix or the whole matrix at once, by setting `cn = matrix.cols*matrix.rows*matrix.channels()` if the matrix data is continuous.

`ConvertData` and `ConvertScaleData` are defined as:

```
typedef void (*ConvertData)(const void* from, void* to, int cn)
typedef void (*ConvertScaleData)(const void* from, void* to,
                                 int cn, double alpha, double beta)
```

**See Also:**

`Mat::convertTo()` , `SparseMat::convertTo()`

## getOptimalDFTSize

Returns the optimal DFT size for a given vector size.

**C++:** int **getOptimalDFTSize**(int **vecsize**)

**Python:** cv2.**getOptimalDFTSize**(vecsize) → retval

**C:** int **cvGetOptimalDFTSize**(int **size0**)

**Python:** cv.**GetOptimalDFTSize**(size0) → int

> **Parameters**
>
> > **vecsize** – vector size.

DFT performance is not a monotonic function of a vector size. Therefore, when you calculate convolution of two arrays or perform the spectral analysis of an array, it usually makes sense to pad the input data with zeros to get a bit larger array that can be transformed much faster than the original one. Arrays whose size is a power-of-two (2, 4, 8, 16, 32, ...) are the fastest to process. Though, the arrays whose size is a product of 2's, 3's, and 5's (for example, 300 = 5*5*3*2*2) are also processed quite efficiently.

The function `getOptimalDFTSize` returns the minimum number `N` that is greater than or equal to `vecsize` so that the DFT of a vector of size `N` can be processed efficiently. In the current implementation $N = 2^p * 3^q * 5^r$ for some integer `p`, `q`, `r`.

The function returns a negative number if `vecsize` is too large (very close to `INT_MAX`).

While the function cannot be used directly to estimate the optimal vector size for DCT transform (since the current DCT implementation supports only even-size vectors), it can be easily processed as `getOptimalDFTSize((vecsize+1)/2)*2`.

**See Also:**

`dft()`, `dct()`, `idft()`, `idct()`, `mulSpectrums()`

## idct

Calculates the inverse Discrete Cosine Transform of a 1D or 2D array.

**C++:** void **idct**(InputArray **src**, OutputArray **dst**, int **flags**=0)

**Python:** cv2.**idct**(src[, dst[, flags ]]) → dst

> **Parameters**
>
> > **src** – input floating-point single-channel array.
> >
> > **dst** – output array of the same size and type as `src`.
> >
> > **flags** – operation flags.

`idct(src, dst, flags)` is equivalent to `dct(src, dst, flags | DCT_INVERSE)`.

**See Also:**

`dct()`, `dft()`, `idft()`, `getOptimalDFTSize()`

## idft

Calculates the inverse Discrete Fourier Transform of a 1D or 2D array.

**C++:** void **idft**(InputArray **src**, OutputArray **dst**, int **flags**=0, int **nonzeroRows**=0)

**Python:** cv2.**idft**(src[, dst[, flags[, nonzeroRows ]]]) → dst

> **Parameters**
>
> > **src** – input floating-point real or complex array.
> >
> > **dst** – output array whose size and type depend on the `flags`.
> >
> > **flags** – operation flags (see `dft()`).
> >
> > **nonzeroRows** – number of `dst` rows to process; the rest of the rows have undefined content (see the convolution sample in `dft()` description.

`idft(src, dst, flags)` is equivalent to `dft(src, dst, flags | DFT_INVERSE)`.

See `dft()` for details.

---

**Note:** None of `dft` and `idft` scales the result by default. So, you should pass `DFT_SCALE` to one of `dft` or `idft` explicitly to make these transforms mutually inverse.

---

**See Also:**

`dft()`, `dct()`, `idct()`, `mulSpectrums()`, `getOptimalDFTSize()`

## inRange

Checks if array elements lie between the elements of two other arrays.

**C++:** void **inRange**(InputArray **src**, InputArray **lowerb**, InputArray **upperb**, OutputArray **dst**)

**Python:** cv2.**inRange**(src, lowerb, upperb[, dst]) → dst

**C:** void **cvInRange**(const CvArr* **src**, const CvArr* **lower**, const CvArr* **upper**, CvArr* **dst**)

**C:** void **cvInRangeS**(const CvArr* **src**, CvScalar **lower**, CvScalar **upper**, CvArr* **dst**)

**Python:** cv.**InRange**(src, lower, upper, dst) → None

**Python:** cv.**InRangeS**(src, lower, upper, dst) → None

> **Parameters**
>
> > **src** – first input array.
> >
> > **lowerb** – inclusive lower boundary array or a scalar.
> >
> > **upperb** – inclusive upper boundary array or a scalar.
> >
> > **dst** – output array of the same size as `src` and CV_8U type.

The function checks the range as follows:

- For every element of a single-channel input array:

$$\texttt{dst}(I) = \texttt{lowerb}(I)_0 \leq \texttt{src}(I)_0 \leq \texttt{upperb}(I)_0$$

- For two-channel arrays:

$$\texttt{dst}(I) = \texttt{lowerb}(I)_0 \leq \texttt{src}(I)_0 \leq \texttt{upperb}(I)_0 \wedge \texttt{lowerb}(I)_1 \leq \texttt{src}(I)_1 \leq \texttt{upperb}(I)_1$$

- and so forth.

That is, `dst` (I) is set to 255 (all `1` -bits) if `src` (I) is within the specified 1D, 2D, 3D, ... box and 0 otherwise.

When the lower and/or upper boundary parameters are scalars, the indexes (`I`) at `lowerb` and `upperb` in the above formulas should be omitted.

## invert

Finds the inverse or pseudo-inverse of a matrix.

**C++:** double **invert**(InputArray **src**, OutputArray **dst**, int **flags**=DECOMP_LU)

**Python:** cv2.**invert**(src[, dst[, flags]]) → retval, dst

---

**C:** double **cvInvert**(const CvArr* **src**, CvArr* **dst**, int **method**=CV_LU )

**Python:** cv.**Invert**(src, dst, method=CV_LU) → float

> **Parameters**

>> **src** – input floating-point M x N matrix.

>> **dst** – output matrix of N x M size and the same type as src.

>> **flags** – inversion method :

>>> – **DECOMP_LU** Gaussian elimination with the optimal pivot element chosen.

>>> – **DECOMP_SVD** singular value decomposition (SVD) method.

>>> – **DECOMP_CHOLESKY** Cholesky decomposition; the matrix must be symmetrical and positively defined.

The function invert inverts the matrix src and stores the result in dst . When the matrix src is singular or non-square, the function calculates the pseudo-inverse matrix (the dst matrix) so that norm(src*dst - I) is minimal, where I is an identity matrix.

In case of the DECOMP_LU method, the function returns non-zero value if the inverse has been successfully calculated and 0 if src is singular.

In case of the DECOMP_SVD method, the function returns the inverse condition number of src (the ratio of the smallest singular value to the largest singular value) and 0 if src is singular. The SVD method calculates a pseudo-inverse matrix if src is singular.

Similarly to DECOMP_LU , the method DECOMP_CHOLESKY works only with non-singular square matrices that should also be symmetrical and positively defined. In this case, the function stores the inverted matrix in dst and returns non-zero. Otherwise, it returns 0.

**See Also:**

solve(), SVD

## log

Calculates the natural logarithm of every array element.

**C++:** void **log**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**log**(src[, dst]) → dst

**C:** void **cvLog**(const CvArr* **src**, CvArr* **dst**)

**Python:** cv.**Log**(src, dst) → None

> **Parameters**

>> **src** – input array.

>> **dst** – output array of the same size and type as src .

The function log calculates the natural logarithm of the absolute value of every element of the input array:

$$\texttt{dst}(I) = \begin{cases} \log|\texttt{src}(I)| & \text{if } \texttt{src}(I) \neq 0 \\ \texttt{C} & \text{otherwise} \end{cases}$$

where C is a large negative number (about -700 in the current implementation). The maximum relative error is about 7e-6 for single-precision input and less than 1e-10 for double-precision input. Special values (NaN, Inf) are not handled.

**See Also:**

exp(), cartToPolar(), polarToCart(), phase(), pow(), sqrt(), magnitude()

## LUT

Performs a look-up table transform of an array.

**C++:** void **LUT**(InputArray **src**, InputArray **lut**, OutputArray **dst**, int **interpolation**=0 )

**Python:** cv2.**LUT**(src, lut[, dst[, interpolation ]]) → dst

**C:** void **cvLUT**(const CvArr* **src**, CvArr* **dst**, const CvArr* **lut**)

**Python:** cv.**LUT**(src, dst, lut) → None

> **Parameters**
>
>> **src** – input array of 8-bit elements.
>>
>> **lut** – look-up table of 256 elements; in case of multi-channel input array, the table should either have a single channel (in this case the same table is used for all channels) or the same number of channels as in the input array.
>>
>> **dst** – output array of the same size and number of channels as src, and the same depth as lut.

The function LUT fills the output array with values from the look-up table. Indices of the entries are taken from the input array. That is, the function processes each element of src as follows:

$$\texttt{dst}(I) \leftarrow \texttt{lut}(\texttt{src}(I) + d)$$

where

$$d = \begin{cases} 0 & \text{if } \texttt{src} \text{ has depth } \texttt{CV\_8U} \\ 128 & \text{if } \texttt{src} \text{ has depth } \texttt{CV\_8S} \end{cases}$$

**See Also:**

convertScaleAbs(), Mat::convertTo()

## magnitude

Calculates the magnitude of 2D vectors.

**C++:** void **magnitude**(InputArray **x**, InputArray **y**, OutputArray **magnitude**)

**Python:** cv2.**magnitude**(x, y[, magnitude ]) → magnitude

> **Parameters**
>
>> **x** – floating-point array of x-coordinates of the vectors.
>>
>> **y** – floating-point array of y-coordinates of the vectors; it must have the same size as x.
>>
>> **magnitude** – output array of the same size and type as x.

The function magnitude calculates the magnitude of 2D vectors formed from the corresponding elements of x and y arrays:

$$\texttt{dst}(I) = \sqrt{\texttt{x}(I)^2 + \texttt{y}(I)^2}$$

**See Also:**

cartToPolar(), polarToCart(), phase(), sqrt()

## Mahalanobis

Calculates the Mahalanobis distance between two vectors.

**C++:** double **Mahalanobis**(InputArray **v1**, InputArray **v2**, InputArray **icovar**)

**Python:** cv2.**Mahalanobis**(v1, v2, icovar) → retval

**C:** double **cvMahalanobis**(const CvArr* **vec1**, const CvArr* **vec2**, const CvArr* **mat**)

**Python:** cv.**Mahalonobis**(vec1, vec2, mat) → None

> **Parameters**
>
> > **vec1** – first 1D input vector.
> >
> > **vec2** – second 1D input vector.
> >
> > **icovar** – inverse covariance matrix.

The function Mahalanobis calculates and returns the weighted distance between two vectors:

$$d(\mathtt{vec1}, \mathtt{vec2}) = \sqrt{\sum_{i,j} \mathtt{icovar(i,j)} \cdot (\mathtt{vec1(I)} - \mathtt{vec2(I)}) \cdot (\mathtt{vec1(j)} - \mathtt{vec2(j)})}$$

The covariance matrix may be calculated using the calcCovarMatrix() function and then inverted using the invert() function (preferably using the DECOMP_SVD method, as the most accurate).

## max

Calculates per-element maximum of two arrays or an array and a scalar.

**C++:** MatExpr **max**(const Mat& **a**, const Mat& **b**)

**C++:** MatExpr **max**(const Mat& **a**, double **s**)

**C++:** MatExpr **max**(double **s**, const Mat& **a**)

**C++:** void **max**(InputArray **src1**, InputArray **src2**, OutputArray **dst**)

**C++:** void **max**(const Mat& **src1**, const Mat& **src2**, Mat& **dst**)

**C++:** void **max**(const Mat& **src1**, double **src2**, Mat& **dst**)

**Python:** cv2.**max**(src1, src2[, dst]) → dst

**C:** void **cvMax**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**)

**C:** void **cvMaxS**(const CvArr* **src**, double **value**, CvArr* **dst**)

**Python:** cv.**Max**(src1, src2, dst) → None

**Python:** cv.**MaxS**(src, value, dst) → None

> **Parameters**
>
> > **src1** – first input array.
> >
> > **src2** – second input array of the same size and type as src1 .
> >
> > **value** – real scalar value.

> **dst** – output array of the same size and type as `src1`.

The functions max calculate the per-element maximum of two arrays:

$$\mathtt{dst}(I) = \max(\mathtt{src1}(I), \mathtt{src2}(I))$$

or array and a scalar:

$$\mathtt{dst}(I) = \max(\mathtt{src1}(I), \mathtt{value})$$

In the second variant, when the input array is multi-channel, each channel is compared with `value` independently.

The first 3 variants of the function listed above are actually a part of *Matrix Expressions* . They return an expression object that can be further either transformed/ assigned to a matrix, or passed to a function, and so on.

**See Also:**

min(), compare(), inRange(), minMaxLoc(), *Matrix Expressions*

## mean

Calculates an average (mean) of array elements.

**C++:** Scalar **mean**(InputArray **src**, InputArray **mask**=noArray())

**Python:** cv2.**mean**(src[, mask]) → retval

**C:** CvScalar **cvAvg**(const CvArr* **arr**, const CvArr* **mask**=NULL )

**Python:** cv.**Avg**(arr, mask=None) → scalar

> **Parameters**
>
> > **src** – input array that should have from 1 to 4 channels so that the result can be stored in Scalar_ .
> >
> > **mask** – optional operation mask.

The function mean calculates the mean value M of array elements, independently for each channel, and return it:

$$\begin{aligned} N &= \sum_{I: \, \mathtt{mask}(I) \neq 0} 1 \\ M_c &= \left( \sum_{I: \, \mathtt{mask}(I) \neq 0} \mathtt{mtx}(I)_c \right) / N \end{aligned}$$

When all the mask elements are 0's, the functions return Scalar::all(0) .

**See Also:**

countNonZero(), meanStdDev(), norm(), minMaxLoc()

## meanStdDev

Calculates a mean and standard deviation of array elements.

**C++:** void **meanStdDev**(InputArray **src**, OutputArray **mean**, OutputArray **stddev**, InputArray **mask**=noArray())

**Python:** cv2.**meanStdDev**(src[, mean[, stddev[, mask]]]) → mean, stddev

**C:** void **cvAvgSdv**(const CvArr* **arr**, CvScalar* **mean**, CvScalar* **std_dev**, const CvArr* **mask**=NULL )

**Python:** cv.**AvgSdv**(arr, mask=None) -> (mean, stdDev)

> **Parameters**

>>> **src** – input array that should have from 1 to 4 channels so that the results can be stored in `Scalar_` 's.

>>> **mean** – output parameter: calculated mean value.

>>> **stddev** – output parameter: calculateded standard deviation.

>>> **mask** – optional operation mask.

The function `meanStdDev` calculates the mean and the standard deviation `M` of array elements independently for each channel and returns it via the output parameters:

$$N = \sum_{I,\text{mask}(I)\neq 0} 1$$
$$\text{mean}_c = \frac{\sum_{I:\ \text{mask}(I)\neq 0} \text{src}(I)_c}{N}$$
$$\text{stddev}_c = \sqrt{\frac{\sum_{I:\ \text{mask}(I)\neq 0} (\text{src}(I)_c - \text{mean}_c)^2}{N}}$$

When all the mask elements are 0's, the functions return `mean=stddev=Scalar::all(0)`.

**Note:** The calculated standard deviation is only the diagonal of the complete normalized covariance matrix. If the full matrix is needed, you can reshape the multi-channel array `M x N` to the single-channel array `M*N x mtx.channels()` (only possible when the matrix is continuous) and then pass the matrix to `calcCovarMatrix()`.

**See Also:**

`countNonZero()`, `mean()`, `norm()`, `minMaxLoc()`, `calcCovarMatrix()`

## merge

Creates one multichannel array out of several single-channel ones.

**C++:** void **merge**(const Mat* **mv**, size_t **count**, OutputArray **dst**)

**C++:** void **merge**(InputArrayOfArrays **mv**, OutputArray **dst**)

**Python:** cv2.**merge**(mv[, dst]) → dst

**C:** void **cvMerge**(const CvArr* **src0**, const CvArr* **src1**, const CvArr* **src2**, const CvArr* **src3**, CvArr* **dst**)

**Python:** cv.**Merge**(src0, src1, src2, src3, dst) → None

>>> **Parameters**

>>>> **mv** – input array or vector of matrices to be merged; all the matrices in `mv` must have the same size and the same depth.

>>>> **count** – number of input matrices when `mv` is a plain C array; it must be greater than zero.

>>>> **dst** – output array of the same size and the same depth as `mv[0]`; The number of channels will be the total number of channels in the matrix array.

The functions `merge` merge several arrays to make a single multi-channel array. That is, each element of the output array will be a concatenation of the elements of the input arrays, where elements of i-th input array are treated as `mv[i].channels()`-element vectors.

The function `split()` does the reverse operation. If you need to shuffle channels in some other advanced way, use `mixChannels()`.

**See Also:**

`mixChannels()`, `split()`, `Mat::reshape()`

---

## min

Calculates per-element minimum of two arrays or an array and a scalar.

**C++:** MatExpr **min**(const Mat& **a**, const Mat& **b**)

**C++:** MatExpr **min**(const Mat& **a**, double **s**)

**C++:** MatExpr **min**(double **s**, const Mat& **a**)

**C++:** void **min**(InputArray **src1**, InputArray **src2**, OutputArray **dst**)

**C++:** void **min**(const Mat& **src1**, const Mat& **src2**, Mat& **dst**)

**C++:** void **min**(const Mat& **src1**, double **src2**, Mat& **dst**)

**Python:** cv2.**min**(src1, src2[, dst]) → dst

**C:** void **cvMin**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**)

**C:** void **cvMinS**(const CvArr* **src**, double **value**, CvArr* **dst**)

**Python:** cv.**Min**(src1, src2, dst) → None

**Python:** cv.**MinS**(src, value, dst) → None

> **Parameters**

>> **src1** – first input array.

>> **src2** – second input array of the same size and type as src1.

>> **value** – real scalar value.

>> **dst** – output array of the same size and type as src1.

The functions min calculate the per-element minimum of two arrays:

$$\texttt{dst}(I) = \min(\texttt{src1}(I), \texttt{src2}(I))$$

or array and a scalar:

$$\texttt{dst}(I) = \min(\texttt{src1}(I), \texttt{value})$$

In the second variant, when the input array is multi-channel, each channel is compared with value independently.

The first three variants of the function listed above are actually a part of *Matrix Expressions* . They return the expression object that can be further either transformed/assigned to a matrix, or passed to a function, and so on.

**See Also:**

max(), compare(), inRange(), minMaxLoc(), *Matrix Expressions*

## minMaxIdx

Finds the global minimum and maximum in an array

**C++:** void **minMaxIdx**(InputArray **src**, double* **minVal**, double* **maxVal**, int* **minIdx**=0, int* **maxIdx**=0, InputArray **mask**=noArray())

> **Parameters**

>> **src** – input single-channel array.

>> **minVal** – pointer to the returned minimum value; NULL is used if not required.

>> **maxVal** – pointer to the returned maximum value; NULL is used if not required.

> **minIdx** – pointer to the returned minimum location (in nD case); `NULL` is used if not required; Otherwise, it must point to an array of `src.dims` elements, the coordinates of the minimum element in each dimension are stored there sequentially.

---

> **Note:** When `minIdx` is not NULL, it must have at least 2 elements (as well as `maxIdx`), even if `src` is a single-row or single-column matrix. In OpenCV (following MATLAB) each array has at least 2 dimensions, i.e. single-column matrix is `Mx1` matrix (and therefore `minIdx`/`maxIdx` will be `(i1,0)`/`(i2,0)`) and single-row matrix is `1xN` matrix (and therefore `minIdx`/`maxIdx` will be `(0,j1)`/`(0,j2)`).

---

> **maxIdx** – pointer to the returned maximum location (in nD case). `NULL` is used if not required.

The function `minMaxIdx` finds the minimum and maximum element values and their positions. The extremums are searched across the whole array or, if `mask` is not an empty array, in the specified array region.

The function does not work with multi-channel arrays. If you need to find minimum or maximum elements across all the channels, use `Mat::reshape()` first to reinterpret the array as single-channel. Or you may extract the particular channel using either `extractImageCOI()`, or `mixChannels()`, or `split()`.

In case of a sparse matrix, the minimum is found among non-zero elements only.

## minMaxLoc

Finds the global minimum and maximum in an array.

**C++:** void **minMaxLoc**(InputArray **src**, double* **minVal**, double* **maxVal**=0, Point* **minLoc**=0, Point* **maxLoc**=0, InputArray **mask**=noArray())

**C++:** void **minMaxLoc**(const SparseMat& **a**, double* **minVal**, double* **maxVal**, int* **minIdx**=0, int* **maxIdx**=0 )

**Python:** cv2.**minMaxLoc**(src[, mask ]) → minVal, maxVal, minLoc, maxLoc

**C:** void **cvMinMaxLoc**(const CvArr* **arr**, double* **min_val**, double* **max_val**, CvPoint* **min_loc**=NULL, CvPoint* **max_loc**=NULL, const CvArr* **mask**=NULL )

**Python:** cv.**MinMaxLoc**(arr, mask=None)-> (minVal, maxVal, minLoc, maxLoc)

> **Parameters**

>> **src** – input single-channel array.

>> **minVal** – pointer to the returned minimum value; `NULL` is used if not required.

>> **maxVal** – pointer to the returned maximum value; `NULL` is used if not required.

>> **minLoc** – pointer to the returned minimum location (in 2D case); `NULL` is used if not required.

>> **maxLoc** – pointer to the returned maximum location (in 2D case); `NULL` is used if not required.

>> **mask** – optional mask used to select a sub-array.

The functions `minMaxLoc` find the minimum and maximum element values and their positions. The extremums are searched across the whole array or, if `mask` is not an empty array, in the specified array region.

The functions do not work with multi-channel arrays. If you need to find minimum or maximum elements across all the channels, use `Mat::reshape()` first to reinterpret the array as single-channel. Or you may extract the particular channel using either `extractImageCOI()`, or `mixChannels()`, or `split()`.

---

**See Also:**

max(), min(), compare(), inRange(), extractImageCOI(), mixChannels(), split(), Mat::reshape()

## mixChannels

Copies specified channels from input arrays to the specified channels of output arrays.

**C++:** void **mixChannels**(const Mat* **src**, size_t **nsrcs**, Mat* **dst**, size_t **ndsts**, const int* **fromTo**, size_t **npairs**)

**C++:** void **mixChannels**(const vector<Mat>& **src**, vector<Mat>& **dst**, const int* **fromTo**, size_t **npairs**)

**Python:** cv2.**mixChannels**(src, dst, fromTo) → None

**C:** void **cvMixChannels**(const CvArr** **src**, int **src_count**, CvArr** **dst**, int **dst_count**, const int* **from_to**, int **pair_count**)

**Python:** cv.**MixChannels**(src, dst, fromTo) → None

> **Parameters**

>> **src** – input array or vector of matricesl; all of the matrices must have the same size and the same depth.

>> **nsrcs** – number of matrices in src.

>> **dst** – output array or vector of matrices; all the matrices *must be allocated*; their size and depth must be the same as in src[0].

>> **ndsts** – number of matrices in dst.

>> **fromTo** – array of index pairs specifying which channels are copied and where; fromTo[k*2] is a 0-based index of the input channel in src, fromTo[k*2+1] is an index of the output channel in dst; the continuous channel numbering is used: the first input image channels are indexed from 0 to src[0].channels()-1, the second input image channels are indexed from src[0].channels() to src[0].channels() + src[1].channels()-1, and so on, the same scheme is used for the output image channels; as a special case, when fromTo[k*2] is negative, the corresponding output channel is filled with zero .

>> **npairs** – number of index pairs in fromTo.

The functions mixChannels provide an advanced mechanism for shuffling image channels.

split() and merge() and some forms of cvtColor() are partial cases of mixChannels .

In the example below, the code splits a 4-channel RGBA image into a 3-channel BGR (with R and B channels swapped) and a separate alpha-channel image:

```
Mat rgba( 100, 100, CV_8UC4, Scalar(1,2,3,4) );
Mat bgr( rgba.rows, rgba.cols, CV_8UC3 );
Mat alpha( rgba.rows, rgba.cols, CV_8UC1 );

// forming an array of matrices is a quite efficient operation,
// because the matrix data is not copied, only the headers
Mat out[] = { bgr, alpha };
// rgba[0] -> bgr[2], rgba[1] -> bgr[1],
// rgba[2] -> bgr[0], rgba[3] -> alpha[0]
int from_to[] = { 0,2, 1,1, 2,0, 3,3 };
mixChannels( &rgba, 1, out, 2, from_to, 4 );
```

---

**Note:** Unlike many other new-style C++ functions in OpenCV (see the introduction section and `Mat::create()` ), `mixChannels` requires the output arrays to be pre-allocated before calling the function.

---

**See Also:**

`split()`, `merge()`, `cvtColor()`

## mulSpectrums

Performs the per-element multiplication of two Fourier spectrums.

**C++:** void **mulSpectrums** (InputArray **a**, InputArray **b**, OutputArray **c**, int **flags**, bool **conjB**=false )

**Python:** cv2.**mulSpectrums** (a, b, flags$\big[$, c$\big[$, conjB $\big]\big]$) $\rightarrow$ c

**C:** void **cvMulSpectrums** (const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, int **flags**)

**Python:** cv.**MulSpectrums** (src1, src2, dst, flags) $\rightarrow$ None

>    **Parameters**

>    >    **src1** – first input array.

>    >    **src2** – second input array of the same size and type as `src1` .

>    >    **dst** – output array of the same size and type as `src1` .

>    >    **flags** – operation flags; currently, the only supported flag is `DFT_ROWS`, which indicates that each row of `src1` and `src2` is an independent 1D Fourier spectrum.

>    >    **conjB** – optional flag that conjugates the second input array before the multiplication (true) or not (false).

The function `mulSpectrums` performs the per-element multiplication of the two CCS-packed or complex matrices that are results of a real or complex Fourier transform.

The function, together with `dft()` and `idft()` , may be used to calculate convolution (pass `conjB=false` ) or correlation (pass `conjB=true` ) of two arrays rapidly. When the arrays are complex, they are simply multiplied (per element) with an optional conjugation of the second-array elements. When the arrays are real, they are assumed to be CCS-packed (see `dft()` for details).

## multiply

Calculates the per-element scaled product of two arrays.

**C++:** void **multiply** (InputArray **src1**, InputArray **src2**, OutputArray **dst**, double **scale**=1, int **dtype**=-1 )

**Python:** cv2.**multiply** (src1, src2$\big[$, dst$\big[$, scale$\big[$, dtype $\big]\big]\big]$) $\rightarrow$ dst

**C:** void **cvMul** (const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, double **scale**=1)

**Python:** cv.**Mul** (src1, src2, dst, scale=1) $\rightarrow$ None

>    **Parameters**

>    >    **src1** – first input array.

>    >    **src2** – second input array of the same size and the same type as `src1`.

>    >    **dst** – output array of the same size and type as `src1`.

>    >    **scale** – optional scale factor.

---

The function `multiply` calculates the per-element product of two arrays:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{scale} \cdot \texttt{src1}(I) \cdot \texttt{src2}(I))$$

There is also a *Matrix Expressions* -friendly variant of the first function. See `Mat::mul()` .

For a not-per-element matrix product, see `gemm()` .

---

**Note:** Saturation is not applied when the output array has the depth `CV_32S`. You may even get result of an incorrect sign in the case of overflow.

---

**See Also:**

`add()`, `subtract()`, `divide()`, *Matrix Expressions*, `scaleAdd()`, `addWeighted()`, `accumulate()`, `accumulateProduct()`, `accumulateSquare()`, `Mat::convertTo()`

## mulTransposed

Calculates the product of a matrix and its transposition.

**C++:** void **mulTransposed**(InputArray **src**, OutputArray **dst**, bool **aTa**, InputArray **delta**=noArray(), double **scale**=1, int **dtype**=-1 )

**Python:** cv2.**mulTransposed**(src, aTa[, dst[, delta[, scale[, dtype ]]]]) → dst

**C:** void **cvMulTransposed**(const CvArr* **src**, CvArr* **dst**, int **order**, const CvArr* **delta**=NULL, double **scale**=1. )

**Python:** cv.**MulTransposed**(src, dst, order, delta=None, scale=1.0) → None

> **Parameters**

>> **src** – input single-channel matrix. Note that unlike `gemm()`, the function can multiply not only floating-point matrices.

>> **dst** – output square matrix.

>> **aTa** – Flag specifying the multiplication ordering. See the description below.

>> **delta** – Optional delta matrix subtracted from `src` before the multiplication. When the matrix is empty ( `delta=noArray()` ), it is assumed to be zero, that is, nothing is subtracted. If it has the same size as `src` , it is simply subtracted. Otherwise, it is "repeated" (see `repeat()` ) to cover the full `src` and then subtracted. Type of the delta matrix, when it is not empty, must be the same as the type of created output matrix. See the `dtype` parameter description below.

>> **scale** – Optional scale factor for the matrix product.

>> **dtype** – Optional type of the output matrix. When it is negative, the output matrix will have the same type as `src` . Otherwise, it will be `type=CV_MAT_DEPTH(dtype)` that should be either `CV_32F` or `CV_64F` .

The function `mulTransposed` calculates the product of `src` and its transposition:

$$\texttt{dst} = \texttt{scale}(\texttt{src} - \texttt{delta})^T(\texttt{src} - \texttt{delta})$$

if `aTa=true` , and

$$\texttt{dst} = \texttt{scale}(\texttt{src} - \texttt{delta})(\texttt{src} - \texttt{delta})^T$$

otherwise. The function is used to calculate the covariance matrix. With zero delta, it can be used as a faster substitute for general matrix product A*B when B=A'

---

**See Also:**

`calcCovarMatrix()`, `gemm()`, `repeat()`, `reduce()`

## norm

Calculates an absolute array norm, an absolute difference norm, or a relative difference norm.

**C++:** double **norm**(InputArray **src1**, int **normType**=NORM_L2, InputArray **mask**=noArray())

**C++:** double **norm**(InputArray **src1**, InputArray **src2**, int **normType**=NORM_L2, InputArray **mask**=noArray() )

**C++:** double **norm**(const SparseMat& **src**, int **normType**)

**Python:** cv2.**norm**(src1[, normType[, mask ]]) → retval

**Python:** cv2.**norm**(src1, src2[, normType[, mask ]]) → retval

**C:** double **cvNorm**(const CvArr* **arr1**, const CvArr* **arr2**=NULL, int **norm_type**=CV_L2, const CvArr* **mask**=NULL )

**Python:** cv.**Norm**(arr1, arr2, normType=CV_L2, mask=None) → float

> Parameters
>
> > **src1** – first input array.
> >
> > **src2** – second input array of the same size and the same type as `src1`.
> >
> > **normType** – type of the norm (see the details below).
> >
> > **mask** – optional operation mask; it must have the same size as `src1` and `CV_8UC1` type.

The functions `norm` calculate an absolute norm of `src1` (when there is no `src2` ):

$$
\mathrm{norm} = \begin{cases} \|\texttt{src1}\|_{L_\infty} = \max_I |\texttt{src1}(I)| & \text{if normType} = \texttt{NORM\_INF} \\ \|\texttt{src1}\|_{L_1} = \sum_I |\texttt{src1}(I)| & \text{if normType} = \texttt{NORM\_L1} \\ \|\texttt{src1}\|_{L_2} = \sqrt{\sum_I \texttt{src1}(I)^2} & \text{if normType} = \texttt{NORM\_L2} \end{cases}
$$

or an absolute or relative difference norm if `src2` is there:

$$
\mathrm{norm} = \begin{cases} \|\texttt{src1} - \texttt{src2}\|_{L_\infty} = \max_I |\texttt{src1}(I) - \texttt{src2}(I)| & \text{if normType} = \texttt{NORM\_INF} \\ \|\texttt{src1} - \texttt{src2}\|_{L_1} = \sum_I |\texttt{src1}(I) - \texttt{src2}(I)| & \text{if normType} = \texttt{NORM\_L1} \\ \|\texttt{src1} - \texttt{src2}\|_{L_2} = \sqrt{\sum_I (\texttt{src1}(I) - \texttt{src2}(I))^2} & \text{if normType} = \texttt{NORM\_L2} \end{cases}
$$

or

$$
\mathrm{norm} = \begin{cases} \frac{\|\texttt{src1} - \texttt{src2}\|_{L_\infty}}{\|\texttt{src2}\|_{L_\infty}} & \text{if normType} = \texttt{NORM\_RELATIVE\_INF} \\ \frac{\|\texttt{src1} - \texttt{src2}\|_{L_1}}{\|\texttt{src2}\|_{L_1}} & \text{if normType} = \texttt{NORM\_RELATIVE\_L1} \\ \frac{\|\texttt{src1} - \texttt{src2}\|_{L_2}}{\|\texttt{src2}\|_{L_2}} & \text{if normType} = \texttt{NORM\_RELATIVE\_L2} \end{cases}
$$

The functions `norm` return the calculated norm.

When the `mask` parameter is specified and it is not empty, the norm is calculated only over the region specified by the mask.

A multi-channel input arrays are treated as a single-channel, that is, the results for all channels are combined.

## normalize

Normalizes the norm or value range of an array.

**C++:** void **normalize**(InputArray **src**, OutputArray **dst**, double **alpha**=1, double **beta**=0, int **norm_type**=NORM_L2, int **dtype**=-1, InputArray **mask**=noArray() )

**C++:** void **normalize**(const SparseMat& **src**, SparseMat& **dst**, double **alpha**, int **normType**)

**Python:** cv2.**normalize**(src[, dst[, alpha[, beta[, norm_type[, dtype[, mask ]]]]]]) → dst

>   **Parameters**

>>      **src** – input array.

>>      **dst** – output array of the same size as `src`.

>>      **alpha** – norm value to normalize to or the lower range boundary in case of the range normalization.

>>      **beta** – upper range boundary in case of the range normalization; it is not used for the norm normalization.

>>      **normType** – normalization type (see the details below).

>>      **dtype** – when negative, the output array has the same type as `src`; otherwise, it has the same number of channels as `src` and the depth =CV_MAT_DEPTH(dtype).

>>      **mask** – optional operation mask.

The functions `normalize` scale and shift the input array elements so that

$$\|\mathtt{dst}\|_{L_p} = \mathtt{alpha}$$

(where p=Inf, 1 or 2) when `normType`=NORM_INF, NORM_L1, or NORM_L2, respectively; or so that

$$\min_{I} \mathtt{dst}(I) = \mathtt{alpha}, \; \max_{I} \mathtt{dst}(I) = \mathtt{beta}$$

when `normType`=NORM_MINMAX (for dense arrays only). The optional mask specifies a sub-array to be normalized. This means that the norm or min-n-max are calculated over the sub-array, and then this sub-array is modified to be normalized. If you want to only use the mask to calculate the norm or min-max but modify the whole array, you can use `norm()` and `Mat::convertTo()`.

In case of sparse matrices, only the non-zero values are analyzed and transformed. Because of this, the range transformation for sparse matrices is not allowed since it can shift the zero level.

**See Also:**

`norm()`, `Mat::convertTo()`, `SparseMat::convertTo()`

## PCA

**class PCA**

Principal Component Analysis class.

The class is used to calculate a special basis for a set of vectors. The basis will consist of eigenvectors of the covariance matrix calculated from the input set of vectors. The class `PCA` can also transform vectors to/from the new coordinate space defined by the basis. Usually, in this new coordinate system, each vector from the original set (and any linear combination of such vectors) can be quite accurately approximated by taking its first few components, corresponding to the eigenvectors of the largest eigenvalues of the covariance matrix. Geometrically it means that you calculate a projection of the vector to a subspace formed by a few eigenvectors corresponding to the dominant

eigenvalues of the covariance matrix. And usually such a projection is very close to the original vector. So, you can represent the original vector from a high-dimensional space with a much shorter vector consisting of the projected vector's coordinates in the subspace. Such a transformation is also known as Karhunen-Loeve Transform, or KLT. See http://en.wikipedia.org/wiki/Principal_component_analysis .

The sample below is the function that takes two matrices. The first function stores a set of vectors (a row per vector) that is used to calculate PCA. The second function stores another "test" set of vectors (a row per vector). First, these vectors are compressed with PCA, then reconstructed back, and then the reconstruction error norm is computed and printed for each vector.

```cpp
PCA compressPCA(InputArray pcaset, int maxComponents,
                const Mat& testset, OutputArray compressed)
{
    PCA pca(pcaset, // pass the data
            Mat(), // there is no pre-computed mean vector,
                   // so let the PCA engine to compute it
            CV_PCA_DATA_AS_ROW, // indicate that the vectors
                                // are stored as matrix rows
                                // (use CV_PCA_DATA_AS_COL if the vectors are
                                // the matrix columns)
            maxComponents // specify how many principal components to retain
            );
    // if there is no test data, just return the computed basis, ready-to-use
    if( !testset.data )
        return pca;
    CV_Assert( testset.cols == pcaset.cols );

    compressed.create(testset.rows, maxComponents, testset.type());

    Mat reconstructed;
    for( int i = 0; i < testset.rows; i++ )
    {
        Mat vec = testset.row(i), coeffs = compressed.row(i);
        // compress the vector, the result will be stored
        // in the i-th row of the output matrix
        pca.project(vec, coeffs);
        // and then reconstruct it
        pca.backProject(coeffs, reconstructed);
        // and measure the error
        printf("%d. diff = %g\n", i, norm(vec, reconstructed, NORM_L2));
    }
    return pca;
}
```

**See Also:**

calcCovarMatrix(), mulTransposed(), SVD, dft(), dct()

## PCA::PCA

PCA constructors

**C++:** PCA::**PCA**()

**C++:** PCA::**PCA**(InputArray **data**, InputArray **mean**, int **flags**, int **maxComponents**=0)

**C++:** PCA::**PCA**(InputArray **data**, InputArray **mean**, int **flags**, double **retainedVariance**)

    **Parameters**

> > **data** – input samples stored as matrix rows or matrix columns.
>
> > **mean** – optional mean value; if the matrix is empty (`noArray()`), the mean is computed from the data.
>
> > **flags** – operation flags; currently the parameter is only used to specify the data layout:
>
> > > – **CV_PCA_DATA_AS_ROW** indicates that the input samples are stored as matrix rows.
> >
> > > – **CV_PCA_DATA_AS_COL** indicates that the input samples are stored as matrix columns.
>
> > **maxComponents** – maximum number of components that PCA should retain; by default, all the components are retained.
>
> > **retainedVariance** – Percentage of variance that PCA should retain. Using this parameter will let the PCA decided how many components to retain but it will always keep at least 2.

The default constructor initializes an empty PCA structure. The other constructors initialize the structure and call `PCA::operator()`.

## PCA::operator ()

Performs Principal Component Analysis of the supplied dataset.

**C++:** PCA& PCA::**operator()**(InputArray **data**, InputArray **mean**, int **flags**, int **maxComponents**=0)

**C++:** PCA& PCA::**computeVar**(InputArray **data**, InputArray **mean**, int **flags**, double **retainedVariance**)

**Python:** cv2.**PCACompute**(data[, mean[, eigenvectors[, maxComponents ] ] ]) → mean, eigenvectors

**Python:** cv2.**PCAComputeVar**(data, retainedVariance[, mean[, eigenvectors ] ]) → mean, eigenvectors

> > **Parameters**

> > > **data** – input samples stored as the matrix rows or as the matrix columns.
> >
> > > **mean** – optional mean value; if the matrix is empty (`noArray()`), the mean is computed from the data.
> >
> > > **flags** – operation flags; currently the parameter is only used to specify the data layout.
> >
> > > > – **CV_PCA_DATA_AS_ROW** indicates that the input samples are stored as matrix rows.
> > >
> > > > – **CV_PCA_DATA_AS_COL** indicates that the input samples are stored as matrix columns.
> >
> > > **maxComponents** – maximum number of components that PCA should retain; by default, all the components are retained.
> >
> > > **retainedVariance** – Percentage of variance that PCA should retain. Using this parameter will let the PCA decided how many components to retain but it will always keep at least 2.

The operator performs PCA of the supplied dataset. It is safe to reuse the same PCA structure for multiple datasets. That is, if the structure has been previously used with another dataset, the existing internal data is reclaimed and the new `eigenvalues`, `eigenvectors`, and `mean` are allocated and computed.

The computed eigenvalues are sorted from the largest to the smallest and the corresponding eigenvectors are stored as `PCA::eigenvectors` rows.

## PCA::project

Projects vector(s) to the principal component subspace.

**C++:** Mat PCA::**project**(InputArray **vec**) const

**C++:** void PCA::**project**(InputArray **vec**, OutputArray **result**) const

**Python:** cv2.**PCAProject**(data, mean, eigenvectors[, result]) → result

>        **Parameters**

>> **vec** – input vector(s); must have the same dimensionality and the same layout as the input data used at PCA phase, that is, if CV_PCA_DATA_AS_ROW are specified, then vec.cols==data.cols (vector dimensionality) and vec.rows is the number of vectors to project, and the same is true for the CV_PCA_DATA_AS_COL case.

>> **result** – output vectors; in case of CV_PCA_DATA_AS_COL, the output matrix has as many columns as the number of input vectors, this means that result.cols==vec.cols and the number of rows match the number of principal components (for example, maxComponents parameter passed to the constructor).

The methods project one or more vectors to the principal component subspace, where each vector projection is represented by coefficients in the principal component basis. The first form of the method returns the matrix that the second form writes to the result. So the first form can be used as a part of expression while the second form can be more efficient in a processing loop.

## PCA::backProject

Reconstructs vectors from their PC projections.

**C++:** Mat PCA::**backProject**(InputArray **vec**) const

**C++:** void PCA::**backProject**(InputArray **vec**, OutputArray **result**) const

**Python:** cv2.**PCABackProject**(data, mean, eigenvectors[, result]) → result

>        **Parameters**

>> **vec** – coordinates of the vectors in the principal component subspace, the layout and size are the same as of PCA::project output vectors.

>> **result** – reconstructed vectors; the layout and size are the same as of PCA::project input vectors.

The methods are inverse operations to PCA::project(). They take PC coordinates of projected vectors and reconstruct the original vectors. Unless all the principal components have been retained, the reconstructed vectors are different from the originals. But typically, the difference is small if the number of components is large enough (but still much smaller than the original vector dimensionality). As a result, PCA is used.

## perspectiveTransform

Performs the perspective matrix transformation of vectors.

**C++:** void **perspectiveTransform**(InputArray **src**, OutputArray **dst**, InputArray **m**)

**Python:** cv2.**perspectiveTransform**(src, m[, dst]) → dst

**C:** void **cvPerspectiveTransform**(const CvArr* **src**, CvArr* **dst**, const CvMat* **mat**)

**Python:** cv.**PerspectiveTransform**(src, dst, mat) → None

---

**Parameters**

**src** – input two-channel or three-channel floating-point array; each element is a 2D/3D vector to be transformed.

**dst** – output array of the same size and type as `src`.

**m** – 3x3 or 4x4 floating-point transformation matrix.

The function `perspectiveTransform` transforms every element of `src` by treating it as a 2D or 3D vector, in the following way:

$$(x, y, z) \rightarrow (x'/w, y'/w, z'/w)$$

where

$$(x', y', z', w') = \mathtt{mat} \cdot \begin{bmatrix} x & y & z & 1 \end{bmatrix}$$

and

$$w = \begin{cases} w' & \text{if } w' \neq 0 \\ \infty & \text{otherwise} \end{cases}$$

Here a 3D vector transformation is shown. In case of a 2D vector transformation, the `z` component is omitted.

**Note:** The function transforms a sparse set of 2D or 3D vectors. If you want to transform an image using perspective transformation, use `warpPerspective()` . If you have an inverse problem, that is, you want to compute the most probable perspective transformation out of several pairs of corresponding points, you can use `getPerspectiveTransform()` or `findHomography()` .

**See Also:**

`transform()`, `warpPerspective()`, `getPerspectiveTransform()`, `findHomography()`

## phase

Calculates the rotation angle of 2D vectors.

**C++:** void **phase**(InputArray **x**, InputArray **y**, OutputArray **angle**, bool **angleInDegrees**=false)

**Python:** `cv2.`**phase**`(x, y`[, angle[, angleInDegrees ]]`) → angle`

**Parameters**

**x** – input floating-point array of x-coordinates of 2D vectors.

**y** – input array of y-coordinates of 2D vectors; it must have the same size and the same type as `x`.

**angle** – output array of vector angles; it has the same size and same type as `x` .

**angleInDegrees** – when true, the function calculates the angle in degrees, otherwise, they are measured in radians.

The function `phase` calculates the rotation angle of each 2D vector that is formed from the corresponding elements of `x` and `y` :

$$\mathtt{angle}(I) = \mathtt{atan2}(\mathtt{y}(I), \mathtt{x}(I))$$

The angle estimation accuracy is about 0.3 degrees. When `x(I)=y(I)=0` , the corresponding `angle(I)` is set to 0.

## polarToCart

Calculates x and y coordinates of 2D vectors from their magnitude and angle.

**C++:** void **polarToCart**(InputArray **magnitude**, InputArray **angle**, OutputArray **x**, OutputArray **y**, bool **angleInDegrees**=false)

**Python:** cv2.**polarToCart**(magnitude, angle[, x[, y[, angleInDegrees ]]]) → x, y

**C:** void **cvPolarToCart**(const CvArr* **magnitude**, const CvArr* **angle**, CvArr* **x**, CvArr* **y**, int **angle_in_degrees**=0 )

**Python:** cv.**PolarToCart**(magnitude, angle, x, y, angleInDegrees=0) → None

> **Parameters**

>> **magnitude** – input floating-point array of magnitudes of 2D vectors; it can be an empty matrix (=Mat()), in this case, the function assumes that all the magnitudes are =1; if it is not empty, it must have the same size and type as angle.

>> **angle** – input floating-point array of angles of 2D vectors.

>> **x** – output array of x-coordinates of 2D vectors; it has the same size and type as angle.

>> **y** – output array of y-coordinates of 2D vectors; it has the same size and type as angle.

>> **angleInDegrees** – when true, the input angles are measured in degrees, otherwise, they are measured in radians.

The function polarToCart calculates the Cartesian coordinates of each 2D vector represented by the corresponding elements of magnitude and angle :

$$x(I) = \mathtt{magnitude}(I)\cos(\mathtt{angle}(I))$$
$$y(I) = \mathtt{magnitude}(I)\sin(\mathtt{angle}(I))$$

The relative accuracy of the estimated coordinates is about 1e-6.

**See Also:**

cartToPolar(), magnitude(), phase(), exp(), log(), pow(), sqrt()

## pow

Raises every array element to a power.

**C++:** void **pow**(InputArray **src**, double **power**, OutputArray **dst**)

**Python:** cv2.**pow**(src, power[, dst ]) → dst

**C:** void **cvPow**(const CvArr* **src**, CvArr* **dst**, double **power**)

**Python:** cv.**Pow**(src, dst, power) → None

> **Parameters**

>> **src** – input array.

>> **power** – exponent of power.

>> **dst** – output array of the same size and type as src.

The function pow raises every element of the input array to power :

$$\mathtt{dst}(I) = \begin{cases} \mathtt{src}(I)^{\mathtt{power}} & \text{if } \mathtt{power} \text{ is integer} \\ |\mathtt{src}(I)|^{\mathtt{power}} & \text{otherwise} \end{cases}$$

So, for a non-integer power exponent, the absolute values of input array elements are used. However, it is possible to get true values for negative values using some extra operations. In the example below, computing the 5th root of array `src` shows:

```
Mat mask = src < 0;
pow(src, 1./5, dst);
subtract(Scalar::all(0), dst, dst, mask);
```

For some values of `power` , such as integer values, 0.5 and -0.5, specialized faster algorithms are used.

Special values (NaN, Inf) are not handled.

**See Also:**

`sqrt()`, `exp()`, `log()`, `cartToPolar()`, `polarToCart()`

## RNG

**class RNG**

Random number generator. It encapsulates the state (currently, a 64-bit integer) and has methods to return scalar random values and to fill arrays with random values. Currently it supports uniform and Gaussian (normal) distributions. The generator uses Multiply-With-Carry algorithm, introduced by G. Marsaglia ( http://en.wikipedia.org/wiki/Multiply-with-carry ). Gaussian-distribution random numbers are generated using the Ziggurat algorithm ( http://en.wikipedia.org/wiki/Ziggurat_algorithm ), introduced by G. Marsaglia and W. W. Tsang.

## RNG::RNG

The constructors

**C++:** RNG::**RNG**()

**C++:** RNG::**RNG**(uint64 **state**)

> **Parameters**
>
> > **state** – 64-bit value used to initialize the RNG.

These are the RNG constructors. The first form sets the state to some pre-defined value, equal to `2**32-1` in the current implementation. The second form sets the state to the specified value. If you passed `state=0` , the constructor uses the above default value instead to avoid the singular random number sequence, consisting of all zeros.

## RNG::next

Returns the next random number.

**C++:** unsigned int RNG::**next**()

The method updates the state using the MWC algorithm and returns the next 32-bit random number.

## RNG::operator T

Returns the next random number of the specified type.

**C++:** RNG::**operator** uchar()

**C++:** RNG::**operator** schar()

**C++:** RNG::**operator** ushort()

**C++:** RNG::**operator** short int()

**C++:** RNG::**operator** int()

**C++:** RNG::**operator** unsigned int()

**C++:** RNG::**operator** float()

**C++:** RNG::**operator** double()

Each of the methods updates the state using the MWC algorithm and returns the next random number of the specified type. In case of integer types, the returned number is from the available value range for the specified type. In case of floating-point types, the returned value is from [0,1) range.

## RNG::operator ()

Returns the next random number.

**C++:** unsigned int RNG::**operator()**()

**C++:** unsigned int RNG::**operator()**(unsigned int **N**)

> **Parameters**

>> **N** – upper non-inclusive boundary of the returned random number.

The methods transform the state using the MWC algorithm and return the next random number. The first form is equivalent to RNG::next() . The second form returns the random number modulo N , which means that the result is in the range [0, N) .

## RNG::uniform

Returns the next random number sampled from the uniform distribution.

**C++:** int RNG::**uniform**(int **a**, int **b**)

**C++:** float RNG::**uniform**(float **a**, float **b**)

**C++:** double RNG::**uniform**(double **a**, double **b**)

> **Parameters**

>> **a** – lower inclusive boundary of the returned random numbers.

>> **b** – upper non-inclusive boundary of the returned random numbers.

The methods transform the state using the MWC algorithm and return the next uniformly-distributed random number of the specified type, deduced from the input parameter type, from the range [a, b) . There is a nuance illustrated by the following sample:

```
RNG rng;

// always produces 0
double a = rng.uniform(0, 1);

// produces double from [0, 1)
double a1 = rng.uniform((double)0, (double)1);

// produces float from [0, 1)
double b = rng.uniform(0.f, 1.f);
```

```
// produces double from [0, 1)
double c = rng.uniform(0., 1.);

// may cause compiler error because of ambiguity:
//  RNG::uniform(0, (int)0.999999)? or RNG::uniform((double)0, 0.99999)?
double d = rng.uniform(0, 0.999999);
```

The compiler does not take into account the type of the variable to which you assign the result of `RNG::uniform` . The only thing that matters to the compiler is the type of a and b parameters. So, if you want a floating-point random number, but the range boundaries are integer numbers, either put dots in the end, if they are constants, or use explicit type cast operators, as in the a1 initialization above.

## RNG::gaussian

Returns the next random number sampled from the Gaussian distribution.

**C++:** double RNG::**gaussian**(double **sigma**)

> **Parameters**
>
> > **sigma** – standard deviation of the distribution.

The method transforms the state using the MWC algorithm and returns the next random number from the Gaussian distribution `N(0,sigma)` . That is, the mean value of the returned random numbers is zero and the standard deviation is the specified `sigma` .

## RNG::fill

Fills arrays with random numbers.

**C++:** void RNG::**fill**(InputOutputArray **mat**, int **distType**, InputArray **a**, InputArray **b**, bool **saturateRange**=false )

> **Parameters**
>
> > **mat** – 2D or N-dimensional matrix; currently matrices with more than 4 channels are not supported by the methods, use `Mat::reshape()` as a possible workaround.
> >
> > **distType** – distribution type, `RNG::UNIFORM` or `RNG::NORMAL`.
> >
> > **a** – first distribution parameter; in case of the uniform distribution, this is an inclusive lower boundary, in case of the normal distribution, this is a mean value.
> >
> > **b** – second distribution parameter; in case of the uniform distribution, this is a non-inclusive upper boundary, in case of the normal distribution, this is a standard deviation (diagonal of the standard deviation matrix or the full standard deviation matrix).
> >
> > **saturateRange** – pre-saturation flag; for uniform distribution only; if true, the method will first convert a and b to the acceptable value range (according to the mat datatype) and then will generate uniformly distributed random numbers within the range `[saturate(a), saturate(b))`, if `saturateRange=false`, the method will generate uniformly distributed random numbers in the original range `[a, b)` and then will saturate them, it means, for example, that `theRNG().fill(mat_8u, RNG::UNIFORM, -DBL_MAX, DBL_MAX)` will likely produce array mostly filled with 0's and 255's, since the range (0, 255) is significantly smaller than `[-DBL_MAX, DBL_MAX)`.

Each of the methods fills the matrix with the random values from the specified distribution. As the new numbers are generated, the RNG state is updated accordingly. In case of multiple-channel images, every channel is filled

independently, which means that RNG cannot generate samples from the multi-dimensional Gaussian distribution with non-diagonal covariance matrix directly. To do that, the method generates samples from multi-dimensional standard Gaussian distribution with zero mean and identity covariation matrix, and then transforms them using `transform()` to get samples from the specified Gaussian distribution.

## randu

Generates a single uniformly-distributed random number or an array of random numbers.

**C++:** template<typename _Tp> _Tp **randu**()

**C++:** void **randu**(InputOutputArray **dst**, InputArray **low**, InputArray **high**)

**Python:** cv2.**randu**(dst, low, high) → None

> **Parameters**
>
> > **dst** – output array of random numbers; the array must be pre-allocated.
> >
> > **low** – inclusive lower boundary of the generated random numbers.
> >
> > **high** – exclusive upper boundary of the generated random numbers.

The template functions `randu` generate and return the next uniformly-distributed random value of the specified type. `randu<int>()` is an equivalent to `(int)theRNG();` , and so on. See RNG description.

The second non-template variant of the function fills the matrix `dst` with uniformly-distributed random numbers from the specified range:

$$\texttt{low}_c \leq \texttt{dst}(I)_c < \texttt{high}_c$$

**See Also:**

RNG, randn(), theRNG()

## randn

Fills the array with normally distributed random numbers.

**C++:** void **randn**(InputOutputArray **dst**, InputArray **mean**, InputArray **stddev**)

**Python:** cv2.**randn**(dst, mean, stddev) → None

> **Parameters**
>
> > **dst** – output array of random numbers; the array must be pre-allocated and have 1 to 4 channels.
> >
> > **mean** – mean value (expectation) of the generated random numbers.
> >
> > **stddev** – standard deviation of the generated random numbers; it can be either a vector (in which case a diagonal standard deviation matrix is assumed) or a square matrix.

The function `randn` fills the matrix `dst` with normally distributed random numbers with the specified mean vector and the standard deviation matrix. The generated random numbers are clipped to fit the value range of the output array data type.

**See Also:**

RNG, randu()

## randShuffle

Shuffles the array elements randomly.

**C++:** void **randShuffle**(InputOutputArray **dst**, double **iterFactor**=1., RNG* **rng**=0 )

**Python:** cv2.**randShuffle**(dst[, iterFactor ]) → None

> **Parameters**
>
>> **dst** – input/output numerical 1D array.
>>
>> **iterFactor** – scale factor that determines the number of random swap operations (see the details below).
>>
>> **rng** – optional random number generator used for shuffling; if it is zero, theRNG() () is used instead.

The function randShuffle shuffles the specified 1D array by randomly choosing pairs of elements and swapping them. The number of such swap operations will be dst.rows*dst.cols*iterFactor .

**See Also:**

RNG, sort()

## reduce

Reduces a matrix to a vector.

**C++:** void **reduce**(InputArray **src**, OutputArray **dst**, int **dim**, int **rtype**, int **dtype**=-1 )

**Python:** cv2.**reduce**(src, dim, rtype[, dst[, dtype ] ]) → dst

**C:** void **cvReduce**(const CvArr* **src**, CvArr* **dst**, int **dim**=-1, int **op**=CV_REDUCE_SUM)

**Python:** cv.**Reduce**(src, dst, dim=-1, op=CV_REDUCE_SUM) → None

> **Parameters**
>
>> **src** – input 2D matrix.
>>
>> **dst** – output vector. Its size and type is defined by dim and dtype parameters.
>>
>> **dim** – dimension index along which the matrix is reduced. 0 means that the matrix is reduced to a single row. 1 means that the matrix is reduced to a single column.
>>
>> **rtype** – reduction operation that could be one of the following:
>>
>> – **CV_REDUCE_SUM**: the output is the sum of all rows/columns of the matrix.
>>
>> – **CV_REDUCE_AVG**: the output is the mean vector of all rows/columns of the matrix.
>>
>> – **CV_REDUCE_MAX**: the output is the maximum (column/row-wise) of all rows/columns of the matrix.
>>
>> – **CV_REDUCE_MIN**: the output is the minimum (column/row-wise) of all rows/columns of the matrix.
>>
>> **dtype** – when negative, the output vector will have the same type as the input matrix, otherwise, its type will be CV_MAKE_TYPE(CV_MAT_DEPTH(dtype), src.channels()).

The function reduce reduces the matrix to a vector by treating the matrix rows/columns as a set of 1D vectors and performing the specified operation on the vectors until a single row/column is obtained. For example, the function can be used to compute horizontal and vertical projections of a raster image. In case of CV_REDUCE_SUM and CV_REDUCE_AVG

, the output may have a larger element bit-depth to preserve accuracy. And multi-channel arrays are also supported in these two reduction modes.

**See Also:**

repeat()

## repeat

Fills the output array with repeated copies of the input array.

**C++:** void **repeat**(InputArray **src**, int **ny**, int **nx**, OutputArray **dst**)

**C++:** Mat **repeat**(const Mat& **src**, int **ny**, int **nx**)

**Python:** cv2.**repeat**(src, ny, nx[, dst]) → dst

**C:** void **cvRepeat**(const CvArr* **src**, CvArr* **dst**)

**Python:** cv.**Repeat**(src, dst) → None

> **Parameters**

>> **src** – input array to replicate.

>> **dst** – output array of the same type as src.

>> **ny** – Flag to specify how many times the src is repeated along the vertical axis.

>> **nx** – Flag to specify how many times the src is repeated along the horizontal axis.

The functions repeat() duplicate the input array one or more times along each of the two axes:

$$\texttt{dst}_{ij} = \texttt{src}_{i \mod \texttt{src.rows}, \ j \mod \texttt{src.cols}}$$

The second variant of the function is more convenient to use with *Matrix Expressions* .

**See Also:**

reduce(), *Matrix Expressions*

## scaleAdd

Calculates the sum of a scaled array and another array.

**C++:** void **scaleAdd**(InputArray **src1**, double **alpha**, InputArray **src2**, OutputArray **dst**)

**Python:** cv2.**scaleAdd**(src1, alpha, src2[, dst]) → dst

**C:** void **cvScaleAdd**(const CvArr* **src1**, CvScalar **scale**, const CvArr* **src2**, CvArr* **dst**)

**Python:** cv.**ScaleAdd**(src1, scale, src2, dst) → None

> **Parameters**

>> **src1** – first input array.

>> **scale** – scale factor for the first array.

>> **src2** – second input array of the same size and type as src1.

>> **dst** – output array of the same size and type as src1.

The function `scaleAdd` is one of the classical primitive linear algebra operations, known as DAXPY or SAXPY in BLAS. It calculates the sum of a scaled array and another array:

$$dst(I) = scale \cdot src1(I) + src2(I)$$

The function can also be emulated with a matrix expression, for example:

```
Mat A(3, 3, CV_64F);
...
A.row(0) = A.row(1)*2 + A.row(2);
```

**See Also:**

add(), addWeighted(), subtract(), Mat::dot(), Mat::convertTo(), *Matrix Expressions*

## setIdentity

Initializes a scaled identity matrix.

**C++:** void **setIdentity**(InputOutputArray **mtx**, const Scalar& **s**=Scalar(1) )

**Python:** cv2.**setIdentity**(mtx$[, s]$) → None

**C:** void **cvSetIdentity**(CvArr* **mat**, CvScalar **value**=cvRealScalar(1))

**Python:** cv.**SetIdentity**(mat, value=1) → None

> **Parameters**
>
>> **mtx** – matrix to initialize (not necessarily square).
>>
>> **value** – value to assign to diagonal elements.

The function setIdentity() initializes a scaled identity matrix:

$$mtx(i,j) = \begin{cases} value & \text{if } i = j \\ 0 & \text{otherwise} \end{cases}$$

The function can also be emulated using the matrix initializers and the matrix expressions:

```
Mat A = Mat::eye(4, 3, CV_32F)*5;
// A will be set to [[5, 0, 0], [0, 5, 0], [0, 0, 5], [0, 0, 0]]
```

**See Also:**

Mat::zeros(), Mat::ones(), *Matrix Expressions*, Mat::setTo(), Mat::operator=()

## solve

Solves one or more linear systems or least-squares problems.

**C++:** bool **solve**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, int **flags**=DECOMP_LU)

**Python:** cv2.**solve**(src1, src2$[, dst[, flags]]$) → retval, dst

**C:** int **cvSolve**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, int **method**=CV_LU)

**Python:** cv.**Solve**(A, B, X, method=CV_LU) → None

> **Parameters**
>
>> **src1** – input matrix on the left-hand side of the system.

>    **src2** – input matrix on the right-hand side of the system.
>
>    **dst** – output solution.
>
>    **flags** – solution (matrix inversion) method.
>
>    - **DECOMP_LU** Gaussian elimination with optimal pivot element chosen.
>    - **DECOMP_CHOLESKY** Cholesky $LL^T$ factorization; the matrix `src1` must be symmetrical and positively defined.
>    - **DECOMP_EIG** eigenvalue decomposition; the matrix `src1` must be symmetrical.
>    - **DECOMP_SVD** singular value decomposition (SVD) method; the system can be over-defined and/or the matrix `src1` can be singular.
>    - **DECOMP_QR** QR factorization; the system can be over-defined and/or the matrix `src1` can be singular.
>    - **DECOMP_NORMAL** while all the previous flags are mutually exclusive, this flag can be used together with any of the previous; it means that the normal equations $src1^T \cdot src1 \cdot dst = src1^T src2$ are solved instead of the original system $src1 \cdot dst = src2$ .

The function `solve` solves a linear system or least-squares problem (the latter is possible with SVD or QR methods, or by specifying the flag `DECOMP_NORMAL` ):

$$dst = \arg\min_X \| src1 \cdot X - src2 \|$$

If `DECOMP_LU` or `DECOMP_CHOLESKY` method is used, the function returns 1 if `src1` (or $src1^T src1$ ) is non-singular. Otherwise, it returns 0. In the latter case, `dst` is not valid. Other methods find a pseudo-solution in case of a singular left-hand side part.

---

**Note:** If you want to find a unity-norm solution of an under-defined singular system $src1 \cdot dst = 0$ , the function `solve` will not do the work. Use `SVD::solveZ()` instead.

---

**See Also:**

`invert()`, `SVD`, `eigen()`

## solveCubic

Finds the real roots of a cubic equation.

**C++:** int **solveCubic**(InputArray **coeffs**, OutputArray **roots**)

**Python:** cv2.**solveCubic**(coeffs[, roots]) → retval, roots

**C:** int **cvSolveCubic**(const CvMat* **coeffs**, CvMat* **roots**)

**Python:** cv.**SolveCubic**(coeffs, roots) → None

>    **Parameters**
>
>    >    **coeffs** – equation coefficients, an array of 3 or 4 elements.
>    >
>    >    **roots** – output array of real roots that has 1 or 3 elements.

The function `solveCubic` finds the real roots of a cubic equation:

- if `coeffs` is a 4-element vector:

$$coeffs[0]x^3 + coeffs[1]x^2 + coeffs[2]x + coeffs[3] = 0$$

- if `coeffs` is a 3-element vector:

$$x^3 + \text{coeffs}[0]x^2 + \text{coeffs}[1]x + \text{coeffs}[2] = 0$$

The roots are stored in the `roots` array.

## solvePoly

Finds the real or complex roots of a polynomial equation.

**C++:** double **solvePoly**(InputArray **coeffs**, OutputArray **roots**, int **maxIters**=300 )

**Python:** cv2.**solvePoly**(coeffs$\big[$, roots$\big[$, maxIters $\big]\big]$) $\rightarrow$ retval, roots

> **Parameters**
>
> > **coeffs** – array of polynomial coefficients.
> >
> > **roots** – output (complex) array of roots.
> >
> > **maxIters** – maximum number of iterations the algorithm does.

The function `solvePoly` finds real and complex roots of a polynomial equation:

$$\text{coeffs}[n]x^n + \text{coeffs}[n-1]x^{n-1} + ... + \text{coeffs}[1]x + \text{coeffs}[0] = 0$$

## sort

Sorts each row or each column of a matrix.

**C++:** void **sort**(InputArray **src**, OutputArray **dst**, int **flags**)

**Python:** cv2.**sort**(src, flags$\big[$, dst $\big]$) $\rightarrow$ dst

> **Parameters**
>
> > **src** – input single-channel array.
> >
> > **dst** – output array of the same size and type as `src`.
> >
> > **flags** – operation flags, a combination of the following values:
> >
> > > – **CV_SORT_EVERY_ROW** each matrix row is sorted independently.
> > >
> > > – **CV_SORT_EVERY_COLUMN** each matrix column is sorted independently; this flag and the previous one are mutually exclusive.
> > >
> > > – **CV_SORT_ASCENDING** each matrix row is sorted in the ascending order.
> > >
> > > – **CV_SORT_DESCENDING** each matrix row is sorted in the descending order; this flag and the previous one are also mutually exclusive.

The function `sort` sorts each matrix row or each matrix column in ascending or descending order. So you should pass two operation flags to get desired behaviour. If you want to sort matrix rows or columns lexicographically, you can use STL `std::sort` generic function with the proper comparison predicate.

**See Also:**

`sortIdx()`, `randShuffle()`

## sortIdx

Sorts each row or each column of a matrix.

**C++:** void **sortIdx**(InputArray **src**, OutputArray **dst**, int **flags**)

**Python:** cv2.**sortIdx**(src, flags[, dst]) → dst

> **Parameters**
>
> > **src** – input single-channel array.
> >
> > **dst** – output integer array of the same size as `src`.
> >
> > **flags** – operation flags that could be a combination of the following values:
> >
> > – **CV_SORT_EVERY_ROW** each matrix row is sorted independently.
> >
> > – **CV_SORT_EVERY_COLUMN** each matrix column is sorted independently; this flag and the previous one are mutually exclusive.
> >
> > – **CV_SORT_ASCENDING** each matrix row is sorted in the ascending order.
> >
> > – **CV_SORT_DESCENDING** each matrix row is sorted in the descending order; his flag and the previous one are also mutually exclusive.

The function `sortIdx` sorts each matrix row or each matrix column in the ascending or descending order. So you should pass two operation flags to get desired behaviour. Instead of reordering the elements themselves, it stores the indices of sorted elements in the output array. For example:

```
Mat A = Mat::eye(3,3,CV_32F), B;
sortIdx(A, B, CV_SORT_EVERY_ROW + CV_SORT_ASCENDING);
// B will probably contain
// (because of equal elements in A some permutations are possible):
// [[1, 2, 0], [0, 2, 1], [0, 1, 2]]
```

**See Also:**

sort(), randShuffle()

## split

Divides a multi-channel array into several single-channel arrays.

**C++:** void **split**(const Mat& **src**, Mat* **mvbegin**)

**C++:** void **split**(InputArray **m**, OutputArrayOfArrays **mv**)

**Python:** cv2.**split**(m[, mv]) → mv

**C:** void **cvSplit**(const CvArr* **src**, CvArr* **dst0**, CvArr* **dst1**, CvArr* **dst2**, CvArr* **dst3**)

**Python:** cv.**Split**(src, dst0, dst1, dst2, dst3) → None

> **Parameters**
>
> > **src** – input multi-channel array.
> >
> > **mv** – output array or vector of arrays; in the first variant of the function the number of arrays must match `src.channels()`; the arrays themselves are reallocated, if needed.

The functions `split` split a multi-channel array into separate single-channel arrays:

$$\texttt{mv}[c](I) = \texttt{src}(I)_c$$

If you need to extract a single channel or do some other sophisticated channel permutation, use `mixChannels()` .

**See Also:**

`merge()`, `mixChannels()`, `cvtColor()`

## sqrt

Calculates a square root of array elements.

**C++:** void **sqrt**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**sqrt**(src[, dst]) → dst

**C:** float **cvSqrt**(float **value**)

**Python:** cv.**Sqrt**(value) → float

> **Parameters**

>> **src** – input floating-point array.

>> **dst** – output array of the same size and type as `src`.

The functions `sqrt` calculate a square root of each input array element. In case of multi-channel arrays, each channel is processed independently. The accuracy is approximately the same as of the built-in `std::sqrt` .

**See Also:**

`pow()`, `magnitude()`

## subtract

Calculates the per-element difference between two arrays or array and a scalar.

**C++:** void **subtract**(InputArray **src1**, InputArray **src2**, OutputArray **dst**, InputArray **mask**=noArray(), int **dtype**=-1)

**Python:** cv2.**subtract**(src1, src2[, dst[, mask[, dtype]]]) → dst

**C:** void **cvSub**(const CvArr* **src1**, const CvArr* **src2**, CvArr* **dst**, const CvArr* **mask**=NULL)

**C:** void **cvSubRS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL )

**C:** void **cvSubS**(const CvArr* **src**, CvScalar **value**, CvArr* **dst**, const CvArr* **mask**=NULL )

**Python:** cv.**Sub**(src1, src2, dst, mask=None) → None

**Python:** cv.**SubRS**(src, value, dst, mask=None) → None

**Python:** cv.**SubS**(src, value, dst, mask=None) → None

> **Parameters**

>> **src1** – first input array or a scalar.

>> **src2** – second input array or a scalar.

>> **dst** – output array of the same size and the same number of channels as the input array.

>> **mask** – optional operation mask; this is an 8-bit single channel array that specifies elements of the output array to be changed.

>> **dtype** – optional depth of the output array (see the details below).

The function `subtract` calculates:

- Difference between two arrays, when both input arrays have the same size and the same number of channels:

$$\mathtt{dst}(I) = \mathtt{saturate}(\mathtt{src1}(I) - \mathtt{src2}(I)) \quad \text{if } \mathtt{mask}(I) \neq 0$$

- Difference between an array and a scalar, when `src2` is constructed from `Scalar` or has the same number of elements as `src1.channels()`:

$$\mathtt{dst}(I) = \mathtt{saturate}(\mathtt{src1}(I) - \mathtt{src2}) \quad \text{if } \mathtt{mask}(I) \neq 0$$

- Difference between a scalar and an array, when `src1` is constructed from `Scalar` or has the same number of elements as `src2.channels()`:

$$\mathtt{dst}(I) = \mathtt{saturate}(\mathtt{src1} - \mathtt{src2}(I)) \quad \text{if } \mathtt{mask}(I) \neq 0$$

- The reverse difference between a scalar and an array in the case of `SubRS`:

$$\mathtt{dst}(I) = \mathtt{saturate}(\mathtt{src2} - \mathtt{src1}(I)) \quad \text{if } \mathtt{mask}(I) \neq 0$$

where `I` is a multi-dimensional index of array elements. In case of multi-channel arrays, each channel is processed independently.

The first function in the list above can be replaced with matrix expressions:

```
dst = src1 - src2;
dst -= src1; // equivalent to subtract(dst, src1, dst);
```

The input arrays and the output array can all have the same or different depths. For example, you can subtract to 8-bit unsigned arrays and store the difference in a 16-bit signed array. Depth of the output array is determined by `dtype` parameter. In the second and third cases above, as well as in the first case, when `src1.depth() == src2.depth()`, `dtype` can be set to the default `-1`. In this case the output array will have the same depth as the input array, be it `src1`, `src2` or both.

---

**Note:** Saturation is not applied when the output array has the depth `CV_32S`. You may even get result of an incorrect sign in the case of overflow.

---

**See Also:**

`add()`, `addWeighted()`, `scaleAdd()`, `Mat::convertTo()`, *Matrix Expressions*

## SVD

**class SVD**

Class for computing Singular Value Decomposition of a floating-point matrix. The Singular Value Decomposition is used to solve least-square problems, under-determined linear systems, invert matrices, compute condition numbers, and so on.

For a faster operation, you can pass `flags=SVD::MODIFY_A|...` to modify the decomposed matrix when it is not necessary to preserve it. If you want to compute a condition number of a matrix or an absolute value of its determinant, you do not need u and vt . You can pass `flags=SVD::NO_UV|...` . Another flag `FULL_UV` indicates that full-size u and vt must be computed, which is not necessary most of the time.

**See Also:**

`invert()`, `solve()`, `eigen()`, `determinant()`

---

## SVD::SVD

The constructors.

**C++:** `SVD::`**`SVD`**`()`

**C++:** `SVD::`**`SVD`**`(`InputArray **src**, int **flags**=0 `)`

> **Parameters**
>
> > **src** – decomposed matrix.
> >
> > **flags** – operation flags.
> >
> > – **SVD::MODIFY_A** use the algorithm to modify the decomposed matrix; it can save space and speed up processing.
> >
> > – **SVD::NO_UV** indicates that only a vector of singular values w is to be processed, while u and vt will be set to empty matrices.
> >
> > – **SVD::FULL_UV** when the matrix is not square, by default the algorithm produces u and vt matrices of sufficiently large size for the further A reconstruction; if, however, FULL_UV flag is specified, u and ``vt``will be full-size square orthogonal matrices.

The first constructor initializes an empty SVD structure. The second constructor initializes an empty SVD structure and then calls `SVD::operator()` .

## SVD::operator ()

Performs SVD of a matrix.

**C++:** SVD& `SVD::`**`operator()`**`(`InputArray **src**, int **flags**=0 `)`

> **Parameters**
>
> > **src** – decomposed matrix.
> >
> > **flags** – operation flags.
> >
> > – **SVD::MODIFY_A** use the algorithm to modify the decomposed matrix; it can save space and speed up processing.
> >
> > – **SVD::NO_UV** use only singular values; the algorithm does not compute u and vt matrices.
> >
> > – **SVD::FULL_UV** when the matrix is not square, by default the algorithm produces u and vt matrices of sufficiently large size for the further A reconstruction; if, however, the FULL_UV flag is specified, u and vt are full-size square orthogonal matrices.

The operator performs the singular value decomposition of the supplied matrix. The u,``vt`` , and the vector of singular values w are stored in the structure. The same SVD structure can be reused many times with different matrices. Each time, if needed, the previous u,``vt`` , and w are reclaimed and the new matrices are created, which is all handled by `Mat::create()` .

## SVD::compute

Performs SVD of a matrix

**C++:** **static** void `SVD::`**`compute`**`(`InputArray **src**, OutputArray **w**, OutputArray **u**, OutputArray **vt**, int **flags**=0 `)`

**C++:** **static** void `SVD::`**`compute`**`(`InputArray **src**, OutputArray **w**, int **flags**=0 `)`

**Python:** cv2.**SVDecomp**(src[, w[, u[, vt[, flags ] ] ] ]) → w, u, vt

**C:** void **cvSVD**(CvArr* **A**, CvArr* **W**, CvArr* **U**=NULL, CvArr* **V**=NULL, int **flags**=0 )

**Python:** cv.**SVD**(A, W, U=None, V=None, flags=0) → None

> **Parameters**

>> **src** – decomposed matrix

>> **w** – calculated singular values

>> **u** – calculated left singular vectors

>> **V** – calculated right singular vectors

>> **vt** – transposed matrix of right singular values

>> **flags** – operation flags - see SVD::SVD().

The methods/functions perform SVD of matrix. Unlike SVD::SVD constructor and SVD::operator(), they store the results to the user-provided matrices.

```
Mat A, w, u, vt;
SVD::compute(A, w, u, vt);
```

## SVD::solveZ

Solves an under-determined singular linear system.

**C++: static** void SVD::**solveZ**(InputArray **src**, OutputArray **dst**)

> **Parameters**

>> **src** – left-hand-side matrix.

>> **dst** – found solution.

The method finds a unit-length solution x of a singular linear system A*x = 0. Depending on the rank of A, there can be no solutions, a single solution or an infinite number of solutions. In general, the algorithm solves the following problem:

$$\text{dst} = \arg \min_{\text{x}:\|\text{x}\|=1} \|\text{src} \cdot \text{x}\|$$

## SVD::backSubst

Performs a singular value back substitution.

**C++:** void SVD::**backSubst**(InputArray **rhs**, OutputArray **dst**) const

**C++: static** void SVD::**backSubst**(InputArray **w**, InputArray **u**, InputArray **vt**, InputArray **rhs**, OutputArray **dst**)

**Python:** cv2.**SVBackSubst**(w, u, vt, rhs[, dst ]) → dst

**C:** void **cvSVBkSb**(const CvArr* **W**, const CvArr* **U**, const CvArr* **V**, const CvArr* **B**, CvArr* **X**, int **flags**)

**Python:** cv.**SVBkSb**(W, U, V, B, X, flags) → None

> **Parameters**

>> **w** – singular values

>> **u** – left singular vectors

**V** – right singular vectors

**vt** – transposed matrix of right singular vectors.

**rhs** – right-hand side of a linear system (u*w*v')*dst = rhs to be solved, where A has been previously decomposed.

**dst** – found solution of the system.

The method calculates a back substitution for the specified right-hand side:

$$x = vt^T \cdot diag(w)^{-1} \cdot u^T \cdot rhs \sim A^{-1} \cdot rhs$$

Using this technique you can either get a very accurate solution of the convenient linear system, or the best (in the least-squares terms) pseudo-solution of an overdetermined linear system.

---

**Note:** Explicit SVD with the further back substitution only makes sense if you need to solve many linear systems with the same left-hand side (for example, src ). If all you need is to solve a single system (possibly with multiple rhs immediately available), simply call solve() add pass DECOMP_SVD there. It does absolutely the same thing.

---

## sum

Calculates the sum of array elements.

**C++:** Scalar **sum**(InputArray **src**)

**Python:** cv2.**sumElems**(src) → retval

**C:** CvScalar **cvSum**(const CvArr* **arr**)

**Python:** cv.**Sum**(arr) → scalar

>        **Parameters**

>                **arr** – input array that must have from 1 to 4 channels.

The functions sum calculate and return the sum of array elements, independently for each channel.

**See Also:**

countNonZero(), mean(), meanStdDev(), norm(), minMaxLoc(), reduce()

## theRNG

Returns the default random number generator.

**C++:** RNG& **theRNG**()

The function theRNG returns the default random number generator. For each thread, there is a separate random number generator, so you can use the function safely in multi-thread environments. If you just need to get a single random number using this generator or initialize an array, you can use randu() or randn() instead. But if you are going to generate many random numbers inside a loop, it is much faster to use this function to retrieve the generator and then use RNG::operator _Tp() .

**See Also:**

RNG, randu(), randn()

## trace

Returns the trace of a matrix.

**C++:** Scalar **trace**(InputArray **mtx**)

**Python:** cv2.**trace**(mtx) → retval

**C:** CvScalar **cvTrace**(const CvArr* **mat**)

**Python:** cv.**Trace**(mat) → scalar

> **Parameters**
>
> > **mat** – input matrix.

The function `trace` returns the sum of the diagonal elements of the matrix `mtx` .

$$\text{tr}(\text{mtx}) = \sum_i \text{mtx}(i, i)$$

## transform

Performs the matrix transformation of every array element.

**C++:** void **transform**(InputArray **src**, OutputArray **dst**, InputArray **m**)

**Python:** cv2.**transform**(src, m[, dst]) → dst

**C:** void **cvTransform**(const CvArr* **src**, CvArr* **dst**, const CvMat* **transmat**, const CvMat* **shiftvec**=NULL )

**Python:** cv.**Transform**(src, dst, transmat, shiftvec=None) → None

> **Parameters**
>
> > **src** – input array that must have as many channels (1 to 4) as `m.cols` or `m.cols-1`.
> >
> > **dst** – output array of the same size and depth as `src`; it has as many channels as `m.rows`.
> >
> > **m** – transformation 2x2 or 2x3 floating-point matrix.
> >
> > **shiftvec** – optional translation vector (when `m` is 2x2)

The function `transform` performs the matrix transformation of every element of the array `src` and stores the results in `dst` :

$$\text{dst}(I) = \text{m} \cdot \text{src}(I)$$

(when `m.cols=src.channels()` ), or

$$\text{dst}(I) = \text{m} \cdot [\text{src}(I); 1]$$

(when `m.cols=src.channels()+1` )

Every element of the `N` -channel array `src` is interpreted as `N` -element vector that is transformed using the `M x N` or `M x (N+1)` matrix `m` to `M`-element vector - the corresponding element of the output array `dst` .

The function may be used for geometrical transformation of `N` -dimensional points, arbitrary linear color space transformation (such as various kinds of RGB to YUV transforms), shuffling the image channels, and so forth.

**See Also:**

`perspectiveTransform()`, `getAffineTransform()`, `estimateRigidTransform()`, `warpAffine()`, `warpPerspective()`

## transpose

Transposes a matrix.

**C++:** void **transpose**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**transpose**(src[, dst]) → dst

**C:** void **cvTranspose**(const CvArr* **src**, CvArr* **dst**)

**Python:** cv.**Transpose**(src, dst) → None

> **Parameters**
>> **src** – input array.
>>
>> **dst** – output array of the same type as src.

The function transpose() transposes the matrix src :

$$dst(i, j) = src(j, i)$$

---

**Note:** No complex conjugation is done in case of a complex matrix. It it should be done separately if needed.

---

## 2.5 Drawing Functions

Drawing functions work with matrices/images of arbitrary depth. The boundaries of the shapes can be rendered with antialiasing (implemented only for 8-bit images for now). All the functions include the parameter color that uses an RGB value (that may be constructed with CV_RGB or the Scalar_ constructor ) for color images and brightness for grayscale images. For color images, the channel ordering is normally *Blue, Green, Red*. This is what imshow(), imread(), and imwrite() expect. So, if you form a color using the Scalar constructor, it should look like:

$$Scalar(blue\_component, green\_component, red\_component[, alpha\_component])$$

If you are using your own image rendering and I/O functions, you can use any channel ordering. The drawing functions process each channel independently and do not depend on the channel order or even on the used color space. The whole image can be converted from BGR to RGB or to a different color space using cvtColor() .

If a drawn figure is partially or completely outside the image, the drawing functions clip it. Also, many drawing functions can handle pixel coordinates specified with sub-pixel accuracy. This means that the coordinates can be passed as fixed-point numbers encoded as integers. The number of fractional bits is specified by the shift parameter and the real point coordinates are calculated as $Point(x, y) \rightarrow Point2f(x * 2^{-shift}, y * 2^{-shift})$ . This feature is especially effective when rendering antialiased shapes.

---

**Note:** The functions do not support alpha-transparency when the target image is 4-channel. In this case, the color[3] is simply copied to the repainted pixels. Thus, if you want to paint semi-transparent shapes, you can paint them in a separate buffer and then blend it with the main image.

---

## circle

Draws a circle.

**C++:** void **circle**(Mat& **img**, Point **center**, int **radius**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0)

**Python:** cv2.**circle**(img, center, radius, color[, thickness[, lineType[, shift ] ] ]) → None

**C:** void **cvCircle**(CvArr* **img**, CvPoint **center**, int **radius**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int **shift**=0 )

**Python:** cv.**Circle**(img, center, radius, color, thickness=1, lineType=8, shift=0) → None

> **Parameters**
>> **img** – Image where the circle is drawn.
>>
>> **center** – Center of the circle.
>>
>> **radius** – Radius of the circle.
>>
>> **color** – Circle color.
>>
>> **thickness** – Thickness of the circle outline, if positive. Negative thickness means that a filled circle is to be drawn.
>>
>> **lineType** – Type of the circle boundary. See the line() description.
>>
>> **shift** – Number of fractional bits in the coordinates of the center and in the radius value.

The function circle draws a simple or filled circle with a given center and radius.

## clipLine

Clips the line against the image rectangle.

**C++:** bool **clipLine**(Size **imgSize**, Point& **pt1**, Point& **pt2**)

**C++:** bool **clipLine**(Rect **imgRect**, Point& **pt1**, Point& **pt2**)

**Python:** cv2.**clipLine**(imgRect, pt1, pt2) → retval, pt1, pt2

**C:** int **cvClipLine**(CvSize **img_size**, CvPoint* **pt1**, CvPoint* **pt2**)

**Python:** cv.**ClipLine**(imgSize, pt1, pt2) -> (point1, point2)

> **Parameters**
>> **imgSize** – Image size. The image rectangle is Rect(0, 0, imgSize.width, imgSize.height) .
>>
>> **imgRect** – Image rectangle.
>>
>> **pt1** – First line point.
>>
>> **pt2** – Second line point.

The functions clipLine calculate a part of the line segment that is entirely within the specified rectangle. They return false if the line segment is completely outside the rectangle. Otherwise, they return true .

## ellipse

Draws a simple or thick elliptic arc or fills an ellipse sector.

**C++:** void **ellipse**(Mat& **img**, Point **center**, Size **axes**, double **angle**, double **startAngle**, double **endAngle**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0)

**C++:** void **ellipse**(Mat& **img**, const RotatedRect& **box**, const Scalar& **color**, int **thickness**=1, int **line-Type**=8)

**Python:** cv2.**ellipse**(img, center, axes, angle, startAngle, endAngle, color$\left[\right.$, thickness$\left[\right.$, lineType$\left[\right.$, shift$\left.\right]\left.\right]$
$\left.\right]$) → None

**Python:** cv2.**ellipse**(img, box, color$\left[\right.$, thickness$\left[\right.$, lineType$\left.\right]\left.\right]$) → None

**C:** void **cvEllipse**(CvArr* **img**, CvPoint **center**, CvSize **axes**, double **angle**, double **start_angle**, double
**end_angle**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int **shift**=0 )

**Python:** cv.**Ellipse**(img, center, axes, angle, start_angle, end_angle, color, thickness=1, lineType=8,
shift=0) → None

**C:** void **cvEllipseBox**(CvArr* **img**, CvBox2D **box**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int
**shift**=0 )

**Python:** cv.**EllipseBox**(img, box, color, thickness=1, lineType=8, shift=0) → None

> **Parameters**
>
> > **img** – Image.
> >
> > **center** – Center of the ellipse.
> >
> > **axes** – Length of the ellipse axes.
> >
> > **angle** – Ellipse rotation angle in degrees.
> >
> > **startAngle** – Starting angle of the elliptic arc in degrees.
> >
> > **endAngle** – Ending angle of the elliptic arc in degrees.
> >
> > **box** – Alternative ellipse representation via `RotatedRect` or `CvBox2D`. This means that the
> > function draws an ellipse inscribed in the rotated rectangle.
> >
> > **color** – Ellipse color.
> >
> > **thickness** – Thickness of the ellipse arc outline, if positive. Otherwise, this indicates that a
> > filled ellipse sector is to be drawn.
> >
> > **lineType** – Type of the ellipse boundary. See the `line()` description.
> >
> > **shift** – Number of fractional bits in the coordinates of the center and values of axes.
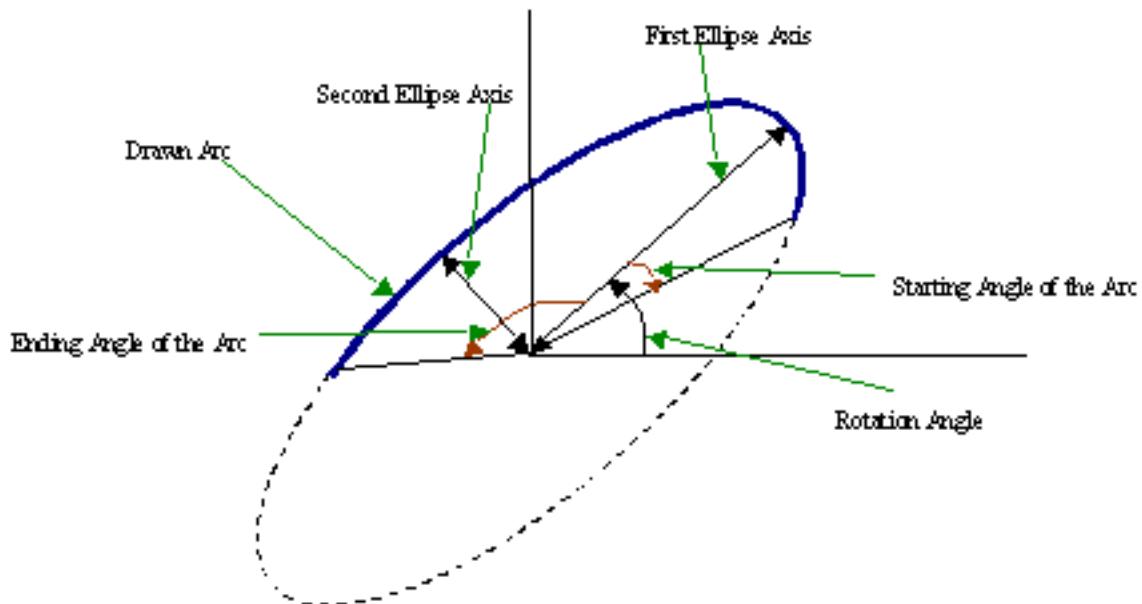
The functions `ellipse` with less parameters draw an ellipse outline, a filled ellipse, an elliptic arc, or a filled ellipse
sector. A piecewise-linear curve is used to approximate the elliptic arc boundary. If you need more control of the
ellipse rendering, you can retrieve the curve using `ellipse2Poly()` and then render it with `polylines()` or fill it
with `fillPoly()` . If you use the first variant of the function and want to draw the whole ellipse, not an arc, pass
`startAngle=0` and `endAngle=360` . The figure below explains the meaning of the parameters.

**Figure 1. Parameters of Elliptic Arc**

## ellipse2Poly

Approximates an elliptic arc with a polyline.

**C++:** void **ellipse2Poly**(Point **center**, Size **axes**, int **angle**, int **arcStart**, int **arcEnd**, int **delta**, vector<Point>& **pts**)

**Python:** cv2.**ellipse2Poly**(center, axes, angle, arcStart, arcEnd, delta) → pts

> **Parameters**

>> **center** – Center of the arc.

>> **axes** – Half-sizes of the arc. See the `ellipse()` for details.

>> **angle** – Rotation angle of the ellipse in degrees. See the `ellipse()` for details.

>> **arcStart** – Starting angle of the elliptic arc in degrees.

>> **arcEnd** – Ending angle of the elliptic arc in degrees.

>> **delta** – Angle between the subsequent polyline vertices. It defines the approximation accuracy.

>> **pts** – Output vector of polyline vertices.

The function `ellipse2Poly` computes the vertices of a polyline that approximates the specified elliptic arc. It is used by `ellipse()`.

## fillConvexPoly

Fills a convex polygon.

**C++:** void **fillConvexPoly**(Mat& **img**, const Point* **pts**, int **npts**, const Scalar& **color**, int **lineType**=8, int **shift**=0)

---

**Python:** cv2.**fillConvexPoly**(img, points, color[, lineType[, shift ]]) → None

**C:** void **cvFillConvexPoly**(CvArr* **img**, const CvPoint* **pts**, int **npts**, CvScalar **color**, int **line_type**=8, int **shift**=0 )

**Python:** cv.**FillConvexPoly**(img, pn, color, lineType=8, shift=0) → None

>       **Parameters**

>> **img** – Image.

>> **pts** – Polygon vertices.

>> **npts** – Number of polygon vertices.

>> **color** – Polygon color.

>> **lineType** – Type of the polygon boundaries. See the `line()` description.

>> **shift** – Number of fractional bits in the vertex coordinates.

The function `fillConvexPoly` draws a filled convex polygon. This function is much faster than the function `fillPoly` . It can fill not only convex polygons but any monotonic polygon without self-intersections, that is, a polygon whose contour intersects every horizontal line (scan line) twice at the most (though, its top-most and/or the bottom edge could be horizontal).

## fillPoly

Fills the area bounded by one or more polygons.

**C++:** void **fillPoly**(Mat& **img**, const Point** **pts**, const int* **npts**, int **ncontours**, const Scalar& **color**, int **lineType**=8, int **shift**=0, Point **offset**=Point() )

**Python:** cv2.**fillPoly**(img, pts, color[, lineType[, shift[, offset ]]]) → None

**C:** void **cvFillPoly**(CvArr* **img**, CvPoint** **pts**, const int* **npts**, int **contours**, CvScalar **color**, int **line_type**=8, int **shift**=0 )

**Python:** cv.**FillPoly**(img, polys, color, lineType=8, shift=0) → None

>       **Parameters**

>> **img** – Image.

>> **pts** – Array of polygons where each polygon is represented as an array of points.

>> **npts** – Array of polygon vertex counters.

>> **ncontours** – Number of contours that bind the filled region.

>> **color** – Polygon color.

>> **lineType** – Type of the polygon boundaries. See the `line()` description.

>> **shift** – Number of fractional bits in the vertex coordinates.

>> **offset** – Optional offset of all points of the contours.

The function `fillPoly` fills an area bounded by several polygonal contours. The function can fill complex areas, for example, areas with holes, contours with self-intersections (some of their parts), and so forth.

## getTextSize

Calculates the width and height of a text string.

**C++:** Size **getTextSize**(const string& **text**, int **fontFace**, double **fontScale**, int **thickness**, int* **baseLine**)

**Python:** cv2.**getTextSize**(text, fontFace, fontScale, thickness) → retval, baseLine

**C:** void **cvGetTextSize**(const char* **text_string**, const CvFont* **font**, CvSize* **text_size**, int* **baseline**)

**Python:** cv.**GetTextSize**(textString, font)-> (textSize, baseline)

> **Parameters**

> > **text** – Input text string.

> > **fontFace** – Font to use. See the `putText()` for details.

> > **fontScale** – Font scale. See the `putText()` for details.

> > **thickness** – Thickness of lines used to render the text. See `putText()` for details.

> > **baseLine** – Output parameter - y-coordinate of the baseline relative to the bottom-most text point.

The function `getTextSize` calculates and returns the size of a box that contains the specified text. That is, the following code renders some text, the tight box surrounding it, and the baseline:

```
// Use "y" to show that the baseLine is about
string text = "Funny text inside the box";
int fontFace = FONT_HERSHEY_SCRIPT_SIMPLEX;
double fontScale = 2;
int thickness = 3;

Mat img(600, 800, CV_8UC3, Scalar::all(0));

int baseline=0;
Size textSize = getTextSize(text, fontFace,
                            fontScale, thickness, &baseline);
baseline += thickness;

// center the text
Point textOrg((img.cols - textSize.width)/2,
              (img.rows + textSize.height)/2);

// draw the box
rectangle(img, textOrg + Point(0, baseline),
          textOrg + Point(textSize.width, -textSize.height),
          Scalar(0,0,255));
// ... and the baseline first
line(img, textOrg + Point(0, thickness),
     textOrg + Point(textSize.width, thickness),
     Scalar(0, 0, 255));

// then put the text itself
putText(img, text, textOrg, fontFace, fontScale,
        Scalar::all(255), thickness, 8);
```

## InitFont

Initializes font structure (OpenCV 1.x API).

**C:** void **cvInitFont**(CvFont* **font**, int **font_face**, double **hscale**, double **vscale**, double **shear**=0, int **thickness**=1, int **line_type**=8 )

> Parameters

>> **font** – Pointer to the font structure initialized by the function

>> **font_face** – Font name identifier. Only a subset of Hershey fonts http://sources.isc.org/utils/misc/hershey-font.txt are supported now:

>>> – **CV_FONT_HERSHEY_SIMPLEX** normal size sans-serif font

>>> – **CV_FONT_HERSHEY_PLAIN** small size sans-serif font

>>> – **CV_FONT_HERSHEY_DUPLEX** normal size sans-serif font (more complex than CV_FONT_HERSHEY_SIMPLEX )

>>> – **CV_FONT_HERSHEY_COMPLEX** normal size serif font

>>> – **CV_FONT_HERSHEY_TRIPLEX** normal size serif font (more complex than CV_FONT_HERSHEY_COMPLEX )

>>> – **CV_FONT_HERSHEY_COMPLEX_SMALL** smaller version of CV_FONT_HERSHEY_COMPLEX

>>> – **CV_FONT_HERSHEY_SCRIPT_SIMPLEX** hand-writing style font

>>> – **CV_FONT_HERSHEY_SCRIPT_COMPLEX** more complex variant of CV_FONT_HERSHEY_SCRIPT_SIMPLEX

>> The parameter can be composited from one of the values above and an optional CV_FONT_ITALIC flag, which indicates italic or oblique font.

>> **hscale** – Horizontal scale. If equal to 1.0f , the characters have the original width depending on the font type. If equal to 0.5f , the characters are of half the original width.

>> **vscale** – Vertical scale. If equal to 1.0f , the characters have the original height depending on the font type. If equal to 0.5f , the characters are of half the original height.

>> **shear** – Approximate tangent of the character slope relative to the vertical line. A zero value means a non-italic font, 1.0f means about a 45 degree slope, etc.

>> **thickness** – Thickness of the text strokes

>> **line_type** – Type of the strokes, see line() description

The function initializes the font structure that can be passed to text rendering functions.

**See Also:**

PutText()

# line

Draws a line segment connecting two points.

**C++:** void **line**(Mat& **img**, Point **pt1**, Point **pt2**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0)

**Python:** cv2.**line**(img, pt1, pt2, color[, thickness[, lineType[, shift]]]) → None

**C:** void **cvLine**(CvArr* **img**, CvPoint **pt1**, CvPoint **pt2**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int **shift**=0 )

**Python:** cv.**Line**(img, pt1, pt2, color, thickness=1, lineType=8, shift=0) → None

> **Parameters**

>> **img** – Image.

>> **pt1** – First point of the line segment.

>> **pt2** – Second point of the line segment.

>> **color** – Line color.

>> **thickness** – Line thickness.

>> **lineType** – Type of the line:

>>> – **8** (or omitted) - 8-connected line.

>>> – **4** - 4-connected line.

>>> – **CV_AA** - antialiased line.

>> **shift** – Number of fractional bits in the point coordinates.

The function `line` draws the line segment between `pt1` and `pt2` points in the image. The line is clipped by the image boundaries. For non-antialiased lines with integer coordinates, the 8-connected or 4-connected Bresenham algorithm is used. Thick lines are drawn with rounding endings. Antialiased lines are drawn using Gaussian filtering. To specify the line color, you may use the macro `CV_RGB(r, g, b)`.

## LineIterator

**class LineIterator**

Class for iterating pixels on a raster line.

```cpp
class LineIterator
{
public:
    // creates iterators for the line connecting pt1 and pt2
    // the line will be clipped on the image boundaries
    // the line is 8-connected or 4-connected
    // If leftToRight=true, then the iteration is always done
    // from the left-most point to the right most,
    // not to depend on the ordering of pt1 and pt2 parameters
    LineIterator(const Mat& img, Point pt1, Point pt2,
                 int connectivity=8, bool leftToRight=false);
    // returns pointer to the current line pixel
    uchar* operator *();
    // move the iterator to the next pixel
    LineIterator& operator ++();
    LineIterator operator ++(int);
    Point pos() const;

    // internal state of the iterator
    uchar* ptr;
    int err, count;
    int minusDelta, plusDelta;
    int minusStep, plusStep;
};
```

The class `LineIterator` is used to get each pixel of a raster line. It can be treated as versatile implementation of the Bresenham algorithm where you can stop at each pixel and do some extra processing, for example, grab pixel values along the line or draw a line with an effect (for example, with XOR operation).

The number of pixels along the line is stored in `LineIterator::count` . The method `LineIterator::pos` returns the current position in the image

```cpp
// grabs pixels along the line (pt1, pt2)
// from 8-bit 3-channel image to the buffer
LineIterator it(img, pt1, pt2, 8);
LineIterator it2 = it;
vector<Vec3b> buf(it.count);

for(int i = 0; i < it.count; i++, ++it)
    buf[i] = *(const Vec3b)*it;

// alternative way of iterating through the line
for(int i = 0; i < it2.count; i++, ++it2)
{
    Vec3b val = img.at<Vec3b>(it2.pos());
    CV_Assert(buf[i] == val);
}
```

## rectangle

Draws a simple, thick, or filled up-right rectangle.

**C++:** void **rectangle**(Mat& **img**, Point **pt1**, Point **pt2**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0)

**C++:** void **rectangle**(Mat& **img**, Rect **rec**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0 )

**Python:** cv2.**rectangle**(img, pt1, pt2, color$\left[\right.$, thickness$\left[\right.$, lineType$\left[\right.$, shift$\left.\right]\left.\right]\left.\right]$) → None

**C:** void **cvRectangle**(CvArr* **img**, CvPoint **pt1**, CvPoint **pt2**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int **shift**=0 )

**Python:** cv.**Rectangle**(img, pt1, pt2, color, thickness=1, lineType=8, shift=0) → None

> **Parameters**
>> **img** – Image.
>>
>> **pt1** – Vertex of the rectangle.
>>
>> **pt2** – Vertex of the rectangle opposite to `pt1` .
>>
>> **rec** – Alternative specification of the drawn rectangle.
>>
>> **color** – Rectangle color or brightness (grayscale image).
>>
>> **thickness** – Thickness of lines that make up the rectangle. Negative values, like `CV_FILLED` , mean that the function has to draw a filled rectangle.
>>
>> **lineType** – Type of the line. See the `line()` description.
>>
>> **shift** – Number of fractional bits in the point coordinates.

The function `rectangle` draws a rectangle outline or a filled rectangle whose two opposite corners are `pt1` and `pt2`, or `r.tl()` and `r.br()-Point(1,1)`.

## polylines

Draws several polygonal curves.

**C++:** void **polylines**(Mat& **img**, const Point** **pts**, const int* **npts**, int **ncontours**, bool **isClosed**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0 )

**C++:** void **polylines**(InputOutputArray **img**, InputArrayOfArrays **pts**, bool **isClosed**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, int **shift**=0 )

**Python:** cv2.**polylines**(img, pts, isClosed, color[, thickness[, lineType[, shift]]]) → None

**C:** void **cvPolyLine**(CvArr* **img**, CvPoint** **pts**, const int* **npts**, int **contours**, int **is_closed**, CvScalar **color**, int **thickness**=1, int **line_type**=8, int **shift**=0 )

**Python:** cv.**PolyLine**(img, polys, is_closed, color, thickness=1, lineType=8, shift=0) → None

> **Parameters**

> > **img** – Image.

> > **pts** – Array of polygonal curves.

> > **npts** – Array of polygon vertex counters.

> > **ncontours** – Number of curves.

> > **isClosed** – Flag indicating whether the drawn polylines are closed or not. If they are closed, the function draws a line from the last vertex of each curve to its first vertex.

> > **color** – Polyline color.

> > **thickness** – Thickness of the polyline edges.

> > **lineType** – Type of the line segments. See the line() description.

> > **shift** – Number of fractional bits in the vertex coordinates.

The function polylines draws one or more polygonal curves.

## putText

Draws a text string.

**C++:** void **putText**(Mat& **img**, const string& **text**, Point **org**, int **fontFace**, double **fontScale**, Scalar **color**, int **thickness**=1, int **lineType**=8, bool **bottomLeftOrigin**=false )

**Python:** cv2.**putText**(img, text, org, fontFace, fontScale, color[, thickness[, lineType[, bottomLeftOrigin]]]) → None

**C:** void **cvPutText**(CvArr* **img**, const char* **text**, CvPoint **org**, const CvFont* **font**, CvScalar **color**)

**Python:** cv.**PutText**(img, text, org, font, color) → None

> **Parameters**

> > **img** – Image.

> > **text** – Text string to be drawn.

> > **org** – Bottom-left corner of the text string in the image.

> > **font** – CvFont structure initialized using InitFont().

> > **fontFace** – Font type. One of FONT_HERSHEY_SIMPLEX, FONT_HERSHEY_PLAIN, FONT_HERSHEY_DUPLEX, FONT_HERSHEY_COMPLEX, FONT_HERSHEY_TRIPLEX, FONT_HERSHEY_COMPLEX_SMALL, FONT_HERSHEY_SCRIPT_SIMPLEX, or FONT_HERSHEY_SCRIPT_COMPLEX, where each of the font ID's can be combined with FONT_HERSHEY_ITALIC to get the slanted letters.

> > **fontScale** – Font scale factor that is multiplied by the font-specific base size.
> >
> > **color** – Text color.
> >
> > **thickness** – Thickness of the lines used to draw a text.
> >
> > **lineType** – Line type. See the `line` for details.
> >
> > **bottomLeftOrigin** – When true, the image data origin is at the bottom-left corner. Otherwise, it is at the top-left corner.

The function `putText` renders the specified text string in the image. Symbols that cannot be rendered using the specified font are replaced by question marks. See `getTextSize()` for a text rendering code example.

## 2.6 XML/YAML Persistence

### XML/YAML file storages. Writing to a file storage.

You can store and then restore various OpenCV data structures to/from XML (http://www.w3c.org/XML) or YAML (http://www.yaml.org) formats. Also, it is possible store and load arbitrarily complex data structures, which include OpenCV data structures, as well as primitive data types (integer and floating-point numbers and text strings) as their elements.

**Use the following procedure to write something to XML or YAML:**

1. Create new `FileStorage` and open it for writing. It can be done with a single call to `FileStorage::FileStorage()` constructor that takes a filename, or you can use the default constructor and then call `FileStorage::open()`. Format of the file (XML or YAML) is determined from the filename extension (".xml" and ".yml"/".yaml", respectively)

2. Write all the data you want using the streaming operator `>>`, just like in the case of STL streams.

3. Close the file using `FileStorage::release()`. `FileStorage` destructor also closes the file.

Here is an example:

```cpp
#include "opencv2/opencv.hpp"
#include <time.h>

using namespace cv;

int main(int, char** argv)
{
    FileStorage fs("test.yml", FileStorage::WRITE);

    fs << "frameCount" << 5;
    time_t rawtime; time(&rawtime);
    fs << "calibrationDate" << asctime(localtime(&rawtime));
    Mat cameraMatrix = (Mat_<double>(3,3) << 1000, 0, 320, 0, 1000, 240, 0, 0, 1);
    Mat distCoeffs = (Mat_<double>(5,1) << 0.1, 0.01, -0.001, 0, 0);
    fs << "cameraMatrix" << cameraMatrix << "distCoeffs" << distCoeffs;
    fs << "features" << "[";
    for( int i = 0; i < 3; i++ )
    {
        int x = rand() % 640;
        int y = rand() % 480;
        uchar lbp = rand() % 256;

        fs << "{:" << "x" << x << "y" << y << "lbp" << "[:";
```

```
        for( int j = 0; j < 8; j++ )
            fs << ((lbp >> j) & 1);
        fs << "]" << "}";
    }
    fs << "]";
    fs.release();
    return 0;
}
```

The sample above stores to XML and integer, text string (calibration date), 2 matrices, and a custom structure "feature", which includes feature coordinates and LBP (local binary pattern) value. Here is output of the sample:

```
%YAML:1.0
frameCount: 5
calibrationDate: "Fri Jun 17 14:09:29 2011\n"
cameraMatrix: !!opencv-matrix
   rows: 3
   cols: 3
   dt: d
   data: [ 1000., 0., 320., 0., 1000., 240., 0., 0., 1. ]
distCoeffs: !!opencv-matrix
   rows: 5
   cols: 1
   dt: d
   data: [ 1.0000000000000001e-01, 1.0000000000000000e-02,
       -1.0000000000000000e-03, 0., 0. ]
features:
   - { x:167, y:49, lbp:[ 1, 0, 0, 1, 1, 0, 1, 1 ] }
   - { x:298, y:130, lbp:[ 0, 0, 0, 1, 0, 0, 1, 1 ] }
   - { x:344, y:158, lbp:[ 1, 1, 0, 0, 0, 0, 1, 0 ] }
```

As an exercise, you can replace ".yml" with ".xml" in the sample above and see, how the corresponding XML file will look like.

**Several things can be noted by looking at the sample code and the output:**

- The produced YAML (and XML) consists of heterogeneous collections that can be nested. There are 2 types of collections: named collections (mappings) and unnamed collections (sequences). In mappings each element has a name and is accessed by name. This is similar to structures and `std::map` in C/C++ and dictionaries in Python. In sequences elements do not have names, they are accessed by indices. This is similar to arrays and `std::vector` in C/C++ and lists, tuples in Python. "Heterogeneous" means that elements of each single collection can have different types.

  Top-level collection in YAML/XML is a mapping. Each matrix is stored as a mapping, and the matrix elements are stored as a sequence. Then, there is a sequence of features, where each feature is represented a mapping, and lbp value in a nested sequence.

- When you write to a mapping (a structure), you write element name followed by its value. When you write to a sequence, you simply write the elements one by one. OpenCV data structures (such as cv::Mat) are written in absolutely the same way as simple C data structures - using "**<<**" operator.

- To write a mapping, you first write the special string "**{**" to the storage, then write the elements as pairs (`fs << <element_name> << <element_value>`) and then write the closing "**}**".

- To write a sequence, you first write the special string "**[**", then write the elements, then write the closing "**]**".

- In YAML (but not XML), mappings and sequences can be written in a compact Python-like inline form. In the sample above matrix elements, as well as each feature, including its lbp value, is stored in such inline

form. To store a mapping/sequence in a compact form, put ":" after the opening character, e.g. use **"{:"** instead of **"{"** and **"[:"** instead of **"["**. When the data is written to XML, those extra ":" are ignored.

## Reading data from a file storage.

To read the previously written XML or YAML file, do the following:

1. Open the file storage using `FileStorage::FileStorage()` constructor or `FileStorage::open()` method. In the current implementation the whole file is parsed and the whole representation of file storage is built in memory as a hierarchy of file nodes (see `FileNode`)

2. Read the data you are interested in. Use `FileStorage::operator [](),` `FileNode::operator [](}` and/or `FileNodeIterator`.

3. Close the storage using `FileStorage::release()`.

Here is how to read the file created by the code sample above:

```cpp
FileStorage fs2("test.yml", FileStorage::READ);

// first method: use (type) operator on FileNode.
int frameCount = (int)fs2["frameCount"];

std::string date;
// second method: use FileNode::operator >>
fs2["calibrationDate"] >> date;

Mat cameraMatrix2, distCoeffs2;
fs2["cameraMatrix"] >> cameraMatrix2;
fs2["distCoeffs"] >> distCoeffs2;

cout << "frameCount: " << frameCount << endl
     << "calibration date: " << date << endl
     << "camera matrix: " << cameraMatrix2 << endl
     << "distortion coeffs: " << distCoeffs2 << endl;

FileNode features = fs2["features"];
FileNodeIterator it = features.begin(), it_end = features.end();
int idx = 0;
std::vector<uchar> lbpval;

// iterate through a sequence using FileNodeIterator
for( ; it != it_end; ++it, idx++ )
{
    cout << "feature #" << idx << ": ";
    cout << "x=" << (int)(*it)["x"] << ", y=" << (int)(*it)["y"] << ", lbp: (";
    // you can also easily read numerical arrays using FileNode >> std::vector operator.
    (*it)["lbp"] >> lbpval;
    for( int i = 0; i < (int)lbpval.size(); i++ )
        cout << " " << (int)lbpval[i];
    cout << ")" << endl;
}
fs.release();
```

## FileStorage

**class** `FileStorage`

---

XML/YAML file storage class that encapsulates all the information necessary for writing or reading data to/from a file.

## FileStorage::FileStorage

The constructors.

**C++:** FileStorage::**FileStorage**()

**C++:** FileStorage::**FileStorage**(const string& **source**, int **flags**, const string& **encoding**=string())

>    **Parameters**

>>    **source** – Name of the file to open or the text string to read the data from. Extension of the file (.xml or .yml/.yaml) determines its format (XML or YAML respectively). Also you can append .gz to work with compressed files, for example myHugeMatrix.xml.gz. If both FileStorage::WRITE and FileStorage::MEMORY flags are specified, source is used just to specify the output file format (e.g. mydata.xml, .yml etc.).

>>    **flags** – Mode of operation. Possible values are:

>>>    – **FileStorage::READ** Open the file for reading.

>>>    – **FileStorage::WRITE** Open the file for writing.

>>>    – **FileStorage::APPEND** Open the file for appending.

>>>    – **FileStorage::MEMORY** Read data from source or write data to the internal buffer (which is returned by FileStorage::release)

>>    **encoding** – Encoding of the file. Note that UTF-16 XML encoding is not supported currently and you should use 8-bit encoding instead of it.

The full constructor opens the file. Alternatively you can use the default constructor and then call FileStorage::open().

## FileStorage::open

Opens a file.

**C++:** bool FileStorage::**open**(const string& **filename**, int **flags**, const string& **encoding**=string())

See description of parameters in FileStorage::FileStorage(). The method calls FileStorage::release() before opening the file.

## FileStorage::isOpened

Checks whether the file is opened.

**C++:** bool FileStorage::**isOpened**() const

>    **Returns** true if the object is associated with the current file and false otherwise.

It is a good practice to call this method after you tried to open a file.

## FileStorage::release

Closes the file and releases all the memory buffers.

**C++:** void FileStorage::**release**()

Call this method after all I/O operations with the storage are finished.

## FileStorage::releaseAndGetString

Closes the file and releases all the memory buffers.

**C++:** string FileStorage::**releaseAndGetString**()

Call this method after all I/O operations with the storage are finished. If the storage was opened for writing data and FileStorage::WRITE was specified

## FileStorage::getFirstTopLevelNode

Returns the first element of the top-level mapping.

**C++:** FileNode FileStorage::**getFirstTopLevelNode**() const

> **Returns** The first element of the top-level mapping.

## FileStorage::root

Returns the top-level mapping

**C++:** FileNode FileStorage::**root**(int **streamidx**=0) const

> **Parameters**
>
> > **streamidx** – Zero-based index of the stream. In most cases there is only one stream in the file. However, YAML supports multiple streams and so there can be several.
>
> **Returns** The top-level mapping.

## FileStorage::operator[]

Returns the specified element of the top-level mapping.

**C++:** FileNode FileStorage::**operator[]**(const string& **nodename**) const

**C++:** FileNode FileStorage::**operator[]**(const char* **nodename**) const

> **Parameters**
>
> > **nodename** – Name of the file node.
>
> **Returns** Node with the given name.

## FileStorage::operator*

Returns the obsolete C FileStorage structure.

**C++:** CvFileStorage* FileStorage::**operator***()

**C++:** const CvFileStorage* FileStorage::**operator***() const

>   **Returns**  Pointer to the underlying C FileStorage structure

## FileStorage::writeRaw

Writes multiple numbers.

**C++:** void FileStorage::**writeRaw**(const string& **fmt**, const uchar* **vec**, size_t **len**)

>   **Parameters**
>
>>   **fmt** – Specification of each array element that has the following format
>>   (`[count]{'u'|'c'|'w'|'s'|'i'|'f'|'d'}`)... where the characters correspond
>>   to fundamental C++ types:
>>
>>   – **u** 8-bit unsigned number
>>
>>   – **c** 8-bit signed number
>>
>>   – **w** 16-bit unsigned number
>>
>>   – **s** 16-bit signed number
>>
>>   – **i** 32-bit signed number
>>
>>   – **f** single precision floating-point number
>>
>>   – **d** double precision floating-point number
>>
>>   – **r** pointer, 32 lower bits of which are written as a signed integer. The type can be used to
>>   store structures with links between the elements.
>>
>>   `count` is the optional counter of values of a given type. For example, `2if` means that
>>   each array element is a structure of 2 integers, followed by a single-precision floating-point
>>   number. The equivalent notations of the above specification are ' `iif` ', ' `2i1f` ' and so
>>   forth. Other examples: `u` means that the array consists of bytes, and `2d` means the array
>>   consists of pairs of doubles.
>>
>>   **vec** – Pointer to the written array.
>>
>>   **len** – Number of the `uchar` elements to write.

Writes one or more numbers of the specified format to the currently written structure. Usually it is more convenient to
use `operator <<()` instead of this method.

## FileStorage::writeObj

Writes the registered C structure (CvMat, CvMatND, CvSeq).

**C++:** void FileStorage::**writeObj**(const string& **name**, const void* **obj**)

>   **Parameters**
>
>>   **name** – Name of the written object.
>>
>>   **obj** – Pointer to the object.

See `Write()` for details.

## FileStorage::getDefaultObjectName

Returns the normalized object name for the specified name of a file.

**C++:** **static** string FileStorage::**getDefaultObjectName**(const string& **filename**)

> **Parameters**
>
> > **filename** – Name of a file
>
> **Returns** The normalized object name.

## operator <<

Writes data to a file storage.

**C++:** template<typename _Tp> FileStorage& **operator<<**(FileStorage& **fs**, const _Tp& **value**)

**C++:** template<typename _Tp> FileStorage& **operator<<**(FileStorage& **fs**, const vector<_Tp>& **vec**)

> **Parameters**
>
> > **fs** – Opened file storage to write data.
> >
> > **value** – Value to be written to the file storage.
> >
> > **vec** – Vector of values to be written to the file storage.

It is the main function to write data to a file storage. See an example of its usage at the beginning of the section.

## operator >>

Reads data from a file storage.

**C++:** template<typename _Tp> void **operator>>**(const FileNode& **n**, _Tp& **value**)

**C++:** template<typename _Tp> void **operator>>**(const FileNode& **n**, vector<_Tp>& **vec**)

**C++:** template<typename _Tp> FileNodeIterator& **operator>>**(FileNodeIterator& **it**, _Tp& **value**)

**C++:** template<typename _Tp> FileNodeIterator& **operator>>**(FileNodeIterator& **it**, vector<_Tp>& **vec**)

> **Parameters**
>
> > **n** – Node from which data will be read.
> >
> > **it** – Iterator from which data will be read.
> >
> > **value** – Value to be read from the file storage.
> >
> > **vec** – Vector of values to be read from the file storage.

It is the main function to read data from a file storage. See an example of its usage at the beginning of the section.

## FileNode

**class** `FileNode`

File Storage Node class. The node is used to store each and every element of the file storage opened for reading. When XML/YAML file is read, it is first parsed and stored in the memory as a hierarchical collection of nodes. Each node can be a "leaf" that is contain a single number or a string, or be a collection of other nodes. There can be named collections (mappings) where each element has a name and it is accessed by a name, and ordered collections (sequences) where elements do not have names but rather accessed by index. Type of the file node can be determined using `FileNode::type()` method.

Note that file nodes are only used for navigating file storages opened for reading. When a file storage is opened for writing, no data is stored in memory after it is written.

## FileNode::FileNode

The constructors.

**C++:** `FileNode::`**`FileNode`**`()`

**C++:** `FileNode::`**`FileNode`**`(const CvFileStorage*` **fs**`, const CvFileNode*` **node**`)`

**C++:** `FileNode::`**`FileNode`**`(const FileNode&` **node**`)`

> **Parameters**
>
> > **fs** – Pointer to the obsolete file storage structure.
> >
> > **node** – File node to be used as initialization for the created file node.

These constructors are used to create a default file node, construct it from obsolete structures or from the another file node.

## FileNode::operator[]

Returns element of a mapping node or a sequence node.

**C++:** `FileNode FileNode::`**`operator[]`**`(const string&` **nodename**`) const`

**C++:** `FileNode FileNode::`**`operator[]`**`(const char*` **nodename**`) const`

**C++:** `FileNode FileNode::`**`operator[]`**`(int` **i**`) const`

> **Parameters**
>
> > **nodename** – Name of an element in the mapping node.
> >
> > **i** – Index of an element in the sequence node.
>
> **Returns** Returns the element with the given identifier.

## FileNode::type

Returns type of the node.

**C++:** `int FileNode::`**`type`**`() const`

> **Returns**
>
> > Type of the node. Possible values are:

- **FileNode::NONE** Empty node.
- **FileNode::INT** Integer.
- **FileNode::REAL** Floating-point number.
- **FileNode::FLOAT** Synonym or REAL.
- **FileNode::STR** Text string in UTF-8 encoding.
- **FileNode::STRING** Synonym for STR.
- **FileNode::REF** Integer of type size_t. Typically used for storing complex dynamic structures where some elements reference the others.
- **FileNode::SEQ** Sequence.
- **FileNode::MAP** Mapping.
- **FileNode::FLOW** Compact representation of a sequence or mapping. Used only by the YAML writer.
- **FileNode::USER** Registered object (e.g. a matrix).
- **FileNode::EMPTY** Empty structure (sequence or mapping).
- **FileNode::NAMED** The node has a name (i.e. it is an element of a mapping).

## FileNode::empty

Checks whether the node is empty.

**C++:** bool FileNode::**empty**() const

> **Returns** true if the node is empty.

## FileNode::isNone

Checks whether the node is a "none" object

**C++:** bool FileNode::**isNone**() const

> **Returns** true if the node is a "none" object.

## FileNode::isSeq

Checks whether the node is a sequence.

**C++:** bool FileNode::**isSeq**() const

> **Returns** true if the node is a sequence.

## FileNode::isMap

Checks whether the node is a mapping.

**C++:** bool FileNode::**isMap**() const

> **Returns** true if the node is a mapping.

### FileNode::isInt

Checks whether the node is an integer.

**C++:** bool FileNode::**isInt**() const

> **Returns** true if the node is an integer.

### FileNode::isReal

Checks whether the node is a floating-point number.

**C++:** bool FileNode::**isReal**() const

> **Returns** true if the node is a floating-point number.

### FileNode::isString

Checks whether the node is a text string.

**C++:** bool FileNode::**isString**() const

> **Returns** true if the node is a text string.

### FileNode::isNamed

Checks whether the node has a name.

**C++:** bool FileNode::**isNamed**() const

> **Returns** true if the node has a name.

### FileNode::name

Returns the node name.

**C++:** string FileNode::**name**() const

> **Returns** The node name or an empty string if the node is nameless.

### FileNode::size

Returns the number of elements in the node.

**C++:** size_t FileNode::**size**() const

> **Returns** The number of elements in the node, if it is a sequence or mapping, or 1 otherwise.

### FileNode::operator int

Returns the node content as an integer.

**C++:** FileNode::**operator** int() const

> **Returns** The node content as an integer. If the node stores a floating-point number, it is rounded.

---

## FileNode::operator float

Returns the node content as float.

**C++:** FileNode::**operator** float() const

> **Returns** The node content as float.

## FileNode::operator double

Returns the node content as double.

**C++:** FileNode::**operator** double() const

> **Returns** The node content as double.

## FileNode::operator string

Returns the node content as text string.

**C++:** FileNode::**operator** string() const

> **Returns** The node content as a text string.

## FileNode::operator*

Returns pointer to the underlying obsolete file node structure.

**C++:** CvFileNode* FileNode::**operator***()

> **Returns** Pointer to the underlying obsolete file node structure.

## FileNode::begin

Returns the iterator pointing to the first node element.

**C++:** FileNodeIterator FileNode::**begin**() const

> **Returns** Iterator pointing to the first node element.

## FileNode::end

Returns the iterator pointing to the element following the last node element.

**C++:** FileNodeIterator FileNode::**end**() const

> **Returns** Iterator pointing to the element following the last node element.

## FileNode::readRaw

Reads node elements to the buffer with the specified format.

**C++:** void FileNode::**readRaw**(const string& **fmt**, uchar* **vec**, size_t **len**) const

> **Parameters**

>
>> **fmt** – Specification of each array element. It has the same format as in `FileStorage::writeRaw()`.
>>
>> **vec** – Pointer to the destination array.
>>
>> **len** – Number of elements to read. If it is greater than number of remaining elements then all of them will be read.

Usually it is more convenient to use `operator >>()` instead of this method.

## FileNode::readObj

Reads the registered object.

**C++:** void* FileNode::**readObj**() const

>> **Returns** Pointer to the read object.

See `Read()` for details.

## FileNodeIterator

class **FileNodeIterator**

The class `FileNodeIterator` is used to iterate through sequences and mappings. A standard STL notation, with `node.begin()`, `node.end()` denoting the beginning and the end of a sequence, stored in `node`. See the data reading sample in the beginning of the section.

## FileNodeIterator::FileNodeIterator

The constructors.

**C++:** FileNodeIterator::**FileNodeIterator**()

**C++:** FileNodeIterator::**FileNodeIterator**(const CvFileStorage* **fs**, const CvFileNode* **node**, size_t **ofs**=0)

**C++:** FileNodeIterator::**FileNodeIterator**(const FileNodeIterator& **it**)

>> **Parameters**
>>
>>> **fs** – File storage for the iterator.
>>>
>>> **node** – File node for the iterator.
>>>
>>> **ofs** – Index of the element in the node. The created iterator will point to this element.
>>>
>>> **it** – Iterator to be used as initialization for the created iterator.

These constructors are used to create a default iterator, set it to specific element in a file node or construct it from another iterator.

## FileNodeIterator::operator*

Returns the currently observed element.

**C++:** FileNode FileNodeIterator::**operator***() const

>> **Returns** Currently observed element.

---

## FileNodeIterator::operator->

Accesses methods of the currently observed element.

**C++:** FileNode FileNodeIterator::**operator->**() const

## FileNodeIterator::operator ++

Moves iterator to the next node.

**C++:** FileNodeIterator& FileNodeIterator::**operator++**()

**C++:** FileNodeIterator FileNodeIterator::**operator++**(int **None**)

## FileNodeIterator::operator –

Moves iterator to the previous node.

**C++:** FileNodeIterator& FileNodeIterator::**operator-**()

**C++:** FileNodeIterator FileNodeIterator::**operator-**(int **None**)

## FileNodeIterator::operator +=

Moves iterator forward by the specified offset.

**C++:** FileNodeIterator& FileNodeIterator::**operator+=**(int **ofs**)

> **Parameters**
>
> > **ofs** – Offset (possibly negative) to move the iterator.

## FileNodeIterator::operator -=

Moves iterator backward by the specified offset (possibly negative).

**C++:** FileNodeIterator& FileNodeIterator::**operator-=**(int **ofs**)

> **Parameters**
>
> > **ofs** – Offset (possibly negative) to move the iterator.

## FileNodeIterator::readRaw

Reads node elements to the buffer with the specified format.

**C++:** FileNodeIterator& FileNodeIterator::**readRaw**(const string& **fmt**, uchar* **vec**, size_t **max-Count**=(size_t)INT_MAX )

> **Parameters**
>
> > **fmt** – Specification of each array element.  It has the same format as in `FileStorage::writeRaw()`.
> >
> > **vec** – Pointer to the destination array.
> >
> > **maxCount** – Number of elements to read. If it is greater than number of remaining elements then all of them will be read.

Usually it is more convenient to use `operator >>()` instead of this method.

# 2.7 XML/YAML Persistence (C API)

The section describes the OpenCV 1.x API for reading and writing data structures to/from XML or YAML files. It is now recommended to use the new C++ interface for reading and writing data.

## CvFileStorage

**struct `CvFileStorage`**

The structure `CvFileStorage` is a "black box" representation of the file storage associated with a file on disk. Several functions that are described below take `CvFileStorage*` as inputs and allow the user to save or to load hierarchical collections that consist of scalar values, standard CXCore objects (such as matrices, sequences, graphs), and user-defined objects.

OpenCV can read and write data in XML (http://www.w3c.org/XML) or YAML (http://www.yaml.org) formats. Below is an example of 3x3 floating-point identity matrix A, stored in XML and YAML files using CXCore functions:

XML:

```
<?xml version="1.0">
<opencv_storage>
<A type_id="opencv-matrix">
  <rows>3</rows>
  <cols>3</cols>
  <dt>f</dt>
  <data>1. 0. 0. 0. 1. 0. 0. 0. 1.</data>
</A>
</opencv_storage>
```

YAML:

```
%YAML:1.0
A: !!opencv-matrix
  rows: 3
  cols: 3
  dt: f
  data: [ 1., 0., 0., 0., 1., 0., 0., 0., 1.]
```

As it can be seen from the examples, XML uses nested tags to represent hierarchy, while YAML uses indentation for that purpose (similar to the Python programming language).

The same functions can read and write data in both formats; the particular format is determined by the extension of the opened file, ".xml" for XML files and ".yml" or ".yaml" for YAML.

## CvFileNode

**struct `CvFileNode`**

> File storage node. When XML/YAML file is read, it is first parsed and stored in the memory as a hierarchical collection of nodes. Each node can be a "leaf", that is, contain a single number or a string, or be a collection of other nodes. Collections are also referenced to as "structures" in the data writing functions. There can be named collections (mappings), where each element has a name and is accessed by a name, and ordered collections (sequences), where elements do not have names, but rather accessed by index.

int **tag**
>    type of the file node:

>    - CV_NODE_NONE - empty node

>    - CV_NODE_INT - an integer

>    - CV_NODE_REAL - a floating-point number

>    - CV_NODE_STR - text string

>    - CV_NODE_SEQ - a sequence

>    - CV_NODE_MAP - a mapping

>    type of the node can be retrieved using `CV_NODE_TYPE(node->tag)` macro.

CvTypeInfo* **info**
>    optional pointer to the user type information. If you look at the matrix representation in XML and YAML, shown above, you may notice `type_id="opencv-matrix"` or `!!opencv-matrix` strings. They are used to specify that the certain element of a file is a representation of a data structure of certain type ("opencv-matrix" corresponds to `CvMat`). When a file is parsed, such type identifiers are passed to `FindType()` to find type information and the pointer to it is stored in the file node. See `CvTypeInfo` for more details.

union **data**
>    the node data, declared as:

```
union
{
    double f; /* scalar floating-point number */
    int i;    /* scalar integer number */
    CvString str; /* text string */
    CvSeq* seq; /* sequence (ordered collection of file nodes) */
    struct CvMap* map; /* map (collection of named file nodes) */
} data;
```

>    Primitive nodes are read using `ReadInt()`, `ReadReal()` and `ReadString()`. Sequences are read by iterating through `node->data.seq` (see "Dynamic Data Structures" section). Mappings are read using `GetFileNodeByName()`. Nodes with the specified type (so that `node->info != NULL`) can be read using `Read()`.

## CvAttrList

**struct CvAttrList**

List of attributes.

```
typedef struct CvAttrList
{
    const char** attr; /* NULL-terminated array of (attribute_name,attribute_value) pairs */
    struct CvAttrList* next; /* pointer to next chunk of the attributes list */
}
CvAttrList;

/* initializes CvAttrList structure */
inline CvAttrList cvAttrList( const char** attr=NULL, CvAttrList* next=NULL );

/* returns attribute value or 0 (NULL) if there is no such attribute */
const char* cvAttrValue( const CvAttrList* attr, const char* attr_name );
```

In the current implementation, attributes are used to pass extra parameters when writing user objects (see `Write()`).
XML attributes inside tags are not supported, aside from the object type specification (`type_id` attribute).

## CvTypeInfo

**struct CvTypeInfo**

Type information.

```
typedef int (CV_CDECL *CvIsInstanceFunc)( const void* structPtr );
typedef void (CV_CDECL *CvReleaseFunc)( void** structDblPtr );
typedef void* (CV_CDECL *CvReadFunc)( CvFileStorage* storage, CvFileNode* node );
typedef void (CV_CDECL *CvWriteFunc)( CvFileStorage* storage,
                                      const char* name,
                                      const void* structPtr,
                                      CvAttrList attributes );
typedef void* (CV_CDECL *CvCloneFunc)( const void* structPtr );


typedef struct CvTypeInfo
{
    int flags; /* not used */
    int header_size; /* sizeof(CvTypeInfo) */
    struct CvTypeInfo* prev; /* previous registered type in the list */
    struct CvTypeInfo* next; /* next registered type in the list */
    const char* type_name; /* type name, written to file storage */

    /* methods */
    CvIsInstanceFunc is_instance; /* checks if the passed object belongs to the type */
    CvReleaseFunc release; /* releases object (memory etc.) */
    CvReadFunc read; /* reads object from file storage */
    CvWriteFunc write; /* writes object to file storage */
    CvCloneFunc clone; /* creates a copy of the object */
}
CvTypeInfo;
```

The structure contains information about one of the standard or user-defined types. Instances of the type may or
may not contain a pointer to the corresponding `CvTypeInfo` structure. In any case, there is a way to find the type
info structure for a given object using the `TypeOf()` function. Alternatively, type info can be found by type name
using `FindType()`, which is used when an object is read from file storage. The user can register a new type with
`RegisterType()` that adds the type information structure into the beginning of the type list. Thus, it is possible to
create specialized types from generic standard types and override the basic methods.

## Clone

Makes a clone of an object.

**C:** void* **cvClone**(const void* **struct_ptr**)

> **Parameters**
>
> > **struct_ptr** – The object to clone

The function finds the type of a given object and calls `clone` with the passed object. Of course, if you know the object
type, for example, `struct_ptr` is `CvMat*`, it is faster to call the specific function, like `CloneMat()`.

## EndWriteStruct

Finishes writing to a file node collection.

**C:** void **cvEndWriteStruct**(CvFileStorage* **fs**)

> **Parameters**
>
> > **fs** – File storage

**See Also:**

StartWriteStruct().

## FindType

Finds a type by its name.

**C:** CvTypeInfo* **cvFindType**(const char* **type_name**)

> **Parameters**
>
> > **type_name** – Type name

The function finds a registered type by its name. It returns NULL if there is no type with the specified name.

## FirstType

Returns the beginning of a type list.

**C:** CvTypeInfo* **cvFirstType**(void **None**)

The function returns the first type in the list of registered types. Navigation through the list can be done via the `prev` and `next` fields of the CvTypeInfo structure.

## GetFileNode

Finds a node in a map or file storage.

**C:** CvFileNode* **cvGetFileNode**(CvFileStorage* **fs**, CvFileNode* **map**, const CvStringHashNode* **key**, int **create_missing**=0 )

> **Parameters**
>
> > **fs** – File storage
> >
> > **map** – The parent map. If it is NULL, the function searches a top-level node. If both `map` and `key` are NULLs, the function returns the root file node - a map that contains top-level nodes.
> >
> > **key** – Unique pointer to the node name, retrieved with GetHashedKey()
> >
> > **create_missing** – Flag that specifies whether an absent node should be added to the map

The function finds a file node. It is a faster version of GetFileNodeByName() (see GetHashedKey() discussion). Also, the function can insert a new node, if it is not in the map yet.

## GetFileNodeByName

Finds a node in a map or file storage.

**C:** CvFileNode* **cvGetFileNodeByName**(const CvFileStorage* **fs**, const CvFileNode* **map**, const char* **name**)

> **Parameters**
>
> > **fs** – File storage
> >
> > **map** – The parent map. If it is NULL, the function searches in all the top-level nodes (streams), starting with the first one.
> >
> > **name** – The file node name

The function finds a file node by `name`. The node is searched either in `map` or, if the pointer is NULL, among the top-level file storage nodes. Using this function for maps and `GetSeqElem()` (or sequence reader) for sequences, it is possible to navigate through the file storage. To speed up multiple queries for a certain key (e.g., in the case of an array of structures) one may use a combination of `GetHashedKey()` and `GetFileNode()`.

## GetFileNodeName

Returns the name of a file node.

**C:** const char* **cvGetFileNodeName**(const CvFileNode* **node**)

> **Parameters**
>
> > **node** – File node

The function returns the name of a file node or NULL, if the file node does not have a name or if `node` is NULL.

## GetHashedKey

Returns a unique pointer for a given name.

**C:** CvStringHashNode* **cvGetHashedKey**(CvFileStorage* **fs**, const char* **name**, int **len**=-1, int **create_missing**=0 )

> **Parameters**
>
> > **fs** – File storage
> >
> > **name** – Literal node name
> >
> > **len** – Length of the name (if it is known apriori), or -1 if it needs to be calculated
> >
> > **create_missing** – Flag that specifies, whether an absent key should be added into the hash table

The function returns a unique pointer for each particular file node name. This pointer can be then passed to the `GetFileNode()` function that is faster than `GetFileNodeByName()` because it compares text strings by comparing pointers rather than the strings' content.

Consider the following example where an array of points is encoded as a sequence of 2-entry maps:

```
points:
  - { x: 10, y: 10 }
  - { x: 20, y: 20 }
  - { x: 30, y: 30 }
  # ...
```

Then, it is possible to get hashed "x" and "y" pointers to speed up decoding of the points.

```c
#include "cxcore.h"

int main( int argc, char** argv )
{
    CvFileStorage* fs = cvOpenFileStorage( "points.yml", 0, CV_STORAGE_READ );
    CvStringHashNode* x_key = cvGetHashedNode( fs, "x", -1, 1 );
    CvStringHashNode* y_key = cvGetHashedNode( fs, "y", -1, 1 );
    CvFileNode* points = cvGetFileNodeByName( fs, 0, "points" );

    if( CV_NODE_IS_SEQ(points->tag) )
    {
        CvSeq* seq = points->data.seq;
        int i, total = seq->total;
        CvSeqReader reader;
        cvStartReadSeq( seq, &reader, 0 );
        for( i = 0; i < total; i++ )
        {
            CvFileNode* pt = (CvFileNode*)reader.ptr;
#if 1 /* faster variant */
            CvFileNode* xnode = cvGetFileNode( fs, pt, x_key, 0 );
            CvFileNode* ynode = cvGetFileNode( fs, pt, y_key, 0 );
            assert( xnode && CV_NODE_IS_INT(xnode->tag) &&
                    ynode && CV_NODE_IS_INT(ynode->tag));
            int x = xnode->data.i; // or x = cvReadInt( xnode, 0 );
            int y = ynode->data.i; // or y = cvReadInt( ynode, 0 );
#elif 1 /* slower variant; does not use x_key & y_key */
            CvFileNode* xnode = cvGetFileNodeByName( fs, pt, "x" );
            CvFileNode* ynode = cvGetFileNodeByName( fs, pt, "y" );
            assert( xnode && CV_NODE_IS_INT(xnode->tag) &&
                    ynode && CV_NODE_IS_INT(ynode->tag));
            int x = xnode->data.i; // or x = cvReadInt( xnode, 0 );
            int y = ynode->data.i; // or y = cvReadInt( ynode, 0 );
#else /* the slowest yet the easiest to use variant */
            int x = cvReadIntByName( fs, pt, "x", 0 /* default value */ );
            int y = cvReadIntByName( fs, pt, "y", 0 /* default value */ );
#endif
            CV_NEXT_SEQ_ELEM( seq->elem_size, reader );
            printf("
        }
    }
    cvReleaseFileStorage( &fs );
    return 0;
}
```

Please note that whatever method of accessing a map you are using, it is still much slower than using plain sequences; for example, in the above example, it is more efficient to encode the points as pairs of integers in a single numeric sequence.

## GetRootFileNode

Retrieves one of the top-level nodes of the file storage.

**C:** CvFileNode* **cvGetRootFileNode**(const CvFileStorage* **fs**, int **stream_index**=0 )

> **Parameters**

> > **fs** – File storage

> **stream_index** – Zero-based index of the stream. See `StartNextStream()` . In most cases, there is only one stream in the file; however, there can be several.

The function returns one of the top-level file nodes. The top-level nodes do not have a name, they correspond to the streams that are stored one after another in the file storage. If the index is out of range, the function returns a NULL pointer, so all the top-level nodes can be iterated by subsequent calls to the function with `stream_index=0,1,...,` until the NULL pointer is returned. This function can be used as a base for recursive traversal of the file storage.

## Load

Loads an object from a file.

**C:** void* **cvLoad**(const char* **filename**, CvMemStorage* **memstorage**=NULL, const char* **name**=NULL, const char** **real_name**=NULL )

**Python:** cv.**Load**(filename, storage=None, name=None) → generic

> **Parameters**
>
> > **filename** – File name
> >
> > **memstorage** – Memory storage for dynamic structures, such as `CvSeq` or `CvGraph` . It is not used for matrices or images.
> >
> > **name** – Optional object name. If it is NULL, the first top-level object in the storage will be loaded.
> >
> > **real_name** – Optional output parameter that will contain the name of the loaded object (useful if `name=NULL` )

The function loads an object from a file. It basically reads the specified file, find the first top-level node and calls `Read()` for that node. If the file node does not have type information or the type information can not be found by the type name, the function returns NULL. After the object is loaded, the file storage is closed and all the temporary buffers are deleted. Thus, to load a dynamic structure, such as a sequence, contour, or graph, one should pass a valid memory storage destination to the function.

## OpenFileStorage

Opens file storage for reading or writing data.

**C:** CvFileStorage* **cvOpenFileStorage**(const char* **filename**, CvMemStorage* **memstorage**, int **flags**, const char* **encoding**=NULL )

> **Parameters**
>
> > **filename** – Name of the file associated with the storage
> >
> > **memstorage** – Memory storage used for temporary data and for storing dynamic structures, such as `CvSeq` or `CvGraph` . If it is NULL, a temporary memory storage is created and used.
> >
> > **flags** – Can be one of the following:
> >
> > – **CV_STORAGE_READ** the storage is open for reading
> >
> > – **CV_STORAGE_WRITE** the storage is open for writing

The function opens file storage for reading or writing data. In the latter case, a new file is created or an existing file is rewritten. The type of the read or written file is determined by the filename extension: `.xml` for XML and `.yml` or `.yaml` for YAML. The function returns a pointer to the `CvFileStorage` structure. If the file cannot be opened then the function returns NULL.

## Read

Decodes an object and returns a pointer to it.

**C:** void* **cvRead**(CvFileStorage* **fs**, CvFileNode* **node**, CvAttrList* **attributes**=NULL )

> **Parameters**
>
> > **fs** – File storage
> >
> > **node** – The root object node
> >
> > **attributes** – Unused parameter

The function decodes a user object (creates an object in a native representation from the file storage subtree) and returns it. The object to be decoded must be an instance of a registered type that supports the read method (see `CvTypeInfo`). The type of the object is determined by the type name that is encoded in the file. If the object is a dynamic structure, it is created either in memory storage and passed to `OpenFileStorage()` or, if a NULL pointer was passed, in temporary memory storage, which is released when `ReleaseFileStorage()` is called. Otherwise, if the object is not a dynamic structure, it is created in a heap and should be released with a specialized function or by using the generic `Release()`.

## ReadByName

Finds an object by name and decodes it.

**C:** void* **cvReadByName**(CvFileStorage* **fs**, const CvFileNode* **map**, const char* **name**, CvAttrList* **attributes**=NULL )

> **Parameters**
>
> > **fs** – File storage
> >
> > **map** – The parent map. If it is NULL, the function searches a top-level node.
> >
> > **name** – The node name
> >
> > **attributes** – Unused parameter

The function is a simple superposition of `GetFileNodeByName()` and `Read()`.

## ReadInt

Retrieves an integer value from a file node.

**C:** int **cvReadInt**(const CvFileNode* **node**, int **default_value**=0 )

> **Parameters**
>
> > **node** – File node
> >
> > **default_value** – The value that is returned if `node` is NULL

The function returns an integer that is represented by the file node. If the file node is NULL, the `default_value` is returned (thus, it is convenient to call the function right after `GetFileNode()` without checking for a NULL pointer). If the file node has type CV_NODE_INT, then node->data.i is returned. If the file node has type CV_NODE_REAL, then node->data.f is converted to an integer and returned. Otherwise the error is reported.

## ReadIntByName

Finds a file node and returns its value.

**C:** int **cvReadIntByName**(const CvFileStorage* **fs**, const CvFileNode* **map**, const char* **name**, int **default_value**=0 )

>   **Parameters**

>   >   **fs** – File storage

>   >   **map** – The parent map. If it is NULL, the function searches a top-level node.

>   >   **name** – The node name

>   >   **default_value** – The value that is returned if the file node is not found

The function is a simple superposition of `GetFileNodeByName()` and `ReadInt()`.

## ReadRawData

Reads multiple numbers.

**C:** void **cvReadRawData**(const CvFileStorage* **fs**, const CvFileNode* **src**, void* **dst**, const char* **dt**)

>   **Parameters**

>   >   **fs** – File storage

>   >   **src** – The file node (a sequence) to read numbers from

>   >   **dst** – Pointer to the destination array

>   >   **dt** – Specification of each array element. It has the same format as in `WriteRawData()`.

The function reads elements from a file node that represents a sequence of scalars.

## ReadRawDataSlice

Initializes file node sequence reader.

**C:** void **cvReadRawDataSlice**(const CvFileStorage* **fs**, CvSeqReader* **reader**, int **count**, void* **dst**, const char* **dt**)

>   **Parameters**

>   >   **fs** – File storage

>   >   **reader** – The sequence reader. Initialize it with `StartReadRawData()`.

>   >   **count** – The number of elements to read

>   >   **dst** – Pointer to the destination array

>   >   **dt** – Specification of each array element. It has the same format as in `WriteRawData()`.

The function reads one or more elements from the file node, representing a sequence, to a user-specified array. The total number of read sequence elements is a product of `total` and the number of components in each array element. For example, if dt=2if, the function will read `total*3` sequence elements. As with any sequence, some parts of the file node sequence can be skipped or read repeatedly by repositioning the reader using `SetSeqReaderPos()`.

## ReadReal

Retrieves a floating-point value from a file node.

**C:** double **cvReadReal**(const CvFileNode* **node**, double **default_value**=0. )

> **Parameters**
>
> > **node** – File node
> >
> > **default_value** – The value that is returned if `node` is NULL

The function returns a floating-point value that is represented by the file node. If the file node is NULL, the `default_value` is returned (thus, it is convenient to call the function right after `GetFileNode()` without checking for a NULL pointer). If the file node has type `CV_NODE_REAL` , then `node->data.f` is returned. If the file node has type `CV_NODE_INT` , then `node-:math:'>'data.f` is converted to floating-point and returned. Otherwise the result is not determined.

## ReadRealByName

Finds a file node and returns its value.

**C:** double **cvReadRealByName**(const CvFileStorage* **fs**, const CvFileNode* **map**, const char* **name**, double **default_value**=0. )

> **Parameters**
>
> > **fs** – File storage
> >
> > **map** – The parent map. If it is NULL, the function searches a top-level node.
> >
> > **name** – The node name
> >
> > **default_value** – The value that is returned if the file node is not found

The function is a simple superposition of `GetFileNodeByName()` and `ReadReal()` .

## ReadString

Retrieves a text string from a file node.

**C:** const char* **cvReadString**(const CvFileNode* **node**, const char* **default_value**=NULL )

> **Parameters**
>
> > **node** – File node
> >
> > **default_value** – The value that is returned if `node` is NULL

The function returns a text string that is represented by the file node. If the file node is NULL, the `default_value` is returned (thus, it is convenient to call the function right after `GetFileNode()` without checking for a NULL pointer). If the file node has type `CV_NODE_STR` , then `node-:math:'>'data.str.ptr` is returned. Otherwise the result is not determined.

## ReadStringByName

Finds a file node by its name and returns its value.

**C:** const char* **cvReadStringByName**(const CvFileStorage* **fs**, const CvFileNode* **map**, const char* **name**, const char* **default_value**=NULL )

> **Parameters**
>
> > **fs** – File storage
> >
> > **map** – The parent map. If it is NULL, the function searches a top-level node.
> >
> > **name** – The node name
> >
> > **default_value** – The value that is returned if the file node is not found

The function is a simple superposition of `GetFileNodeByName()` and `ReadString()` .

## RegisterType

Registers a new type.

**C:** void **cvRegisterType**(const CvTypeInfo* **info**)

> **Parameters**
>
> > **info** – Type info structure

The function registers a new type, which is described by `info` . The function creates a copy of the structure, so the user should delete it after calling the function.

## Release

Releases an object.

**C:** void **cvRelease**(void** **struct_ptr**)

> **Parameters**
>
> > **struct_ptr** – Double pointer to the object

The function finds the type of a given object and calls `release` with the double pointer.

## ReleaseFileStorage

Releases file storage.

**C:** void **cvReleaseFileStorage**(CvFileStorage** **fs**)

> **Parameters**
>
> > **fs** – Double pointer to the released file storage

The function closes the file associated with the storage and releases all the temporary structures. It must be called after all I/O operations with the storage are finished.

## Save

Saves an object to a file.

**C:** void **cvSave**(const char* **filename**, const void* **struct_ptr**, const char* **name**=NULL, const char* **comment**=NULL, CvAttrList **attributes**=cvAttrList() )

**Python:** cv.**Save**(filename, structPtr, name=None, comment=None) → None

> **Parameters**

> **filename** – File name
>
> **struct_ptr** – Object to save
>
> **name** – Optional object name. If it is NULL, the name will be formed from `filename` .
>
> **comment** – Optional comment to put in the beginning of the file
>
> **attributes** – Optional attributes passed to `Write()`

The function saves an object to a file. It provides a simple interface to `Write()` .

## StartNextStream

Starts the next stream.

**C:** void **cvStartNextStream**(CvFileStorage* **fs**)

> **Parameters**
>
> > **fs** – File storage

The function finishes the currently written stream and starts the next stream. In the case of XML the file with multiple streams looks like this:

```
<opencv_storage>
<!-- stream #1 data -->
</opencv_storage>
<opencv_storage>
<!-- stream #2 data -->
</opencv_storage>
...
```

The YAML file will look like this:

```
%YAML:1.0
# stream #1 data
...
---
# stream #2 data
```

This is useful for concatenating files or for resuming the writing process.

## StartReadRawData

Initializes the file node sequence reader.

**C:** void **cvStartReadRawData**(const CvFileStorage* **fs**, const CvFileNode* **src**, CvSeqReader* **reader**)

> **Parameters**
>
> > **fs** – File storage
> >
> > **src** – The file node (a sequence) to read numbers from
> >
> > **reader** – Pointer to the sequence reader

The function initializes the sequence reader to read data from a file node. The initialized reader can be then passed to `ReadRawDataSlice()`.

## StartWriteStruct

Starts writing a new structure.

**C:** void **cvStartWriteStruct**(CvFileStorage* **fs**, const char* **name**, int **struct_flags**, const char* **type_name**=NULL, CvAttrList **attributes**=cvAttrList() )

> **Parameters**
>
> > **fs** – File storage
> >
> > **name** – Name of the written structure. The structure can be accessed by this name when the storage is read.
> >
> > **struct_flags** – A combination one of the following values:
> >
> > - **CV_NODE_SEQ** the written structure is a sequence (see discussion of `CvFileStorage` ), that is, its elements do not have a name.
> > - **CV_NODE_MAP** the written structure is a map (see discussion of `CvFileStorage` ), that is, all its elements have names.
> >
> > One and only one of the two above flags must be specified
> >
> > - **CV_NODE_FLOW** the optional flag that makes sense only for YAML streams. It means that the structure is written as a flow (not as a block), which is more compact. It is recommended to use this flag for structures or arrays whose elements are all scalars.
> >
> > **type_name** – Optional parameter - the object type name. In case of XML it is written as a `type_id` attribute of the structure opening tag. In the case of YAML it is written after a colon following the structure name (see the example in `CvFileStorage` description). Mainly it is used with user objects. When the storage is read, the encoded type name is used to determine the object type (see `CvTypeInfo` and `FindType()` ).
> >
> > **attributes** – This parameter is not used in the current implementation

The function starts writing a compound structure (collection) that can be a sequence or a map. After all the structure fields, which can be scalars or structures, are written, `EndWriteStruct()` should be called. The function can be used to group some objects or to implement the `write` function for a some user object (see `CvTypeInfo`).

## TypeOf

Returns the type of an object.

**C:** CvTypeInfo* **cvTypeOf**(const void* **struct_ptr**)

> **Parameters**
>
> > **struct_ptr** – The object pointer

The function finds the type of a given object. It iterates through the list of registered types and calls the `is_instance` function/method for every type info structure with that object until one of them returns non-zero or until the whole list has been traversed. In the latter case, the function returns NULL.

## UnregisterType

Unregisters the type.

**C:** void **cvUnregisterType**(const char* **type_name**)

> **Parameters**

> > > **type_name** – Name of an unregistered type

The function unregisters a type with a specified name. If the name is unknown, it is possible to locate the type info by an instance of the type using `TypeOf()` or by iterating the type list, starting from `FirstType()`, and then calling `cvUnregisterType(info->typeName)`.

## Write

Writes an object to file storage.

**C:** void **cvWrite**(CvFileStorage* **fs**, const char* **name**, const void* **ptr**, CvAttrList **attributes**=cvAttrList() )

> > **Parameters**

> > > **fs** – File storage

> > > **name** – Name of the written object. Should be NULL if and only if the parent structure is a sequence.

> > > **ptr** – Pointer to the object

> > > **attributes** – The attributes of the object. They are specific for each particular type (see the discussion below).

The function writes an object to file storage. First, the appropriate type info is found using `TypeOf()`. Then, the `write` method associated with the type info is called.

Attributes are used to customize the writing procedure. The standard types support the following attributes (all the `dt` attributes have the same format as in `WriteRawData()`):

1. CvSeq

    • **header_dt** description of user fields of the sequence header that follow CvSeq, or CvChain (if the sequence is a Freeman chain) or CvContour (if the sequence is a contour or point sequence)

    • **dt** description of the sequence elements.

    • **recursive** if the attribute is present and is not equal to "0" or "false", the whole tree of sequences (contours) is stored.

2. CvGraph

    • **header_dt** description of user fields of the graph header that follows CvGraph;

    • **vertex_dt** description of user fields of graph vertices

    • **edge_dt** description of user fields of graph edges (note that the edge weight is always written, so there is no need to specify it explicitly)

Below is the code that creates the YAML file shown in the `CvFileStorage` description:

```c
#include "cxcore.h"

int main( int argc, char** argv )
{
    CvMat* mat = cvCreateMat( 3, 3, CV_32F );
    CvFileStorage* fs = cvOpenFileStorage( "example.yml", 0, CV_STORAGE_WRITE );

    cvSetIdentity( mat );
    cvWrite( fs, "A", mat, cvAttrList(0,0) );

    cvReleaseFileStorage( &fs );
    cvReleaseMat( &mat );
```

```
    return 0;
}
```

## WriteComment

Writes a comment.

**C:** void **cvWriteComment**(CvFileStorage* **fs**, const char* **comment**, int **eol_comment**)

> **Parameters**
>
>> **fs** – File storage
>>
>> **comment** – The written comment, single-line or multi-line
>>
>> **eol_comment** – If non-zero, the function tries to put the comment at the end of current line. If the flag is zero, if the comment is multi-line, or if it does not fit at the end of the current line, the comment starts a new line.

The function writes a comment into file storage. The comments are skipped when the storage is read.

## WriteFileNode

Writes a file node to another file storage.

**C:** void **cvWriteFileNode**(CvFileStorage* **fs**, const char* **new_node_name**, const CvFileNode* **node**, int **embed**)

> **Parameters**
>
>> **fs** – Destination file storage
>>
>> **new_node_name** – New name of the file node in the destination file storage. To keep the existing name, use `cvGetFileNodeName()`
>>
>> **node** – The written node
>>
>> **embed** – If the written node is a collection and this parameter is not zero, no extra level of hierarchy is created. Instead, all the elements of node are written into the currently written structure. Of course, map elements can only be embedded into another map, and sequence elements can only be embedded into another sequence.

The function writes a copy of a file node to file storage. Possible applications of the function are merging several file storages into one and conversion between XML and YAML formats.

## WriteInt

Writes an integer value.

**C:** void **cvWriteInt**(CvFileStorage* **fs**, const char* **name**, int **value**)

> **Parameters**
>
>> **fs** – File storage
>>
>> **name** – Name of the written value. Should be NULL if and only if the parent structure is a sequence.
>>
>> **value** – The written value

The function writes a single integer value (with or without a name) to the file storage.

## WriteRawData

Writes multiple numbers.

**C:** void **cvWriteRawData**(CvFileStorage* **fs**, const void* **src**, int **len**, const char* **dt**)

>       **Parameters**

>>              **fs** – File storage

>>              **src** – Pointer to the written array

>>              **len** – Number of the array elements to write

>>              **dt** – Specification of each array element that has the following format (`[count]{'u'|'c'|'w'|'s'|'i'|'f'|'d'})...` where the characters correspond to fundamental C types:

>>              – **u** 8-bit unsigned number

>>              – **c** 8-bit signed number

>>              – **w** 16-bit unsigned number

>>              – **s** 16-bit signed number

>>              – **i** 32-bit signed number

>>              – **f** single precision floating-point number

>>              – **d** double precision floating-point number

>>              – **r pointer, 32 lower bits of which are written as a signed integer. The type can be used to store structures w** example, `2if` means that each array element is a structure of 2 integers, followed by a single-precision floating-point number. The equivalent notations of the above specification are ' iif ', ' 2i1f ' and so forth. Other examples: u means that the array consists of bytes, and `2d` means the array consists of pairs of doubles.

The function writes an array, whose elements consist of single or multiple numbers. The function call can be replaced with a loop containing a few `WriteInt()` and `WriteReal()` calls, but a single call is more efficient. Note that because none of the elements have a name, they should be written to a sequence rather than a map.

## WriteReal

Writes a floating-point value.

**C:** void **cvWriteReal**(CvFileStorage* **fs**, const char* **name**, double **value**)

>       **Parameters**

>>              **fs** – File storage

>>              **name** – Name of the written value. Should be NULL if and only if the parent structure is a sequence.

>>              **value** – The written value

The function writes a single floating-point value (with or without a name) to file storage. Special values are encoded as follows: NaN (Not A Number) as .NaN, infinity as +.Inf or -.Inf.

The following example shows how to use the low-level writing functions to store custom structures, such as termination criteria, without registering a new type.

```
void write_termcriteria( CvFileStorage* fs, const char* struct_name,
                         CvTermCriteria* termcrit )
{
    cvStartWriteStruct( fs, struct_name, CV_NODE_MAP, NULL, cvAttrList(0,0));
    cvWriteComment( fs, "termination criteria", 1 ); // just a description
    if( termcrit->type & CV_TERMCRIT_ITER )
        cvWriteInteger( fs, "max_iterations", termcrit->max_iter );
    if( termcrit->type & CV_TERMCRIT_EPS )
        cvWriteReal( fs, "accuracy", termcrit->epsilon );
    cvEndWriteStruct( fs );
}
```

## WriteString

Writes a text string.

**C:** void **cvWriteString**(CvFileStorage* **fs**, const char* **name**, const char* **str**, int **quote**=0 )

> **Parameters**
>
> > **fs** – File storage
> >
> > **name** – Name of the written string . Should be NULL if and only if the parent structure is a sequence.
> >
> > **str** – The written text string
> >
> > **quote** – If non-zero, the written string is put in quotes, regardless of whether they are required. Otherwise, if the flag is zero, quotes are used only when they are required (e.g. when the string starts with a digit or contains spaces).

The function writes a text string to file storage.

# 2.8 Clustering

## kmeans

Finds centers of clusters and groups input samples around the clusters.

**C++:** double **kmeans**(InputArray **data**, int **K**, InputOutputArray **bestLabels**, TermCriteria **criteria**, int **attempts**, int **flags**, OutputArray **centers**=noArray() )

**Python:** cv2.**kmeans**(data, K, criteria, attempts, flags[, bestLabels[, centers ]]) → retval, bestLabels, centers

**C:** int **cvKMeans2**(const CvArr* **samples**, int **cluster_count**, CvArr* **labels**, CvTermCriteria **termcrit**, int **attempts**=1, CvRNG* **rng**=0, int **flags**=0, CvArr* **_centers**=0, double* **compactness**=0 )

**Python:** cv.**KMeans2**(samples, nclusters, labels, termcrit, attempts=1, flags=0, centers=None) → float

> **Parameters**
>
> > **samples** – Floating-point matrix of input samples, one row per sample.
> >
> > **cluster_count** – Number of clusters to split the set by.
> >
> > **labels** – Input/output integer array that stores the cluster indices for every sample.

> **criteria** – The algorithm termination criteria, that is, the maximum number of iterations and/or the desired accuracy. The accuracy is specified as `criteria.epsilon`. As soon as each of the cluster centers moves by less than `criteria.epsilon` on some iteration, the algorithm stops.
>
> **attempts** – Flag to specify the number of times the algorithm is executed using different initial labellings. The algorithm returns the labels that yield the best compactness (see the last function parameter).
>
> **rng** – CvRNG state initialized by RNG().
>
> **flags** – Flag that can take the following values:
>
> – **KMEANS_RANDOM_CENTERS** Select random initial centers in each attempt.
>
> – **KMEANS_PP_CENTERS** Use `kmeans++` center initialization by Arthur and Vassilvit-skii [Arthur2007].
>
> – **KMEANS_USE_INITIAL_LABELS** During the first (and possibly the only) attempt, use the user-supplied labels instead of computing them from the initial centers. For the second and further attempts, use the random or semi-random centers. Use one of `KMEANS_*_CENTERS` flag to specify the exact method.
>
> **centers** – Output matrix of the cluster centers, one row per each cluster center.
>
> **compactness** – The returned value that is described below.

The function `kmeans` implements a k-means algorithm that finds the centers of `cluster_count` clusters and groups the input samples around the clusters. As an output, $\texttt{labels}_i$ contains a 0-based cluster index for the sample stored in the $i^{\text{th}}$ row of the `samples` matrix.

The function returns the compactness measure that is computed as

$$\sum_i \| \texttt{samples}_i - \texttt{centers}_{\texttt{labels}_i} \|^2$$

after every attempt. The best (minimum) value is chosen and the corresponding labels and the compactness value are returned by the function. Basically, you can use only the core of the function, set the number of attempts to 1, initialize labels each time using a custom algorithm, pass them with the ( `flags` = `KMEANS_USE_INITIAL_LABELS` ) flag, and then choose the best (most-compact) clustering.

## partition

Splits an element set into equivalency classes.

**C++:** template<typename _Tp, class _EqPredicate> int **partition**(const vector<_Tp>& **vec**, vector<int>& **labels**, _EqPredicate **predicate**=_EqPredicate())

> **Parameters**
>
> > **vec** – Set of elements stored as a vector.
> >
> > **labels** – Output vector of labels. It contains as many elements as `vec`. Each label `labels[i]` is a 0-based cluster index of `vec[i]` .
> >
> > **predicate** – Equivalence predicate (pointer to a boolean function of two arguments or an instance of the class that has the method `bool operator()(const _Tp& a, const _Tp& b)` ). The predicate returns `true` when the elements are certainly in the same class, and returns `false` if they may or may not be in the same class.

The generic function `partition` implements an $O(N^2)$ algorithm for splitting a set of N elements into one or more equivalency classes, as described in http://en.wikipedia.org/wiki/Disjoint-set_data_structure . The function returns the number of equivalency classes.

## 2.9 Utility and System Functions and Macros

### alignPtr

Aligns a pointer to the specified number of bytes.

**C++:** template<typename _Tp> _Tp* **alignPtr**(_Tp* **ptr**, int **n**=sizeof(_Tp))

>       **Parameters**

>               **ptr** – Aligned pointer.

>               **n** – Alignment size that must be a power of two.

The function returns the aligned pointer of the same type as the input pointer:

$$(\_Tp*)(((size\_t)ptr + n-1) \& -n)$$

### alignSize

Aligns a buffer size to the specified number of bytes.

**C++:** size_t **alignSize**(size_t **sz**, int **n**)

>       **Parameters**

>               **sz** – Buffer size to align.

>               **n** – Alignment size that must be a power of two.

The function returns the minimum number that is greater or equal to `sz` and is divisible by `n` :

$$(sz + n-1) \& -n$$

### allocate

Allocates an array of elements.

**C++:** template<typename _Tp> _Tp* **allocate**(size_t **n**)

>       **Parameters**

>               **n** – Number of elements to allocate.

The generic function `allocate` allocates a buffer for the specified number of elements. For each element, the default constructor is called.

### deallocate

Deallocates an array of elements.

**C++:** template<typename _Tp> void **deallocate**(_Tp* **ptr**, size_t **n**)

>       **Parameters**

> **ptr** – Pointer to the deallocated buffer.
>
> **n** – Number of elements in the buffer.

The generic function `deallocate` deallocates the buffer allocated with `allocate()` . The number of elements must match the number passed to `allocate()` .

## fastAtan2

Calculates the angle of a 2D vector in degrees.

**C++:** float **fastAtan2**(float **y**, float **x**)

**Python:** cv2.**fastAtan2**(y, x) → retval

**C:** float **cvFastArctan**(float **y**, float **x**)

**Python:** cv.**FastArctan**(y, x) → float

> **Parameters**
>
> > **x** – x-coordinate of the vector.
> >
> > **y** – y-coordinate of the vector.

The function `fastAtan2` calculates the full-range angle of an input 2D vector. The angle is measured in degrees and varies from 0 to 360 degrees. The accuracy is about 0.3 degrees.

## cubeRoot

Computes the cube root of an argument.

**C++:** float **cubeRoot**(float **val**)

**Python:** cv2.**cubeRoot**(val) → retval

**C:** float **cvCbrt**(float **value**)

**Python:** cv.**Cbrt**(value) → float

> **Parameters**
>
> > **val** – A function argument.

The function `cubeRoot` computes $\sqrt[3]{\texttt{val}}$. Negative arguments are handled correctly. NaN and Inf are not handled. The accuracy approaches the maximum possible accuracy for single-precision data.

## Ceil

Rounds floating-point number to the nearest integer not smaller than the original.

**C:** int **cvCeil**(double **value**)

**Python:** cv.**Ceil**(value) → int

> **Parameters**
>
> > **value** – floating-point number. If the value is outside of `INT_MIN` ... `INT_MAX` range, the result is not defined.

The function computes an integer `i` such that:

$$i - 1 < \texttt{value} \le i$$

## Floor

Rounds floating-point number to the nearest integer not larger than the original.

**C:** int **cvFloor**(double **value**)

**Python:** cv.**Floor**(value) → int

> **Parameters**
>
> > **value** – floating-point number. If the value is outside of INT_MIN ... INT_MAX range, the result is not defined.

The function computes an integer i such that:

$$i \leq \text{value} < i + 1$$

## Round

Rounds floating-point number to the nearest integer

**C:** int **cvRound**(double **value**)

**Python:** cv.**Round**(value) → int

> **Parameters**
>
> > **value** – floating-point number. If the value is outside of INT_MIN ... INT_MAX range, the result is not defined.

## IsInf

Determines if the argument is Infinity.

**C:** int **cvIsInf**(double **value**)

**Python:** cv.**IsInf**(value) → int

> **Parameters**
>
> > **value** – The input floating-point value

The function returns 1 if the argument is a plus or minus infinity (as defined by IEEE754 standard) and 0 otherwise.

## IsNaN

Determines if the argument is Not A Number.

**C:** int **cvIsNaN**(double **value**)

**Python:** cv.**IsNaN**(value) → int

> **Parameters**
>
> > **value** – The input floating-point value

The function returns 1 if the argument is Not A Number (as defined by IEEE754 standard), 0 otherwise.

## CV_Assert

Checks a condition at runtime and throws exception if it fails

**C++:** `CV_Assert`(expr **None**)

The macros `CV_Assert` (and `CV_DbgAssert`) evaluate the specified expression. If it is 0, the macros raise an error (see `error()` ). The macro `CV_Assert` checks the condition in both Debug and Release configurations while `CV_DbgAssert` is only retained in the Debug configuration.

## error

Signals an error and raises an exception.

**C++:** void **error**(const Exception& **exc**)

**C:** void **cvError**(int **status**, const char* **func_name**, const char* **err_msg**, const char* **file_name**, int **line**)

   **Parameters**

     **exc** – Exception to throw.

     **status** – Error code. Normally, it is a negative value. The list of pre-defined error codes can be found in `cxerror.h` .

     **err_msg** – Text of the error message.

     **args** – `printf` -like formatted error message in parentheses.

The function and the helper macros `CV_Error` and `CV_Error_`:

```
#define CV_Error( code, msg ) error(...)
#define CV_Error_( code, args ) error(...)
```

call the error handler. Currently, the error handler prints the error code ( `exc.code` ), the context ( `exc.file`,‘‘exc.line‘‘ ), and the error message `exc.err` to the standard error stream `stderr` . In the Debug configuration, it then provokes memory access violation, so that the execution stack and all the parameters can be analyzed by the debugger. In the Release configuration, the exception `exc` is thrown.

The macro `CV_Error_` can be used to construct an error message on-fly to include some dynamic information, for example:

```
// note the extra parentheses around the formatted text message
CV_Error_(CV_StsOutOfRange,
    ("the matrix element (
    i, j, mtx.at<float>(i,j)))
```

## Exception

**class Exception** : **public** `std::exception`

Exception class passed to an error.

```
class  Exception
{
public:
    // various constructors and the copy operation
    Exception() { code = 0; line = 0; }
    Exception(int _code, const string& _err,
            const string& _func, const string& _file, int _line);
```

```
Exception(const Exception& exc);
Exception& operator = (const Exception& exc);

// the error code
int code;
// the error text message
string err;
// function name where the error happened
string func;
// the source file name where the error happened
string file;
// the source file line where the error happened
int line;
};
```

The class `Exception` encapsulates all or almost all necessary information about the error happened in the program. The exception is usually constructed and thrown implicitly via `CV_Error` and `CV_Error_` macros. See `error()` .

## fastMalloc

Allocates an aligned memory buffer.

**C++:** void* **fastMalloc**(size_t **bufSize**)

**C:** void* **cvAlloc**(size_t **size**)

>> **Parameters**

>>>> **size** – Allocated buffer size.

The function allocates the buffer of the specified size and returns it. When the buffer size is 16 bytes or more, the returned buffer is aligned to 16 bytes.

## fastFree

Deallocates a memory buffer.

**C++:** void **fastFree**(void* **ptr**)

**C:** void **cvFree**(void** **pptr**)

>> **Parameters**

>>>> **ptr** – Pointer to the allocated buffer.

>>>> **pptr** – Double pointer to the allocated buffer

The function deallocates the buffer allocated with `fastMalloc()` . If NULL pointer is passed, the function does nothing. C version of the function clears the pointer *pptr* to avoid problems with double memory deallocation.

## format

Returns a text string formatted using the `printf`-like expression.

**C++:** string **format**(const char* **fmt**, **...**)

>> **Parameters**

>>>> **fmt** – `printf` -compatible formatting specifiers.

The function acts like `sprintf` but forms and returns an STL string. It can be used to form an error message in the `Exception` constructor.

## checkHardwareSupport

Returns true if the specified feature is supported by the host hardware.

**C++:** bool **checkHardwareSupport**(int **feature**)

**C:** int **cvCheckHardwareSupport**(int **feature**)

**Python:** cv2.**checkHardwareSupport**(feature) → retval

>> **Parameters**

>>> **feature** – The feature of interest, one of:

>>>> – CV_CPU_MMX - MMX

>>>> – CV_CPU_SSE - SSE

>>>> – CV_CPU_SSE2 - SSE 2

>>>> – CV_CPU_SSE3 - SSE 3

>>>> – CV_CPU_SSSE3 - SSSE 3

>>>> – CV_CPU_SSE4_1 - SSE 4.1

>>>> – CV_CPU_SSE4_2 - SSE 4.2

>>>> – CV_CPU_POPCNT - POPCOUNT

>>>> – CV_CPU_AVX - AVX

The function returns true if the host hardware supports the specified feature. When user calls `setUseOptimized(false)`, the subsequent calls to `checkHardwareSupport()` will return false until `setUseOptimized(true)` is called. This way user can dynamically switch on and off the optimized code in OpenCV.

## getNumThreads

Returns the number of threads used by OpenCV.

**C++:** int **getNumThreads**()

The function returns the number of threads that is used by OpenCV.

**See Also:**

setNumThreads(), getThreadNum()

## getThreadNum

Returns the index of the currently executed thread.

**C++:** int **getThreadNum**()

The function returns a 0-based index of the currently executed thread. The function is only valid inside a parallel OpenMP region. When OpenCV is built without OpenMP support, the function always returns 0.

**See Also:**

setNumThreads(), getNumThreads() .

## getTickCount

Returns the number of ticks.

**C++:** int64 **getTickCount**()

**Python:** cv2.**getTickCount**() → retval

The function returns the number of ticks after the certain event (for example, when the machine was turned on). It can be used to initialize RNG() or to measure a function execution time by reading the tick count before and after the function call. See also the tick frequency.

## getTickFrequency

Returns the number of ticks per second.

**C++:** double **getTickFrequency**()

**Python:** cv2.**getTickFrequency**() → retval

The function returns the number of ticks per second. That is, the following code computes the execution time in seconds:

```
double t = (double)getTickCount();
// do something ...
t = ((double)getTickCount() - t)/getTickFrequency();
```

## getCPUTickCount

Returns the number of CPU ticks.

**C++:** int64 **getCPUTickCount**()

**Python:** cv2.**getCPUTickCount**() → retval

The function returns the current number of CPU ticks on some architectures (such as x86, x64, PowerPC). On other platforms the function is equivalent to getTickCount. It can also be used for very accurate time measurements, as well as for RNG initialization. Note that in case of multi-CPU systems a thread, from which getCPUTickCount is called, can be suspended and resumed at another CPU with its own counter. So, theoretically (and practically) the subsequent calls to the function do not necessary return the monotonously increasing values. Also, since a modern CPU varies the CPU frequency depending on the load, the number of CPU clocks spent in some code cannot be directly converted to time units. Therefore, getTickCount is generally a preferable solution for measuring execution time.

## saturate_cast

Template function for accurate conversion from one primitive type to another.

**C++:** template<...> _Tp **saturate_cast**(_Tp2 **v**)

> **Parameters**

>> **v** – Function parameter.

The functions saturate_cast resemble the standard C++ cast operations, such as static_cast<T>() and others. They perform an efficient and accurate conversion from one primitive type to another (see the introduction chapter). saturate in the name means that when the input value v is out of the range of the target type, the result is not formed just by taking low bits of the input, but instead the value is clipped. For example:

```
uchar a = saturate_cast<uchar>(-100); // a = 0 (UCHAR_MIN)
short b = saturate_cast<short>(33333.33333); // b = 32767 (SHRT_MAX)
```

Such clipping is done when the target type is `unsigned char`, `signed char`, `unsigned short` or `signed short`. For 32-bit integers, no clipping is done.

When the parameter is a floating-point value and the target type is an integer (8-, 16- or 32-bit), the floating-point value is first rounded to the nearest integer and then clipped if needed (when the target type is 8- or 16-bit).

This operation is used in the simplest or most complex image processing functions in OpenCV.

**See Also:**

add(), subtract(), multiply(), divide(), Mat::convertTo()

## setNumThreads

Sets the number of threads used by OpenCV.

**C++:** void **setNumThreads**(int **nthreads**)

>    **Parameters**

>        **nthreads** – Number of threads used by OpenCV.

The function sets the number of threads used by OpenCV in parallel OpenMP regions. If `nthreads=0`, the function uses the default number of threads that is usually equal to the number of the processing cores.

**See Also:**

getNumThreads(), getThreadNum()

## setUseOptimized

Enables or disables the optimized code.

**C++:** int **cvUseOptimized**(int **on_off**)

**Python:** cv2.**setUseOptimized**(onoff) → None

**C:** int **cvUseOptimized**(int **on_off**)

>    **Parameters**

>        **on_off** – The boolean flag specifying whether the optimized code should be used (on_off=true) or not (on_off=false).

The function can be used to dynamically turn on and off optimized code (code that uses SSE2, AVX, and other instructions on the platforms that support it). It sets a global flag that is further checked by OpenCV functions. Since the flag is not checked in the inner OpenCV loops, it is only safe to call the function on the very top level in your application where you can be sure that no other OpenCV function is currently executed.

By default, the optimized code is enabled unless you disable it in CMake. The current status can be retrieved using `useOptimized`.

## useOptimized

Returns the status of optimized code usage.

**C++:** bool **useOptimized**()

**Python:** cv2.**useOptimized**() → retval

The function returns `true` if the optimized code is enabled. Otherwise, it returns `false`.

# IMGPROC. IMAGE PROCESSING

## 3.1 Image Filtering

Functions and classes described in this section are used to perform various linear or non-linear filtering operations on 2D images (represented as Mat()'s). It means that for each pixel location $(x, y)$ in the source image (normally, rectangular), its neighborhood is considered and used to compute the response. In case of a linear filter, it is a weighted sum of pixel values. In case of morphological operations, it is the minimum or maximum values, and so on. The computed response is stored in the destination image at the same location $(x, y)$ . It means that the output image will be of the same size as the input image. Normally, the functions support multi-channel arrays, in which case every channel is processed independently. Therefore, the output image will also have the same number of channels as the input one.

Another common feature of the functions and classes described in this section is that, unlike simple arithmetic functions, they need to extrapolate values of some non-existing pixels. For example, if you want to smooth an image using a Gaussian $3 \times 3$ filter, then, when processing the left-most pixels in each row, you need pixels to the left of them, that is, outside of the image. You can let these pixels be the same as the left-most image pixels ("replicated border" extrapolation method), or assume that all the non-existing pixels are zeros ("constant border" extrapolation method), and so on. OpenCV enables you to specify the extrapolation method. For details, see the function borderInterpolate() and discussion of the borderType parameter in the section and various functions below.

```
/*
 Various border types, image boundaries are denoted with '|'

 * BORDER_REPLICATE:     aaaaaa|abcdefgh|hhhhhhh
 * BORDER_REFLECT:       fedcba|abcdefgh|hgfedcb
 * BORDER_REFLECT_101:   gfedcb|abcdefgh|gfedcba
 * BORDER_WRAP:          cdefgh|abcdefgh|abcdefg
 * BORDER_CONSTANT:      iiiiii|abcdefgh|iiiiiii  with some specified 'i'
*/
```

## BaseColumnFilter

**class BaseColumnFilter**

Base class for filters with single-column kernels.

```
class BaseColumnFilter
{
public:
    virtual ~BaseColumnFilter();

    // To be overriden by the user.
```

```
    //
    // runs a filtering operation on the set of rows,
    // "dstcount + ksize - 1" rows on input,
    // "dstcount" rows on output,
    // each input and output row has "width" elements
    // the filtered rows are written into "dst" buffer.
    virtual void operator()(const uchar** src, uchar* dst, int dststep,
                            int dstcount, int width) = 0;
    // resets the filter state (may be needed for IIR filters)
    virtual void reset();

    int ksize; // the aperture size
    int anchor; // position of the anchor point,
                // normally not used during the processing
};
```

The class `BaseColumnFilter` is a base class for filtering data using single-column kernels. Filtering does not have to be a linear operation. In general, it could be written as follows:

$$dst(x, y) = F(src[y](x),\ src[y+1](x),\ ...,\ src[y+ksize-1](x))$$

where F is a filtering function but, as it is represented as a class, it can produce any side effects, memorize previously processed data, and so on. The class only defines an interface and is not used directly. Instead, there are several functions in OpenCV (and you can add more) that return pointers to the derived classes that implement specific filtering operations. Those pointers are then passed to the `FilterEngine` constructor. While the filtering operation interface uses the `uchar` type, a particular implementation is not limited to 8-bit data.

**See Also:**

BaseRowFilter, BaseFilter, FilterEngine, getColumnSumFilter(), getLinearColumnFilter(), getMorphologyColumnFilter()

## BaseFilter

**class BaseFilter**

Base class for 2D image filters.

```
class BaseFilter
{
public:
    virtual ~BaseFilter();

    // To be overriden by the user.
    //
    // runs a filtering operation on the set of rows,
    // "dstcount + ksize.height - 1" rows on input,
    // "dstcount" rows on output,
    // each input row has "(width + ksize.width-1)*cn" elements
    // each output row has "width*cn" elements.
    // the filtered rows are written into "dst" buffer.
    virtual void operator()(const uchar** src, uchar* dst, int dststep,
                            int dstcount, int width, int cn) = 0;
    // resets the filter state (may be needed for IIR filters)
    virtual void reset();
    Size ksize;
    Point anchor;
};
```

The class `BaseFilter` is a base class for filtering data using 2D kernels. Filtering does not have to be a linear operation. In general, it could be written as follows:

$$\texttt{dst}(x, y) = F(\texttt{src}[y](x), \texttt{src}[y](x+1), ..., \texttt{src}[y](x+\texttt{ksize.width}-1),$$
$$\texttt{src}[y+1](x), \texttt{src}[y+1](x+1), ..., \texttt{src}[y+1](x+\texttt{ksize.width}-1),$$
$$.....................................................................................................$$
$$\texttt{src}[y+\texttt{ksize.height-1}](x),$$
$$\texttt{src}[y+\texttt{ksize.height-1}](x+1),$$
$$...\texttt{src}[y+\texttt{ksize.height-1}](x+\texttt{ksize.width}-1))$$

where $F$ is a filtering function. The class only defines an interface and is not used directly. Instead, there are several functions in OpenCV (and you can add more) that return pointers to the derived classes that implement specific filtering operations. Those pointers are then passed to the `FilterEngine` constructor. While the filtering operation interface uses the `uchar` type, a particular implementation is not limited to 8-bit data.

**See Also:**

BaseColumnFilter, BaseRowFilter, FilterEngine, getLinearFilter(), getMorphologyFilter()

## BaseRowFilter

**class BaseRowFilter**

Base class for filters with single-row kernels.

```
class BaseRowFilter
{
public:
    virtual ~BaseRowFilter();

    // To be overriden by the user.
    //
    // runs filtering operation on the single input row
    // of "width" element, each element is has "cn" channels.
    // the filtered row is written into "dst" buffer.
    virtual void operator()(const uchar* src, uchar* dst,
                            int width, int cn) = 0;
    int ksize, anchor;
};
```

The class `BaseRowFilter` is a base class for filtering data using single-row kernels. Filtering does not have to be a linear operation. In general, it could be written as follows:

$$\texttt{dst}(x, y) = F(\texttt{src}[y](x), \texttt{src}[y](x+1), ..., \texttt{src}[y](x+\texttt{ksize.width}-1))$$

where $F$ is a filtering function. The class only defines an interface and is not used directly. Instead, there are several functions in OpenCV (and you can add more) that return pointers to the derived classes that implement specific filtering operations. Those pointers are then passed to the `FilterEngine` constructor. While the filtering operation interface uses the `uchar` type, a particular implementation is not limited to 8-bit data.

**See Also:**

BaseColumnFilter, BaseFilter, FilterEngine, getLinearRowFilter(), getMorphologyRowFilter(), getRowSumFilter()

## FilterEngine

**class FilterEngine**

Generic image filtering class.

```cpp
class FilterEngine
{
public:
    // empty constructor
    FilterEngine();
    // builds a 2D non-separable filter (!_filter2D.empty()) or
    // a separable filter (!_rowFilter.empty() && !_columnFilter.empty())
    // the input data type will be "srcType", the output data type will be "dstType",
    // the intermediate data type is "bufType".
    // _rowBorderType and _columnBorderType determine how the image
    // will be extrapolated beyond the image boundaries.
    // _borderValue is only used when _rowBorderType and/or _columnBorderType
    // == BORDER_CONSTANT
    FilterEngine(const Ptr<BaseFilter>& _filter2D,
                 const Ptr<BaseRowFilter>& _rowFilter,
                 const Ptr<BaseColumnFilter>& _columnFilter,
                 int srcType, int dstType, int bufType,
                 int _rowBorderType=BORDER_REPLICATE,
                 int _columnBorderType=-1, // use _rowBorderType by default
                 const Scalar& _borderValue=Scalar());
    virtual ~FilterEngine();
    // separate function for the engine initialization
    void init(const Ptr<BaseFilter>& _filter2D,
            const Ptr<BaseRowFilter>& _rowFilter,
            const Ptr<BaseColumnFilter>& _columnFilter,
            int srcType, int dstType, int bufType,
            int _rowBorderType=BORDER_REPLICATE, int _columnBorderType=-1,
            const Scalar& _borderValue=Scalar());
    // starts filtering of the ROI in an image of size "wholeSize".
    // returns the starting y-position in the source image.
    virtual int start(Size wholeSize, Rect roi, int maxBufRows=-1);
    // alternative form of start that takes the image
    // itself instead of "wholeSize". Set isolated to true to pretend that
    // there are no real pixels outside of the ROI
    // (so that the pixels are extrapolated using the specified border modes)
    virtual int start(const Mat& src, const Rect& srcRoi=Rect(0,0,-1,-1),
                      bool isolated=false, int maxBufRows=-1);
    // processes the next portion of the source image,
    // "srcCount" rows starting from "src" and
    // stores the results in "dst".
    // returns the number of produced rows
    virtual int proceed(const uchar* src, int srcStep, int srcCount,
                        uchar* dst, int dstStep);
    // higher-level function that processes the whole
    // ROI or the whole image with a single call
    virtual void apply( const Mat& src, Mat& dst,
                        const Rect& srcRoi=Rect(0,0,-1,-1),
                        Point dstOfs=Point(0,0),
                        bool isolated=false);
    bool isSeparable() const { return filter2D.empty(); }
    // how many rows from the input image are not yet processed
    int remainingInputRows() const;
    // how many output rows are not yet produced
    int remainingOutputRows() const;
    ...
    // the starting and the ending rows in the source image
    int startY, endY;
```

```
    // pointers to the filters
    Ptr<BaseFilter> filter2D;
    Ptr<BaseRowFilter> rowFilter;
    Ptr<BaseColumnFilter> columnFilter;
};
```

The class `FilterEngine` can be used to apply an arbitrary filtering operation to an image. It contains all the necessary intermediate buffers, computes extrapolated values of the "virtual" pixels outside of the image, and so on. Pointers to the initialized `FilterEngine` instances are returned by various `create*Filter` functions (see below) and they are used inside high-level functions such as `filter2D()`, `erode()`, `dilate()`, and others. Thus, the class plays a key role in many of OpenCV filtering functions.

This class makes it easier to combine filtering operations with other operations, such as color space conversions, thresholding, arithmetic operations, and others. By combining several operations together you can get much better performance because your data will stay in cache. For example, see below the implementation of the Laplace operator for floating-point images, which is a simplified implementation of `Laplacian()` :

```
void laplace_f(const Mat& src, Mat& dst)
{
    CV_Assert( src.type() == CV_32F );
    dst.create(src.size(), src.type());

    // get the derivative and smooth kernels for d2I/dx2.
    // for d2I/dy2 consider using the same kernels, just swapped
    Mat kd, ks;
    getSobelKernels( kd, ks, 2, 0, ksize, false, ktype );

    // process 10 source rows at once
    int DELTA = std::min(10, src.rows);
    Ptr<FilterEngine> Fxx = createSeparableLinearFilter(src.type(),
        dst.type(), kd, ks, Point(-1,-1), 0, borderType, borderType, Scalar() );
    Ptr<FilterEngine> Fyy = createSeparableLinearFilter(src.type(),
        dst.type(), ks, kd, Point(-1,-1), 0, borderType, borderType, Scalar() );

    int y = Fxx->start(src), dsty = 0, dy = 0;
    Fyy->start(src);
    const uchar* sptr = src.data + y*src.step;

    // allocate the buffers for the spatial image derivatives;
    // the buffers need to have more than DELTA rows, because at the
    // last iteration the output may take max(kd.rows-1,ks.rows-1)
    // rows more than the input.
    Mat Ixx( DELTA + kd.rows - 1, src.cols, dst.type() );
    Mat Iyy( DELTA + kd.rows - 1, src.cols, dst.type() );

    // inside the loop always pass DELTA rows to the filter
    // (note that the "proceed" method takes care of possibe overflow, since
    // it was given the actual image height in the "start" method)
    // on output you can get:
    //  * < DELTA rows (initial buffer accumulation stage)
    //  * = DELTA rows (settled state in the middle)
    //  * > DELTA rows (when the input image is over, generate
    //                 "virtual" rows using the border mode and filter them)
    // this variable number of output rows is dy.
    // dsty is the current output row.
    // sptr is the pointer to the first input row in the portion to process
    for( ; dsty < dst.rows; sptr += DELTA*src.step, dsty += dy )
    {
```

```
            Fxx->proceed( sptr, (int)src.step, DELTA, Ixx.data, (int)Ixx.step );
            dy = Fyy->proceed( sptr, (int)src.step, DELTA, d2y.data, (int)Iyy.step );
            if( dy > 0 )
            {
                Mat dstripe = dst.rowRange(dsty, dsty + dy);
                add(Ixx.rowRange(0, dy), Iyy.rowRange(0, dy), dstripe);
            }
        }
    }
}
```

If you do not need that much control of the filtering process, you can simply use the `FilterEngine::apply` method. The method is implemented as follows:

```
void FilterEngine::apply(const Mat& src, Mat& dst,
    const Rect& srcRoi, Point dstOfs, bool isolated)
{
    // check matrix types
    CV_Assert( src.type() == srcType && dst.type() == dstType );

    // handle the "whole image" case
    Rect _srcRoi = srcRoi;
    if( _srcRoi == Rect(0,0,-1,-1) )
        _srcRoi = Rect(0,0,src.cols,src.rows);

    // check if the destination ROI is inside dst.
    // and FilterEngine::start will check if the source ROI is inside src.
    CV_Assert( dstOfs.x >= 0 && dstOfs.y >= 0 &&
        dstOfs.x + _srcRoi.width <= dst.cols &&
        dstOfs.y + _srcRoi.height <= dst.rows );

    // start filtering
    int y = start(src, _srcRoi, isolated);

    // process the whole ROI. Note that "endY - startY" is the total number
    // of the source rows to process
    // (including the possible rows outside of srcRoi but inside the source image)
    proceed( src.data + y*src.step,
            (int)src.step, endY - startY,
            dst.data + dstOfs.y*dst.step +
            dstOfs.x*dst.elemSize(), (int)dst.step );
}
```

Unlike the earlier versions of OpenCV, now the filtering operations fully support the notion of image ROI, that is, pixels outside of the ROI but inside the image can be used in the filtering operations. For example, you can take a ROI of a single pixel and filter it. This will be a filter response at that particular pixel. However, it is possible to emulate the old behavior by passing `isolated=false` to `FilterEngine::start` or `FilterEngine::apply`. You can pass the ROI explicitly to `FilterEngine::apply` or construct new matrix headers:

```
// compute dI/dx derivative at src(x,y)

// method 1:
// form a matrix header for a single value
float val1 = 0;
Mat dst1(1,1,CV_32F,&val1);

Ptr<FilterEngine> Fx = createDerivFilter(CV_32F, CV_32F,
                    1, 0, 3, BORDER_REFLECT_101);
Fx->apply(src, Rect(x,y,1,1), Point(), dst1);
```

```
// method 2:
// form a matrix header for a single value
float val2 = 0;
Mat dst2(1,1,CV_32F,&val2);

Mat pix_roi(src, Rect(x,y,1,1));
Sobel(pix_roi, dst2, dst2.type(), 1, 0, 3, 1, 0, BORDER_REFLECT_101);

printf("method1 =
```

Explore the data types. As it was mentioned in the `BaseFilter` description, the specific filters can process data of any type, despite that `Base*Filter::operator()` only takes `uchar` pointers and no information about the actual types. To make it all work, the following rules are used:

- In case of separable filtering, `FilterEngine::rowFilter` is applied first. It transforms the input image data (of type `srcType` ) to the intermediate results stored in the internal buffers (of type `bufType` ). Then, these intermediate results are processed as *single-channel data* with `FilterEngine::columnFilter` and stored in the output image (of type `dstType` ). Thus, the input type for `rowFilter` is `srcType` and the output type is `bufType` . The input type for `columnFilter` is `CV_MAT_DEPTH(bufType)` and the output type is `CV_MAT_DEPTH(dstType)` .

- In case of non-separable filtering, `bufType` must be the same as `srcType` . The source data is copied to the temporary buffer, if needed, and then just passed to `FilterEngine::filter2D` . That is, the input type for `filter2D` is `srcType` (= `bufType` ) and the output type is `dstType` .

**See Also:**

BaseColumnFilter, BaseFilter, BaseRowFilter, createBoxFilter(), createDerivFilter(), createGaussianFilter(), createLinearFilter(), createMorphologyFilter(), createSeparableLinearFilter()

## bilateralFilter

Applies the bilateral filter to an image.

**C++:** void **bilateralFilter**(InputArray **src**, OutputArray **dst**, int **d**, double **sigmaColor**, double **sigmaSpace**, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**bilateralFilter**(src, d, sigmaColor, sigmaSpace[, dst[, borderType ] ]) → dst

> **Parameters**

> > **src** – Source 8-bit or floating-point, 1-channel or 3-channel image.

> > **dst** – Destination image of the same size and type as `src` .

> > **d** – Diameter of each pixel neighborhood that is used during filtering. If it is non-positive, it is computed from `sigmaSpace` .

> > **sigmaColor** – Filter sigma in the color space. A larger value of the parameter means that farther colors within the pixel neighborhood (see `sigmaSpace` ) will be mixed together, resulting in larger areas of semi-equal color.

> > **sigmaSpace** – Filter sigma in the coordinate space. A larger value of the parameter means that farther pixels will influence each other as long as their colors are close enough (see `sigmaColor` ). When d>0 , it specifies the neighborhood size regardless of `sigmaSpace` . Otherwise, d is proportional to `sigmaSpace` .

The function applies bilateral filtering to the input image, as described in http://www.dai.ed.ac.uk/CVonline/LOCAL_COPIES/MANDUCHI1/Bilateral_Filtering.html bilateralFilter

can reduce unwanted noise very well while keeping edges fairly sharp. However, it is very slow compared to most filters.

*Sigma values*: For simplicity, you can set the 2 sigma values to be the same. If they are small (< 10), the filter will not have much effect, whereas if they are large (> 150), they will have a very strong effect, making the image look "cartoonish".

*Filter size*: Large filters (d > 5) are very slow, so it is recommended to use d=5 for real-time applications, and perhaps d=9 for offline applications that need heavy noise filtering.

This filter does not work inplace.

## blur

Blurs an image using the normalized box filter.

**C++:** void **blur**(InputArray **src**, OutputArray **dst**, Size **ksize**, Point **anchor**=Point(-1,-1), int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**blur**(src, ksize[, dst[, anchor[, borderType ]]]) → dst

> **Parameters**
>
> > **src** – input image; it can have any number of channels, which are processed independently, but the depth should be `CV_8U`, `CV_16U`, `CV_16S`, `CV_32F` or `CV_64F`.
> >
> > **dst** – output image of the same size and type as `src`.
> >
> > **ksize** – blurring kernel size.
> >
> > **anchor** – anchor point; default value `Point(-1,-1)` means that the anchor is at the kernel center.
> >
> > **borderType** – border mode used to extrapolate pixels outside of the image.

The function smoothes an image using the kernel:

$$K = \frac{1}{\texttt{ksize.width*ksize.height}} \begin{bmatrix} 1 & 1 & 1 & \cdots & 1 & 1 \\ 1 & 1 & 1 & \cdots & 1 & 1 \\ & & \cdots\cdots\cdots & & \\ 1 & 1 & 1 & \cdots & 1 & 1 \end{bmatrix}$$

The call `blur(src, dst, ksize, anchor, borderType)` is equivalent to `boxFilter(src, dst, src.type(), anchor, true, borderType)` .

**See Also:**

`boxFilter()`, `bilateralFilter()`, `GaussianBlur()`, `medianBlur()`

## borderInterpolate

Computes the source location of an extrapolated pixel.

**C++:** int **borderInterpolate**(int **p**, int **len**, int **borderType**)

**Python:** cv2.**borderInterpolate**(p, len, borderType) → retval

> **Parameters**
>
> > **p** – 0-based coordinate of the extrapolated pixel along one of the axes, likely <0 or >= `len` .
> >
> > **len** – Length of the array along the corresponding axis.

> **borderType** – Border type, one of the `BORDER_*` , except for `BORDER_TRANSPARENT` and `BORDER_ISOLATED` . When `borderType==BORDER_CONSTANT` , the function always returns -1, regardless of `p` and `len` .

The function computes and returns the coordinate of a donor pixel corresponding to the specified extrapolated pixel when using the specified extrapolation border mode. For example, if you use `BORDER_WRAP` mode in the horizontal direction, `BORDER_REFLECT_101` in the vertical direction and want to compute value of the "virtual" pixel `Point(-5, 100)` in a floating-point image `img` , it looks like:

```
float val = img.at<float>(borderInterpolate(100, img.rows, BORDER_REFLECT_101),
                          borderInterpolate(-5, img.cols, BORDER_WRAP));
```

Normally, the function is not called directly. It is used inside `FilterEngine` and `copyMakeBorder()` to compute tables for quick extrapolation.

See Also:

`FilterEngine`, `copyMakeBorder()`

## boxFilter

Blurs an image using the box filter.

**C++:** void **boxFilter**(InputArray **src**, OutputArray **dst**, int **ddepth**, Size **ksize**, Point **anchor**=Point(-1,-1), bool **normalize**=true, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**boxFilter**(src, ddepth, ksize[, dst[, anchor[, normalize[, borderType ]]]]) → dst

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image of the same size and type as `src`.
> >
> > **ksize** – blurring kernel size.
> >
> > **anchor** – anchor point; default value `Point(-1,-1)` means that the anchor is at the kernel center.
> >
> > **normalize** – flag, specifying whether the kernel is normalized by its area or not.
> >
> > **borderType** – border mode used to extrapolate pixels outside of the image.

The function smoothes an image using the kernel:

$$
K = \alpha \begin{bmatrix} 1 & 1 & 1 & \cdots & 1 & 1 \\ 1 & 1 & 1 & \cdots & 1 & 1 \\ & & \cdots\cdots\cdots & & \\ 1 & 1 & 1 & \cdots & 1 & 1 \end{bmatrix}
$$

where

$$
\alpha = \begin{cases} \frac{1}{\texttt{ksize.width*ksize.height}} & \text{when } \texttt{normalize=true} \\ 1 & \text{otherwise} \end{cases}
$$

Unnormalized box filter is useful for computing various integral characteristics over each pixel neighborhood, such as covariance matrices of image derivatives (used in dense optical flow algorithms, and so on). If you need to compute pixel sums over variable-size windows, use `integral()` .

See Also:

`blur()`, `bilateralFilter()`, `GaussianBlur()`, `medianBlur()`, `integral()`

## buildPyramid

Constructs the Gaussian pyramid for an image.

**C++:** void **buildPyramid**(InputArray **src**, OutputArrayOfArrays **dst**, int **maxlevel**, int **border-Type**=BORDER_DEFAULT )

> **Parameters**
>
>> **src** – Source image. Check `pyrDown()` for the list of supported types.
>>
>> **dst** – Destination vector of `maxlevel+1` images of the same type as `src` . `dst[0]` will be the same as `src` . `dst[1]` is the next pyramid layer, a smoothed and down-sized `src` , and so on.
>>
>> **maxlevel** – 0-based index of the last (the smallest) pyramid layer. It must be non-negative.

The function constructs a vector of images and builds the Gaussian pyramid by recursively applying `pyrDown()` to the previously built pyramid layers, starting from `dst[0]==src` .

## copyMakeBorder

Forms a border around an image.

**C++:** void **copyMakeBorder**(InputArray **src**, OutputArray **dst**, int **top**, int **bottom**, int **left**, int **right**, int **borderType**, const Scalar& **value**=Scalar() )

**Python:** cv2.**copyMakeBorder**(src, top, bottom, left, right, borderType[, dst[, value ]]) → dst

**C:** void **cvCopyMakeBorder**(const CvArr* **src**, CvArr* **dst**, CvPoint **offset**, int **bordertype**, CvScalar **value**=cvScalarAll(0) )

**Python:** cv.**CopyMakeBorder**(src, dst, offset, bordertype, value=(0, 0, 0, 0)) → None

> **Parameters**
>
>> **src** – Source image.
>>
>> **dst** – Destination image of the same type as `src` and the size `Size(src.cols+left+right, src.rows+top+bottom)` .
>>
>> **top** –
>>
>> **bottom** –
>>
>> **left** –
>>
>> **right** – Parameter specifying how many pixels in each direction from the source image rectangle to extrapolate. For example, `top=1, bottom=1, left=1, right=1` mean that 1 pixel-wide border needs to be built.
>>
>> **borderType** – Border type. See `borderInterpolate()` for details.
>>
>> **value** – Border value if `borderType==BORDER_CONSTANT` .

The function copies the source image into the middle of the destination image. The areas to the left, to the right, above and below the copied source image will be filled with extrapolated pixels. This is not what `FilterEngine` or filtering functions based on it do (they extrapolate pixels on-fly), but what other more complex functions, including your own, may do to simplify image boundary handling.

The function supports the mode when `src` is already in the middle of `dst` . In this case, the function does not copy `src` itself but simply constructs the border, for example:

```
// let border be the same in all directions
int border=2;
// constructs a larger image to fit both the image and the border
Mat gray_buf(rgb.rows + border*2, rgb.cols + border*2, rgb.depth());
// select the middle part of it w/o copying data
Mat gray(gray_canvas, Rect(border, border, rgb.cols, rgb.rows));
// convert image from RGB to grayscale
cvtColor(rgb, gray, CV_RGB2GRAY);
// form a border in-place
copyMakeBorder(gray, gray_buf, border, border,
               border, border, BORDER_REPLICATE);
// now do some custom filtering ...
...
```

**Note:** When the source image is a part (ROI) of a bigger image, the function will try to use the pixels outside of the ROI to form a border. To disable this feature and always do extrapolation, as if src was not a ROI, use borderType | BORDER_ISOLATED.

**See Also:**

borderInterpolate()

## createBoxFilter

Returns a box filter engine.

**C++:** Ptr<FilterEngine> **createBoxFilter**(int **srcType**, int **dstType**, Size **ksize**, Point **anchor**=Point(-1,-1), bool **normalize**=true, int **borderType**=BORDER_DEFAULT)

**C++:** Ptr<BaseRowFilter> **getRowSumFilter**(int **srcType**, int **sumType**, int **ksize**, int **anchor**=-1)

**C++:** Ptr<BaseColumnFilter> **getColumnSumFilter**(int **sumType**, int **dstType**, int **ksize**, int **anchor**=-1, double **scale**=1)

> Parameters
>> **srcType** – Source image type.
>>
>> **sumType** – Intermediate horizontal sum type that must have as many channels as srcType .
>>
>> **dstType** – Destination image type that must have as many channels as srcType .
>>
>> **ksize** – Aperture size.
>>
>> **anchor** – Anchor position with the kernel. Negative values mean that the anchor is at the kernel center.
>>
>> **normalize** – Flag specifying whether the sums are normalized or not. See boxFilter() for details.
>>
>> **scale** – Another way to specify normalization in lower-level getColumnSumFilter .
>>
>> **borderType** – Border type to use. See borderInterpolate() .

The function is a convenience function that retrieves the horizontal sum primitive filter with getRowSumFilter() , vertical sum filter with getColumnSumFilter() , constructs new FilterEngine , and passes both of the primitive filters there. The constructed filter engine can be used for image filtering with normalized or unnormalized box filter.

The function itself is used by blur() and boxFilter() .

**See Also:**

`FilterEngine`, `blur()`, `boxFilter()`

## createDerivFilter

Returns an engine for computing image derivatives.

**C++:** Ptr<FilterEngine> **createDerivFilter**(int **srcType**, int **dstType**, int **dx**, int **dy**, int **ksize**, int **border-Type**=BORDER_DEFAULT )

>> **Parameters**

>>>> **srcType** – Source image type.

>>>> **dstType** – Destination image type that must have as many channels as `srcType` .

>>>> **dx** – Derivative order in respect of x.

>>>> **dy** – Derivative order in respect of y.

>>>> **ksize** – Aperture size See `getDerivKernels()` .

>>>> **borderType** – Border type to use. See `borderInterpolate()` .

The function `createDerivFilter()` is a small convenience function that retrieves linear filter coefficients for computing image derivatives using `getDerivKernels()` and then creates a separable linear filter with `createSeparableLinearFilter()` . The function is used by `Sobel()` and `Scharr()` .

**See Also:**

`createSeparableLinearFilter()`, `getDerivKernels()`, `Scharr()`, `Sobel()`

## createGaussianFilter

Returns an engine for smoothing images with the Gaussian filter.

**C++:** Ptr<FilterEngine> **createGaussianFilter**(int **type**, Size **ksize**, double **sigma1**, double **sigma2**=0, int **borderType**=BORDER_DEFAULT )

>> **Parameters**

>>>> **type** – Source and destination image type.

>>>> **ksize** – Aperture size. See `getGaussianKernel()` .

>>>> **sigma1** – Gaussian sigma in the horizontal direction. See `getGaussianKernel()` .

>>>> **sigma2** – Gaussian sigma in the vertical direction. If 0, then `sigma2` ← `sigma1` .

>>>> **borderType** – Border type to use. See `borderInterpolate()` .

The function `createGaussianFilter()` computes Gaussian kernel coefficients and then returns a separable linear filter for that kernel. The function is used by `GaussianBlur()` . Note that while the function takes just one data type, both for input and output, you can pass this limitation by calling `getGaussianKernel()` and then `createSeparableLinearFilter()` directly.

**See Also:**

`createSeparableLinearFilter()`, `getGaussianKernel()`, `GaussianBlur()`

# createLinearFilter

Creates a non-separable linear filter engine.

**C++:** Ptr<FilterEngine> **createLinearFilter**(int **srcType**, int **dstType**, InputArray **kernel**, Point **_anchor**=Point(-1,-1), double **delta**=0, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1, const Scalar& **borderValue**=Scalar() )

**C++:** Ptr<BaseFilter> **getLinearFilter**(int **srcType**, int **dstType**, InputArray **kernel**, Point **anchor**=Point(-1,-1), double **delta**=0, int **bits**=0)

> **Parameters**
>
>> **srcType** – Source image type.
>>
>> **dstType** – Destination image type that must have as many channels as `srcType` .
>>
>> **kernel** – 2D array of filter coefficients.
>>
>> **anchor** – Anchor point within the kernel. Special value `Point(-1,-1)` means that the anchor is at the kernel center.
>>
>> **delta** – Value added to the filtered results before storing them.
>>
>> **bits** – Number of the fractional bits. The parameter is used when the kernel is an integer matrix representing fixed-point filter coefficients.
>>
>> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.
>>
>> **columnBorderType** – Pixel extrapolation method in the horizontal direction.
>>
>> **borderValue** – Border value used in case of a constant border.

The function returns a pointer to a 2D linear filter for the specified kernel, the source array type, and the destination array type. The function is a higher-level function that calls `getLinearFilter` and passes the retrieved 2D filter to the `FilterEngine` constructor.

**See Also:**

`createSeparableLinearFilter()`, `FilterEngine`, `filter2D()`

# createMorphologyFilter

Creates an engine for non-separable morphological operations.

**C++:** Ptr<FilterEngine> **createMorphologyFilter**(int **op**, int **type**, InputArray **kernel**, Point **anchor**=Point(-1,-1), int **rowBorderType**=BORDER_CONSTANT, int **columnBorderType**=-1, const Scalar& **borderValue**=morphologyDefaultBorderValue() )

**C++:** Ptr<BaseFilter> **getMorphologyFilter**(int **op**, int **type**, InputArray **kernel**, Point **anchor**=Point(-1,-1) )

**C++:** Ptr<BaseRowFilter> **getMorphologyRowFilter**(int **op**, int **type**, int **ksize**, int **anchor**=-1 )

**C++:** Ptr<BaseColumnFilter> **getMorphologyColumnFilter**(int **op**, int **type**, int **ksize**, int **anchor**=-1 )

**C++:** Scalar **morphologyDefaultBorderValue**()

> **Parameters**
>
>> **op** – Morphology operation ID, `MORPH_ERODE` or `MORPH_DILATE` .

> **type** – Input/output image type. The number of channels can be arbitrary. The depth should be one of `CV_8U`, `CV_16U`, `CV_16S`, `CV_32F‘ or ‘'CV_64F`.
>
> **kernel** – 2D 8-bit structuring element for a morphological operation. Non-zero elements indicate the pixels that belong to the element.
>
> **ksize** – Horizontal or vertical structuring element size for separable morphological operations.
>
> **anchor** – Anchor position within the structuring element. Negative values mean that the anchor is at the kernel center.
>
> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.
>
> **columnBorderType** – Pixel extrapolation method in the horizontal direction.
>
> **borderValue** – Border value in case of a constant border. The default value, `morphologyDefaultBorderValue` , has a special meaning. It is transformed $+\inf$ for the erosion and to $-\inf$ for the dilation, which means that the minimum (maximum) is effectively computed only over the pixels that are inside the image.

The functions construct primitive morphological filtering operations or a filter engine based on them. Normally it is enough to use `createMorphologyFilter()` or even higher-level `erode()`, `dilate()` , or `morphologyEx()` . Note that `createMorphologyFilter()` analyzes the structuring element shape and builds a separable morphological filter engine when the structuring element is square.

**See Also:**

`erode()`, `dilate()`, `morphologyEx()`, `FilterEngine`

## createSeparableLinearFilter

Creates an engine for a separable linear filter.

**C++:** Ptr<FilterEngine> **createSeparableLinearFilter**(int **srcType**, int **dstType**, InputArray **rowKernel**, InputArray **columnKernel**, Point **anchor**=Point(-1,-1), double **delta**=0, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1, const Scalar& **borderValue**=Scalar() )

**C++:** Ptr<BaseColumnFilter> **getLinearColumnFilter**(int **bufType**, int **dstType**, InputArray **kernel**, int **anchor**, int **symmetryType**, double **delta**=0, int **bits**=0 )

**C++:** Ptr<BaseRowFilter> **getLinearRowFilter**(int **srcType**, int **bufType**, InputArray **kernel**, int **anchor**, int **symmetryType**)

> **Parameters**
>
> > **srcType** – Source array type.
> >
> > **dstType** – Destination image type that must have as many channels as `srcType` .
> >
> > **bufType** – Intermediate buffer type that must have as many channels as `srcType` .
> >
> > **rowKernel** – Coefficients for filtering each row.
> >
> > **columnKernel** – Coefficients for filtering each column.
> >
> > **anchor** – Anchor position within the kernel. Negative values mean that anchor is positioned at the aperture center.

> **delta** – Value added to the filtered results before storing them.
>
> **bits** – Number of the fractional bits. The parameter is used when the kernel is an integer matrix representing fixed-point filter coefficients.
>
> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.
>
> **columnBorderType** – Pixel extrapolation method in the horizontal direction.
>
> **borderValue** – Border value used in case of a constant border.
>
> **symmetryType** – Type of each row and column kernel. See `getKernelType()` .

The functions construct primitive separable linear filtering operations or a filter engine based on them. Normally it is enough to use `createSeparableLinearFilter()` or even higher-level `sepFilter2D()` . The function `createMorphologyFilter()` is smart enough to figure out the `symmetryType` for each of the two kernels, the intermediate `bufType` and, if filtering can be done in integer arithmetics, the number of `bits` to encode the filter coefficients. If it does not work for you, it is possible to call `getLinearColumnFilter`,``getLinearRowFilter`` directly and then pass them to the `FilterEngine` constructor.

**See Also:**

`sepFilter2D()`, `createLinearFilter()`, `FilterEngine`, `getKernelType()`

## dilate

Dilates an image by using a specific structuring element.

**C++:** void **dilate**(InputArray **src**, OutputArray **dst**, InputArray **kernel**, Point **anchor**=Point(-1,-1), int **iterations**=1, int **borderType**=BORDER_CONSTANT, const Scalar& **borderValue**=morphologyDefaultBorderValue() )

**Python:** cv2.**dilate**(src, kernel[, dst[, anchor[, iterations[, borderType[, borderValue ]]]]]) → dst

**C:** void **cvDilate**(const CvArr* **src**, CvArr* **dst**, IplConvKernel* **element**=NULL, int **iterations**=1 )

**Python:** cv.**Dilate**(src, dst, element=None, iterations=1) → None

> **Parameters**
>
> > **src** – input image; the number of channels can be arbitrary, but the depth should be one of CV_8U, CV_16U, CV_16S, CV_32F' or ''CV_64F.
> >
> > **dst** – output image of the same size and type as `src`.
> >
> > **element** – structuring element used for dilation; if `element=Mat()` , a 3 x 3 rectangular structuring element is used.
> >
> > **anchor** – position of the anchor within the element; default value (-1, -1) means that the anchor is at the element center.
> >
> > **iterations** – number of times dilation is applied.
> >
> > **borderType** – pixel extrapolation method (see `borderInterpolate()` for details).
> >
> > **borderValue** – border value in case of a constant border (see `createMorphologyFilter()` for details).

The function dilates the source image using the specified structuring element that determines the shape of a pixel neighborhood over which the maximum is taken:

$$\mathtt{dst}(x,y) = \max_{(x',y'):\,\mathtt{element}(x',y')\neq 0} \mathtt{src}(x+x', y+y')$$

The function supports the in-place mode. Dilation can be applied several ( `iterations` ) times. In case of multi-channel images, each channel is processed independently.

**See Also:**

erode(), morphologyEx(), createMorphologyFilter()

## erode

Erodes an image by using a specific structuring element.

**C++:** void **erode**(InputArray **src**, OutputArray **dst**, InputArray **kernel**, Point **anchor**=Point(-1,-1), int **iterations**=1, int **borderType**=BORDER_CONSTANT, const Scalar& **borderValue**=morphologyDefaultBorderValue() )

**Python:** cv2.**erode**(src, kernel[, dst[, anchor[, iterations[, borderType[, borderValue ]]]]]) → dst

**C:** void **cvErode**(const CvArr* **src**, CvArr* **dst**, IplConvKernel* **element**=NULL, int **iterations**=1)

**Python:** cv.**Erode**(src, dst, element=None, iterations=1) → None

> **Parameters**
>
> > **src** – input image; the number of channels can be arbitrary, but the depth should be one of `CV_8U, CV_16U, CV_16S, CV_32F‘ or ‘‘CV_64F`.
> >
> > **dst** – output image of the same size and type as `src`.
> >
> > **element** – structuring element used for erosion; if `element=Mat()`, a `3 x 3` rectangular structuring element is used.
> >
> > **anchor** – position of the anchor within the element; default value (`-1, -1`) means that the anchor is at the element center.
> >
> > **iterations** – number of times erosion is applied.
> >
> > **borderType** – pixel extrapolation method (see borderInterpolate() for details).
> >
> > **borderValue** – border value in case of a constant border (see createMorphologyFilter() for details).

The function erodes the source image using the specified structuring element that determines the shape of a pixel neighborhood over which the minimum is taken:

$$\mathtt{dst}(x,y) = \min_{(x',y'):\, \mathtt{element}(x',y') \neq 0} \mathtt{src}(x+x', y+y')$$

The function supports the in-place mode. Erosion can be applied several ( `iterations` ) times. In case of multi-channel images, each channel is processed independently.

**See Also:**

dilate(), morphologyEx(), createMorphologyFilter()

## filter2D

Convolves an image with the kernel.

**C++:** void **filter2D**(InputArray **src**, OutputArray **dst**, int **ddepth**, InputArray **kernel**, Point **anchor**=Point(-1,-1), double **delta**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**filter2D**(src, ddepth, kernel[, dst[, anchor[, delta[, borderType ]]]]) → dst

**C:** void **cvFilter2D**(const CvArr* **src**, CvArr* **dst**, const CvMat* **kernel**, CvPoint **anchor**=cvPoint(-1,-1) )

**Python:** cv.**Filter2D**(src, dst, kernel, anchor=(-1, -1)) → None

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image of the same size and the same number of channels as `src`.
> >
> > **ddepth** –
> >
> > **desired depth of the destination image; if it is negative, it will be the same as `src.depth()`; the following con**
> >
> > > – `src.depth() = CV_8U, ddepth = -1/CV_16S/CV_32F/CV_64F`
> > >
> > > – `src.depth() = CV_16U/CV_16S, ddepth = -1/CV_32F/CV_64F`
> > >
> > > – `src.depth() = CV_32F, ddepth = -1/CV_32F/CV_64F`
> > >
> > > – `src.depth() = CV_64F, ddepth = -1/CV_64F`
> >
> > when `ddepth=-1`, the output image will have the same depth as the source.
> >
> > **kernel** – convolution kernel (or rather a correlation kernel), a single-channel floating point matrix; if you want to apply different kernels to different channels, split the image into separate color planes using `split()` and process them individually.
> >
> > **anchor** – anchor of the kernel that indicates the relative position of a filtered point within the kernel; the anchor should lie within the kernel; default value (-1,-1) means that the anchor is at the kernel center.
> >
> > **delta** – optional value added to the filtered pixels before storing them in `dst`.
> >
> > **borderType** – pixel extrapolation method (see `borderInterpolate()` for details).

The function applies an arbitrary linear filter to an image. In-place operation is supported. When the aperture is partially outside the image, the function interpolates outlier pixel values according to the specified border mode.

The function does actually compute correlation, not the convolution:

$$\text{dst}(x, y) = \sum_{\substack{0 \le x' < \text{kernel.cols,} \\ 0 \le y' < \text{kernel.rows}}} \text{kernel}(x', y') * \text{src}(x + x' - \text{anchor.x}, y + y' - \text{anchor.y})$$

That is, the kernel is not mirrored around the anchor point. If you need a real convolution, flip the kernel using `flip()` and set the new anchor to (`kernel.cols - anchor.x - 1, kernel.rows - anchor.y - 1`).

The function uses the DFT-based algorithm in case of sufficiently large kernels (~``11 x 11`` or larger) and the direct algorithm (that uses the engine retrieved by `createLinearFilter()` ) for small kernels.

**See Also:**

`sepFilter2D()`, `createLinearFilter()`, `dft()`, `matchTemplate()`

## GaussianBlur

Blurs an image using a Gaussian filter.

**C++:** void **GaussianBlur**(InputArray **src**, OutputArray **dst**, Size **ksize**, double **sigmaX**, double **sigmaY**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**GaussianBlur**(src, ksize, sigmaX[, dst[, sigmaY[, borderType ]]]) → dst

> **Parameters**
>
> > **src** – input image; the image can have any number of channels, which are processed independently, but the depth should be CV_8U, CV_16U, CV_16S, CV_32F or CV_64F.
> >
> > **dst** – output image of the same size and type as src.
> >
> > **ksize** – Gaussian kernel size. ksize.width and ksize.height can differ but they both must be positive and odd. Or, they can be zero's and then they are computed from sigma* .
> >
> > **sigmaX** – Gaussian kernel standard deviation in X direction.
> >
> > **sigmaY** – Gaussian kernel standard deviation in Y direction; if sigmaY is zero, it is set to be equal to sigmaX, if both sigmas are zeros, they are computed from ksize.width and ksize.height , respectively (see getGaussianKernel() for details); to fully control the result regardless of possible future modifications of all this semantics, it is recommended to specify all of ksize, sigmaX, and sigmaY.
> >
> > **borderType** – pixel extrapolation method (see borderInterpolate() for details).

The function convolves the source image with the specified Gaussian kernel. In-place filtering is supported.

**See Also:**

sepFilter2D(), filter2D(), blur(), boxFilter(), bilateralFilter(), medianBlur()

## getDerivKernels

Returns filter coefficients for computing spatial image derivatives.

**C++:** void **getDerivKernels**(OutputArray **kx**, OutputArray **ky**, int **dx**, int **dy**, int **ksize**, bool **normalize**=false, int **ktype**=CV_32F )

**Python:** cv2.**getDerivKernels**(dx, dy, ksize[, kx[, ky[, normalize[, ktype ] ] ] ]) → kx, ky

> **Parameters**
>
> > **kx** – Output matrix of row filter coefficients. It has the type ktype .
> >
> > **ky** – Output matrix of column filter coefficients. It has the type ktype .
> >
> > **dx** – Derivative order in respect of x.
> >
> > **dy** – Derivative order in respect of y.
> >
> > **ksize** – Aperture size. It can be CV_SCHARR , 1, 3, 5, or 7.
> >
> > **normalize** – Flag indicating whether to normalize (scale down) the filter coefficients or not. Theoretically, the coefficients should have the denominator $= 2^{ksize*2-dx-dy-2}$ . If you are going to filter floating-point images, you are likely to use the normalized kernels. But if you compute derivatives of an 8-bit image, store the results in a 16-bit image, and wish to preserve all the fractional bits, you may want to set normalize=false .
> >
> > **ktype** – Type of filter coefficients. It can be CV_32f or CV_64F .

The function computes and returns the filter coefficients for spatial image derivatives. When ksize=CV_SCHARR , the Scharr $3 \times 3$ kernels are generated (see Scharr() ). Otherwise, Sobel kernels are generated (see Sobel() ). The filters are normally passed to sepFilter2D() or to createSeparableLinearFilter() .

## getGaussianKernel

Returns Gaussian filter coefficients.

**C++:** Mat **getGaussianKernel**(int **ksize**, double **sigma**, int **ktype**=CV_64F )

**Python:** cv2.**getGaussianKernel**(ksize, sigma[, ktype ]) → retval

>  **Parameters**

>>  **ksize** – Aperture size. It should be odd ( `ksize` $\mod 2 = 1$ ) and positive.

>>  **sigma** – Gaussian standard deviation. If it is non-positive, it is computed from `ksize` as `sigma = 0.3*((ksize-1)*0.5 - 1) + 0.8` .

>>  **ktype** – Type of filter coefficients. It can be `CV_32f` or `CV_64F` .

The function computes and returns the $\texttt{ksize} \times 1$ matrix of Gaussian filter coefficients:

$$G_i = \alpha * e^{-(i-(\texttt{ksize}-1)/2)^2/(2*\texttt{sigma})^2},$$

where $i = 0..\texttt{ksize} - 1$ and $\alpha$ is the scale factor chosen so that $\sum_i G_i = 1$.

Two of such generated kernels can be passed to `sepFilter2D()` or to `createSeparableLinearFilter()`. Those functions automatically recognize smoothing kernels (a symmetrical kernel with sum of weights equal to 1) and handle them accordingly. You may also use the higher-level `GaussianBlur()`.

**See Also:**

`sepFilter2D()`, `createSeparableLinearFilter()`, `getDerivKernels()`, `getStructuringElement()`, `GaussianBlur()`

## getKernelType

Returns the kernel type.

**C++:** int **getKernelType**(InputArray **kernel**, Point **anchor**)

>  **Parameters**

>>  **kernel** – 1D array of the kernel coefficients to analyze.

>>  **anchor** – Anchor position within the kernel.

The function analyzes the kernel coefficients and returns the corresponding kernel type:

- **KERNEL_GENERAL** The kernel is generic. It is used when there is no any type of symmetry or other properties.

- **KERNEL_SYMMETRICAL** The kernel is symmetrical: $\texttt{kernel}_i == \texttt{kernel}_{\texttt{ksize}-i-1}$ , and the anchor is at the center.

- **KERNEL_ASYMMETRICAL** The kernel is asymmetrical: $\texttt{kernel}_i == -\texttt{kernel}_{\texttt{ksize}-i-1}$ , and the anchor is at the center.

- **KERNEL_SMOOTH** All the kernel elements are non-negative and summed to 1. For example, the Gaussian kernel is both smooth kernel and symmetrical, so the function returns `KERNEL_SMOOTH | KERNEL_SYMMETRICAL` .

- **KERNEL_INTEGER** All the kernel coefficients are integer numbers. This flag can be combined with `KERNEL_SYMMETRICAL` or `KERNEL_ASYMMETRICAL` .

## getStructuringElement

Returns a structuring element of the specified size and shape for morphological operations.

**C++:** Mat **getStructuringElement**(int **shape**, Size **ksize**, Point **anchor**=Point(-1,-1))

**Python:** cv2.**getStructuringElement**(shape, ksize[, anchor]) → retval

**C:** IplConvKernel* **cvCreateStructuringElementEx**(int **cols**, int **rows**, int **anchor_x**, int **anchor_y**, int **shape**, int* **values**=NULL )

**Python:** cv.**CreateStructuringElementEx**(cols, rows, anchorX, anchorY, shape, values=None) → kernel

> **Parameters**

>> **shape** – Element shape that could be one of the following:

>> – **MORPH_RECT** - a rectangular structuring element:

$$E_{ij} = 1$$

>> – **MORPH_ELLIPSE** - an elliptic structuring element, that is, a filled ellipse inscribed into the rectangle Rect(0, 0, esize.width, 0.esize.height)

>> – **MORPH_CROSS** - a cross-shaped structuring element:

$$E_{ij} = \begin{cases} 1 & \text{if i=anchor.y or j=anchor.x} \\ 0 & \text{otherwise} \end{cases}$$

>> – **CV_SHAPE_CUSTOM** - custom structuring element (OpenCV 1.x API)

>> **ksize** – Size of the structuring element.

>> **cols** – Width of the structuring element

>> **rows** – Height of the structuring element

>> **anchor** – Anchor position within the element. The default value $(-1, -1)$ means that the anchor is at the center. Note that only the shape of a cross-shaped element depends on the anchor position. In other cases the anchor just regulates how much the result of the morphological operation is shifted.

>> **anchor_x** – x-coordinate of the anchor

>> **anchor_y** – y-coordinate of the anchor

>> **values** – integer array of cols``*``rows elements that specifies the custom shape of the structuring element, when shape=CV_SHAPE_CUSTOM.

The function constructs and returns the structuring element that can be further passed to createMorphologyFilter(), erode(), dilate() or morphologyEx() . But you can also construct an arbitrary binary mask yourself and use it as the structuring element.

---

**Note:** When using OpenCV 1.x C API, the created structuring element IplConvKernel* element must be released in the end using cvReleaseStructuringElement(&element).

---

## medianBlur

Blurs an image using the median filter.

**C++:** void **medianBlur**(InputArray **src**, OutputArray **dst**, int **ksize**)

---

**Python:** cv2.**medianBlur**(src, ksize[, dst ]) → dst

> **Parameters**
>
> > **src** – input 1-, 3-, or 4-channel image; when `ksize` is 3 or 5, the image depth should be `CV_8U`, `CV_16U`, or `CV_32F`, for larger aperture sizes, it can only be `CV_8U`.
> >
> > **dst** – destination array of the same size and type as `src`.
> >
> > **ksize** – aperture linear size; it must be odd and greater than 1, for example: 3, 5, 7 ...

The function smoothes an image using the median filter with the `ksize` × `ksize` aperture. Each channel of a multi-channel image is processed independently. In-place operation is supported.

**See Also:**

bilateralFilter(), blur(), boxFilter(), GaussianBlur()

## morphologyEx

Performs advanced morphological transformations.

**C++:** void **morphologyEx**(InputArray **src**, OutputArray **dst**, int **op**, InputArray **kernel**, Point **anchor**=Point(-1,-1), int **iterations**=1, int **borderType**=BORDER_CONSTANT, const Scalar& **borderValue**=morphologyDefaultBorderValue() )

**Python:** cv2.**morphologyEx**(src, op, kernel[, dst[, anchor[, iterations[, borderType[, borderValue ]]]]]) → dst

**C:** void **cvMorphologyEx**(const CvArr* **src**, CvArr* **dst**, CvArr* **temp**, IplConvKernel* **element**, int **operation**, int **iterations**=1 )

**Python:** cv.**MorphologyEx**(src, dst, temp, element, operation, iterations=1) → None

> **Parameters**
>
> > **src** – Source image. The number of channels can be arbitrary. The depth should be one of `CV_8U`, `CV_16U`, `CV_16S`, `CV_32F'` or `''CV_64F`.
> >
> > **dst** – Destination image of the same size and type as `src` .
> >
> > **element** – Structuring element.
> >
> > **op** – Type of a morphological operation that can be one of the following:
> >
> > – **MORPH_OPEN** - an opening operation
> >
> > – **MORPH_CLOSE** - a closing operation
> >
> > – **MORPH_GRADIENT** - a morphological gradient
> >
> > – **MORPH_TOPHAT** - "top hat"
> >
> > – **MORPH_BLACKHAT** - "black hat"
> >
> > **iterations** – Number of times erosion and dilation are applied.
> >
> > **borderType** – Pixel extrapolation method. See borderInterpolate() for details.
> >
> > **borderValue** – Border value in case of a constant border. The default value has a special meaning. See createMorphologyFilter() for details.

The function can perform advanced morphological transformations using an erosion and dilation as basic operations.

Opening operation:

$$dst = \mathrm{open}(src, \mathtt{element}) = \mathrm{dilate}(\mathrm{erode}(src, \mathtt{element}))$$

---

Closing operation:

$$dst = close(src, element) = erode(dilate(src, element))$$

Morphological gradient:

$$dst = morph\_grad(src, element) = dilate(src, element) - erode(src, element)$$

"Top hat":

$$dst = tophat(src, element) = src - open(src, element)$$

"Black hat":

$$dst = blackhat(src, element) = close(src, element) - src$$

Any of the operations can be done in-place. In case of multi-channel images, each channel is processed independently.

**See Also:**

dilate(), erode(), createMorphologyFilter()

## Laplacian

Calculates the Laplacian of an image.

**C++:** void **Laplacian**(InputArray **src**, OutputArray **dst**, int **ddepth**, int **ksize**=1, double **scale**=1, double **delta**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**Laplacian**(src, ddepth[, dst[, ksize[, scale[, delta[, borderType ]]]]]) → dst

**C:** void **cvLaplace**(const CvArr* **src**, CvArr* **dst**, int **aperture_size**=3 )

**Python:** cv.**Laplace**(src, dst, apertureSize=3) → None

> **Parameters**

>> **src** – Source image.

>> **dst** – Destination image of the same size and the same number of channels as src .

>> **ddepth** – Desired depth of the destination image.

>> **ksize** – Aperture size used to compute the second-derivative filters. See getDerivKernels() for details. The size must be positive and odd.

>> **scale** – Optional scale factor for the computed Laplacian values. By default, no scaling is applied. See getDerivKernels() for details.

>> **delta** – Optional delta value that is added to the results prior to storing them in dst .

>> **borderType** – Pixel extrapolation method. See borderInterpolate() for details.

The function calculates the Laplacian of the source image by adding up the second x and y derivatives calculated using the Sobel operator:

$$dst = \Delta src = \frac{\partial^2 src}{\partial x^2} + \frac{\partial^2 src}{\partial y^2}$$

This is done when ksize > 1 . When ksize == 1 , the Laplacian is computed by filtering the image with the following $3 \times 3$ aperture:

$$\begin{bmatrix} 0 & 1 & 0 \\ 1 & -4 & 1 \\ 0 & 1 & 0 \end{bmatrix}$$

**See Also:**

Sobel(), Scharr()

## pyrDown

Blurs an image and downsamples it.

**C++:** void **pyrDown**(InputArray **src**, OutputArray **dst**, const Size& **dstsize**=Size(), int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**pyrDown**(src[, dst[, dstsize[, borderType ] ] ]) → dst

**C:** void **cvPyrDown**(const CvArr* **src**, CvArr* **dst**, int **filter**=CV_GAUSSIAN_5x5 )

**Python:** cv.**PyrDown**(src, dst, filter=CV_GAUSSIAN_5X5) → None

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image; it has the specified size and the same type as src.
> >
> > **dstsize** – size of the output image; by default, it is computed as Size((src.cols+1)/2, (src.rows+1)/2), but in any case, the following conditions should be satisfied:
> >
> > $$|dstsize.width * 2 - src.cols| \leq 2$$
> > $$|dstsize.height * 2 - src.rows| \leq 2$$

The function performs the downsampling step of the Gaussian pyramid construction. First, it convolves the source image with the kernel:

$$\frac{1}{256} \begin{bmatrix} 1 & 4 & 6 & 4 & 1 \\ 4 & 16 & 24 & 16 & 4 \\ 6 & 24 & 36 & 24 & 6 \\ 4 & 16 & 24 & 16 & 4 \\ 1 & 4 & 6 & 4 & 1 \end{bmatrix}$$

Then, it downsamples the image by rejecting even rows and columns.

## pyrUp

Upsamples an image and then blurs it.

**C++:** void **pyrUp**(InputArray **src**, OutputArray **dst**, const Size& **dstsize**=Size(), int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**pyrUp**(src[, dst[, dstsize[, borderType ] ] ]) → dst

**C:** **cvPyrUp**(const CvArr* **src**, CvArr* **dst**, int **filter**=CV_GAUSSIAN_5x5 )

**Python:** cv.**PyrUp**(src, dst, filter=CV_GAUSSIAN_5X5) → None

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image. It has the specified size and the same type as src .

> **dstsize** – size of the output image; by default, it is computed as `Size(src.cols*2, (src.rows*2)`, but in any case, the following conditions should be satisfied:

$$|dstsize.width - src.cols * 2| \le (\texttt{dstsize.width} \mod 2)$$
$$|dstsize.height - src.rows * 2| \le (\texttt{dstsize.height} \mod 2)$$

The function performs the upsampling step of the Gaussian pyramid construction, though it can actually be used to construct the Laplacian pyramid. First, it upsamples the source image by injecting even zero rows and columns and then convolves the result with the same kernel as in `pyrDown()` multiplied by 4.

## pyrMeanShiftFiltering

Performs initial step of meanshift segmentation of an image.

**C++:** void **pyrMeanShiftFiltering**(InputArray **src**, OutputArray **dst**, double **sp**, double **sr**, int **maxLevel**=1, TermCriteria **termcrit**=TermCriteria( TermCriteria::MAX_ITER+TermCriteria::EPS,5,1) )

**Python:** cv2.**pyrMeanShiftFiltering**(src, sp, sr[, dst[, maxLevel[, termcrit ] ] ]) → dst

**C:** void **cvPyrMeanShiftFiltering**(const CvArr* **src**, CvArr* **dst**, double **sp**, double **sr**, int **max_level**=1, CvTermCriteria **termcrit**=cvTermCriteria(CV_TERMCRIT_ITER+CV_TERMCRIT_EPS,5,1))

**Python:** cv.**PyrMeanShiftFiltering**(src, dst, sp, sr, max_level=1, termcrit=(CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 5, 1)) → None

> **Parameters**

> > **src** – The source 8-bit, 3-channel image.

> > **dst** – The destination image of the same format and the same size as the source.

> > **sp** – The spatial window radius.

> > **sr** – The color window radius.

> > **maxLevel** – Maximum level of the pyramid for the segmentation.

> > **termcrit** – Termination criteria: when to stop meanshift iterations.

The function implements the filtering stage of meanshift segmentation, that is, the output of the function is the filtered "posterized" image with color gradients and fine-grain texture flattened. At every pixel (`X,Y`) of the input image (or down-sized input image, see below) the function executes meanshift iterations, that is, the pixel (`X,Y`) neighborhood in the joint space-color hyperspace is considered:

$$(x, y) : X - \mathsf{sp} \le x \le X + \mathsf{sp}, Y - \mathsf{sp} \le y \le Y + \mathsf{sp}, \|(R, G, B) - (r, g, b)\| \le \mathsf{sr}$$

where (`R,G,B`) and (`r,g,b`) are the vectors of color components at (`X,Y`) and (`x,y`), respectively (though, the algorithm does not depend on the color space used, so any 3-component color space can be used instead). Over the neighborhood the average spatial value (`X',Y'`) and average color vector (`R',G',B'`) are found and they act as the neighborhood center on the next iteration:

$$(X, Y) \ (X', Y'), (R, G, B) \ (R', G', B').$$

After the iterations over, the color components of the initial pixel (that is, the pixel from where the iterations started) are set to the final value (average color at the last iteration):

$$I(X, Y) < -(R*, G*, B*)$$

When `maxLevel > 0`, the gaussian pyramid of `maxLevel+1` levels is built, and the above procedure is run on the smallest layer first. After that, the results are propagated to the larger layer and the iterations are run again only on those pixels where the layer colors differ by more than `sr` from the lower-resolution layer of the pyramid. That makes boundaries of color regions sharper. Note that the results will be actually different from the ones obtained by running the meanshift procedure on the whole original image (i.e. when `maxLevel==0`).

## sepFilter2D

Applies a separable linear filter to an image.

**C++:** void **sepFilter2D**(InputArray **src**, OutputArray **dst**, int **ddepth**, InputArray **kernelX**, InputArray **kernelY**, Point **anchor**=Point(-1,-1), double **delta**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**sepFilter2D**(src, ddepth, kernelX, kernelY[, dst[, anchor[, delta[, borderType ]]]]) → dst

> **Parameters**
>
>> **src** – Source image.
>>
>> **dst** – Destination image of the same size and the same number of channels as `src` .
>>
>> **ddepth** –
>>
>> **Destination image depth. The following combination of `src.depth()` and ddepth are supported:**
>>
>>> – `src.depth() = CV_8U, ddepth = -1/CV_16S/CV_32F/CV_64F`
>>>
>>> – `src.depth() = CV_16U/CV_16S, ddepth = -1/CV_32F/CV_64F`
>>>
>>> – `src.depth() = CV_32F, ddepth = -1/CV_32F/CV_64F`
>>>
>>> – `src.depth() = CV_64F, ddepth = -1/CV_64F`
>>
>> when `ddepth=-1`, the destination image will have the same depth as the source.
>>
>> **kernelX** – Coefficients for filtering each row.
>>
>> **kernelY** – Coefficients for filtering each column.
>>
>> **anchor** – Anchor position within the kernel. The default value $(-1, 1)$ means that the anchor is at the kernel center.
>>
>> **delta** – Value added to the filtered results before storing them.
>>
>> **borderType** – Pixel extrapolation method. See `borderInterpolate()` for details.

The function applies a separable linear filter to the image. That is, first, every row of `src` is filtered with the 1D kernel `kernelX` . Then, every column of the result is filtered with the 1D kernel `kernelY` . The final result shifted by `delta` is stored in `dst` .

**See Also:**

`createSeparableLinearFilter()`, `filter2D()`, `Sobel()`, `GaussianBlur()`, `boxFilter()`, `blur()`

## Smooth

Smooths the image in one of several ways.

**C:** void **cvSmooth**(const CvArr* **src**, CvArr* **dst**, int **smoothtype**=CV_GAUSSIAN, int **size1**=3, int **size2**=0, double **sigma1**=0, double **sigma2**=0 )

**Python:** cv.**Smooth**(src, dst, smoothtype=CV_GAUSSIAN, param1=3, param2=0, param3=0, param4=0) → None

> **Parameters**
>
>> **src** – The source image
>>
>> **dst** – The destination image
>>
>> **smoothtype** – Type of the smoothing:
>>
>> – **CV_BLUR_NO_SCALE** linear convolution with `size1` × `size2` box kernel (all 1's). If you want to smooth different pixels with different-size box kernels, you can use the integral image that is computed using `integral()`
>>
>> – **CV_BLUR** linear convolution with `size1` × `size2` box kernel (all 1's) with subsequent scaling by $1/(\text{size1} \cdot \text{size2})$
>>
>> – **CV_GAUSSIAN** linear convolution with a `size1` × `size2` Gaussian kernel
>>
>> – **CV_MEDIAN** median filter with a `size1` × `size1` square aperture
>>
>> – **CV_BILATERAL** bilateral filter with a `size1` × `size1` square aperture, color sigma= `sigma1` and spatial sigma= `sigma2` . If `size1=0` , the aperture square side is set to `cvRound(sigma2*1.5)*2+1` . Information about bilateral filtering can be found at http://www.dai.ed.ac.uk/CVonline/LOCAL_COPIES/MANDUCHI1/Bilateral_Filtering.html
>>
>> **size1** – The first parameter of the smoothing operation, the aperture width. Must be a positive odd number (1, 3, 5, ...)
>>
>> **size2** – The second parameter of the smoothing operation, the aperture height. Ignored by `CV_MEDIAN` and `CV_BILATERAL` methods. In the case of simple scaled/non-scaled and Gaussian blur if `size2` is zero, it is set to `size1` . Otherwise it must be a positive odd number.
>>
>> **sigma1** – In the case of a Gaussian parameter this parameter may specify Gaussian σ (standard deviation). If it is zero, it is calculated from the kernel size:
>>
>> $$\sigma = 0.3(n/2 - 1) + 0.8 \quad \text{where} \quad n = \begin{array}{l} \text{size1 for horizontal kernel} \\ \text{size2 for vertical kernel} \end{array}$$
>>
>> Using standard sigma for small kernels ( $3 \times 3$ to $7 \times 7$ ) gives better speed. If `sigma1` is not zero, while `size1` and `size2` are zeros, the kernel size is calculated from the sigma (to provide accurate enough operation).

The function smooths an image using one of several methods. Every of the methods has some features and restrictions listed below:

- Blur with no scaling works with single-channel images only and supports accumulation of 8-bit to 16-bit format (similar to `Sobel()` and `Laplacian()`) and 32-bit floating point to 32-bit floating-point format.

- Simple blur and Gaussian blur support 1- or 3-channel, 8-bit and 32-bit floating point images. These two methods can process images in-place.

- Median and bilateral filters work with 1- or 3-channel 8-bit images and can not process images in-place.

---

**Note:** The function is now obsolete. Use GaussianBlur(), blur(), medianBlur() or bilateralFilter().

---

## Sobel

Calculates the first, second, third, or mixed image derivatives using an extended Sobel operator.

**C++:** void **Sobel**(InputArray **src**, OutputArray **dst**, int **ddepth**, int **dx**, int **dy**, int **ksize**=3, double **scale**=1, double **delta**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**Sobel**(src, ddepth, dx, dy[, dst[, ksize[, scale[, delta[, borderType ]]]]]) → dst

**C:** void **cvSobel**(const CvArr* **src**, CvArr* **dst**, int **xorder**, int **yorder**, int **aperture_size**=3 )

**Python:** cv.**Sobel**(src, dst, xorder, yorder, apertureSize=3) → None

> Parameters
>> **src** – input image.
>>
>> **dst** – output image of the same size and the same number of channels as src .
>>
>> **ddepth** –
>>
>> **output image depth; the following combinations of src.depth() and ddepth are supported:**
>>
>>> – src.depth() = CV_8U, ddepth = -1/CV_16S/CV_32F/CV_64F
>>>
>>> – src.depth() = CV_16U/CV_16S, ddepth = -1/CV_32F/CV_64F
>>>
>>> – src.depth() = CV_32F, ddepth = -1/CV_32F/CV_64F
>>>
>>> – src.depth() = CV_64F, ddepth = -1/CV_64F
>>
>> when ddepth=-1, the destination image will have the same depth as the source; in the case of 8-bit input images it will result in truncated derivatives.
>>
>> **xorder** – order of the derivative x.
>>
>> **yorder** – order of the derivative y.
>>
>> **ksize** – size of the extended Sobel kernel; it must be 1, 3, 5, or 7.
>>
>> **scale** – optional scale factor for the computed derivative values; by default, no scaling is applied (see getDerivKernels() for details).
>>
>> **delta** – optional delta value that is added to the results prior to storing them in dst.
>>
>> **borderType** – pixel extrapolation method (see borderInterpolate() for details).

In all cases except one, the ksize $\times$ ksize separable kernel is used to calculate the derivative. When ksize = 1, the $3 \times 1$ or $1 \times 3$ kernel is used (that is, no Gaussian smoothing is done). ksize = 1 can only be used for the first or the second x- or y- derivatives.

There is also the special value ksize = CV_SCHARR (-1) that corresponds to the $3 \times 3$ Scharr filter that may give more accurate results than the $3 \times 3$ Sobel. The Scharr aperture is

$$\begin{bmatrix} -3 & 0 & 3 \\ -10 & 0 & 10 \\ -3 & 0 & 3 \end{bmatrix}$$

for the x-derivative, or transposed for the y-derivative.

---

The function calculates an image derivative by convolving the image with the appropriate kernel:

$$\texttt{dst} = \frac{\partial^{xorder+yorder}\texttt{src}}{\partial x^{xorder}\partial y^{yorder}}$$

The Sobel operators combine Gaussian smoothing and differentiation, so the result is more or less resistant to the noise. Most often, the function is called with ( xorder = 1, yorder = 0, ksize = 3) or ( xorder = 0, yorder = 1, ksize = 3) to calculate the first x- or y- image derivative. The first case corresponds to a kernel of:

$$\begin{bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{bmatrix}$$

The second case corresponds to a kernel of:

$$\begin{bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix}$$

**See Also:**

`Scharr()`, `Laplacian()`, `sepFilter2D()`, `filter2D()`, `GaussianBlur()`, `cartToPolar()`

## Scharr

Calculates the first x- or y- image derivative using Scharr operator.

**C++:** void **Scharr**(InputArray **src**, OutputArray **dst**, int **ddepth**, int **dx**, int **dy**, double **scale**=1, double **delta**=0, int **borderType**=BORDER_DEFAULT )

**Python:** cv2.**Scharr**(src, ddepth, dx, dy$[$, dst$[$, scale$[$, delta$[$, borderType $]]]]$) $\rightarrow$ dst

    **Parameters**

        **src** – input image.

        **dst** – output image of the same size and the same number of channels as `src`.

        **ddepth** – output image depth (see `Sobel()` for the list of supported combination of `src.depth()` and ddepth).

        **dx** – order of the derivative x.

        **dy** – order of the derivative y.

        **scale** – optional scale factor for the computed derivative values; by default, no scaling is applied (see `getDerivKernels()` for details).

        **delta** – optional delta value that is added to the results prior to storing them in `dst`.

        **borderType** – pixel extrapolation method (see `borderInterpolate()` for details).

The function computes the first x- or y- spatial image derivative using the Scharr operator. The call

```
Scharr(src, dst, ddepth, dx, dy, scale, delta, borderType)
```

is equivalent to

```
Sobel(src, dst, ddepth, dx, dy, CV_SCHARR, scale, delta, borderType).
```

**See Also:**

`cartToPolar()`

## 3.2 Geometric Image Transformations

The functions in this section perform various geometrical transformations of 2D images. They do not change the image content but deform the pixel grid and map this deformed grid to the destination image. In fact, to avoid sampling artifacts, the mapping is done in the reverse order, from destination to the source. That is, for each pixel $(x, y)$ of the destination image, the functions compute coordinates of the corresponding "donor" pixel in the source image and copy the pixel value:

$$dst(x, y) = src(f_x(x, y), f_y(x, y))$$

In case when you specify the forward mapping $\langle g_x, g_y \rangle : src \rightarrow dst$ , the OpenCV functions first compute the corresponding inverse mapping $\langle f_x, f_y \rangle : dst \rightarrow src$ and then use the above formula.

The actual implementations of the geometrical transformations, from the most generic `remap()` and to the simplest and the fastest `resize()` , need to solve two main problems with the above formula:

- Extrapolation of non-existing pixels. Similarly to the filtering functions described in the previous section, for some $(x, y)$ , either one of $f_x(x, y)$ , or $f_y(x, y)$ , or both of them may fall outside of the image. In this case, an extrapolation method needs to be used. OpenCV provides the same selection of extrapolation methods as in the filtering functions. In addition, it provides the method `BORDER_TRANSPARENT` . This means that the corresponding pixels in the destination image will not be modified at all.

- Interpolation of pixel values. Usually $f_x(x, y)$ and $f_y(x, y)$ are floating-point numbers. This means that $\langle f_x, f_y \rangle$ can be either an affine or perspective transformation, or radial lens distortion correction, and so on. So, a pixel value at fractional coordinates needs to be retrieved. In the simplest case, the coordinates can be just rounded to the nearest integer coordinates and the corresponding pixel can be used. This is called a nearest-neighbor interpolation. However, a better result can be achieved by using more sophisticated interpolation methods , where a polynomial function is fit into some neighborhood of the computed pixel $(f_x(x, y), f_y(x, y))$ , and then the value of the polynomial at $(f_x(x, y), f_y(x, y))$ is taken as the interpolated pixel value. In OpenCV, you can choose between several interpolation methods. See `resize()` for details.

### convertMaps

Converts image transformation maps from one representation to another.

**C++:** void **convertMaps**(InputArray **map1**, InputArray **map2**, OutputArray **dstmap1**, OutputArray **dstmap2**, int **dstmap1type**, bool **nninterpolation**=false )

**Python:** cv2.**convertMaps**(map1, map2, dstmap1type[, dstmap1[, dstmap2[, nninterpolation ]]]) → dstmap1, dstmap2

> **Parameters**
>
> > **map1** – The first input map of type `CV_16SC2` , `CV_32FC1` , or `CV_32FC2` .
> >
> > **map2** – The second input map of type `CV_16UC1` , `CV_32FC1` , or none (empty matrix), respectively.
> >
> > **dstmap1** – The first output map that has the type `dstmap1type` and the same size as `src` .
> >
> > **dstmap2** – The second output map.
> >
> > **dstmap1type** – Type of the first output map that should be `CV_16SC2` , `CV_32FC1` , or `CV_32FC2` .
> >
> > **nninterpolation** – Flag indicating whether the fixed-point maps are used for the nearest-neighbor or for a more complex interpolation.

The function converts a pair of maps for `remap()` from one representation to another. The following options ( `(map1.type(), map2.type()) → (dstmap1.type(), dstmap2.type())` ) are supported:

- (CV_32FC1, CV_32FC1) → (CV_16SC2, CV_16UC1) . This is the most frequently used conversion operation, in which the original floating-point maps (see `remap()` ) are converted to a more compact and much faster fixed-point representation. The first output array contains the rounded coordinates and the second array (created only when `nninterpolation=false` ) contains indices in the interpolation tables.

- (CV_32FC2) → (CV_16SC2, CV_16UC1) . The same as above but the original maps are stored in one 2-channel matrix.

- Reverse conversion. Obviously, the reconstructed floating-point maps will not be exactly the same as the originals.

**See Also:**

`remap()`, `undistort()`, `initUndistortRectifyMap()`

## getAffineTransform

Calculates an affine transform from three pairs of the corresponding points.

**C++:** Mat **getAffineTransform**(InputArray **src**, InputArray **dst**)

**C++:** Mat **getAffineTransform**(const Point2f **src**[], const Point2f **dst**[])

**Python:** cv2.**getAffineTransform**(src, dst) → retval

**C:** CvMat* **cvGetAffineTransform**(const CvPoint2D32f* **src**, const CvPoint2D32f* **dst**, CvMat* **map_matrix**)

**Python:** cv.**GetAffineTransform**(src, dst, mapMatrix) → None

      **Parameters**

           **src** – Coordinates of triangle vertices in the source image.

           **dst** – Coordinates of the corresponding triangle vertices in the destination image.

The function calculates the $2 \times 3$ matrix of an affine transform so that:

$$\begin{bmatrix} x_i' \\ y_i' \end{bmatrix} = \texttt{map\_matrix} \cdot \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

where

$$dst(i) = (x_i', y_i'), src(i) = (x_i, y_i), i = 0, 1, 2$$

**See Also:**

`warpAffine()`, `transform()`

## getPerspectiveTransform

Calculates a perspective transform from four pairs of the corresponding points.

**C++:** Mat **getPerspectiveTransform**(InputArray **src**, InputArray **dst**)

**C++:** Mat **getPerspectiveTransform**(const Point2f **src**[], const Point2f **dst**[])

**Python:** cv2.**getPerspectiveTransform**(src, dst) → retval

**C:** CvMat* **cvGetPerspectiveTransform**(const CvPoint2D32f* **src**, const CvPoint2D32f* **dst**, CvMat* **map_matrix**)

**Python:** cv.**GetPerspectiveTransform**(src, dst, mapMatrix) → None

> **Parameters**
>
> > **src** – Coordinates of quadrangle vertices in the source image.
> >
> > **dst** – Coordinates of the corresponding quadrangle vertices in the destination image.

The function calculates the $3 \times 3$ matrix of a perspective transform so that:

$$\begin{bmatrix} t_i x_i' \\ t_i y_i' \\ t_i \end{bmatrix} = \texttt{map\_matrix} \cdot \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

where

$$dst(i) = (x_i', y_i'), src(i) = (x_i, y_i), i = 0, 1, 2, 3$$

**See Also:**

findHomography(), warpPerspective(), perspectiveTransform()

## getRectSubPix

Retrieves a pixel rectangle from an image with sub-pixel accuracy.

**C++:** void **getRectSubPix**(InputArray **image**, Size **patchSize**, Point2f **center**, OutputArray **patch**, int **patchType**=-1 )

**Python:** cv2.**getRectSubPix**(image, patchSize, center[, patch[, patchType ]]) → patch

**C:** void **cvGetRectSubPix**(const CvArr* **src**, CvArr* **dst**, CvPoint2D32f **center**)

**Python:** cv.**GetRectSubPix**(src, dst, center) → None

> **Parameters**
>
> > **src** – Source image.
> >
> > **patchSize** – Size of the extracted patch.
> >
> > **center** – Floating point coordinates of the center of the extracted rectangle within the source image. The center must be inside the image.
> >
> > **dst** – Extracted patch that has the size `patchSize` and the same number of channels as `src`.
> >
> > **patchType** – Depth of the extracted pixels. By default, they have the same depth as `src` .

The function getRectSubPix extracts pixels from `src` :

$$dst(x, y) = src(x + \texttt{center.x} - (\texttt{dst.cols} - 1) * 0.5, y + \texttt{center.y} - (\texttt{dst.rows} - 1) * 0.5)$$

where the values of the pixels at non-integer coordinates are retrieved using bilinear interpolation. Every channel of multi-channel images is processed independently. While the center of the rectangle must be inside the image, parts of the rectangle may be outside. In this case, the replication border mode (see borderInterpolate() ) is used to extrapolate the pixel values outside of the image.

**See Also:**

warpAffine(), warpPerspective()

## getRotationMatrix2D

Calculates an affine matrix of 2D rotation.

**C++:** Mat **getRotationMatrix2D**(Point2f **center**, double **angle**, double **scale**)

**Python:** cv2.**getRotationMatrix2D**(center, angle, scale) → retval

**C:** CvMat* **cv2DRotationMatrix**(CvPoint2D32f **center**, double **angle**, double **scale**, CvMat* **map_matrix**)

**Python:** cv.**GetRotationMatrix2D**(center, angle, scale, mapMatrix) → None

> **Parameters**

>> **center** – Center of the rotation in the source image.

>> **angle** – Rotation angle in degrees. Positive values mean counter-clockwise rotation (the coordinate origin is assumed to be the top-left corner).

>> **scale** – Isotropic scale factor.

>> **map_matrix** – The output affine transformation, 2x3 floating-point matrix.

The function calculates the following matrix:

$$\begin{bmatrix} \alpha & \beta & (1-\alpha) \cdot \texttt{center.x} - \beta \cdot \texttt{center.y} \\ -\beta & \alpha & \beta \cdot \texttt{center.x} + (1-\alpha) \cdot \texttt{center.y} \end{bmatrix}$$

where

$$\alpha = \texttt{scale} \cdot \cos \texttt{angle},$$
$$\beta = \texttt{scale} \cdot \sin \texttt{angle}$$

The transformation maps the rotation center to itself. If this is not the target, adjust the shift.

**See Also:**

getAffineTransform(), warpAffine(), transform()

## invertAffineTransform

Inverts an affine transformation.

**C++:** void **invertAffineTransform**(InputArray **M**, OutputArray **iM**)

**Python:** cv2.**invertAffineTransform**(M[, iM]) → iM

> **Parameters**

>> **M** – Original affine transformation.

>> **iM** – Output reverse affine transformation.

The function computes an inverse affine transformation represented by $2 \times 3$ matrix M:

$$\begin{bmatrix} a_{11} & a_{12} & b_1 \\ a_{21} & a_{22} & b_2 \end{bmatrix}$$

The result is also a $2 \times 3$ matrix of the same type as M.

## LogPolar

Remaps an image to log-polar space.

**C:** void **cvLogPolar**(const CvArr* **src**, CvArr* **dst**, CvPoint2D32f **center**, double **M**, int **flags**=CV_INTER_LINEAR+CV_WARP_FILL_OUTLIERS )

**Python:** cv.**LogPolar**(src, dst, center, M, flags=CV_INNER_LINEAR+CV_WARP_FILL_OUTLIERS) → None

> **Parameters**
>> **src** – Source image
>>
>> **dst** – Destination image
>>
>> **center** – The transformation center; where the output precision is maximal
>>
>> **M** – Magnitude scale parameter. See below
>>
>> **flags** – A combination of interpolation methods and the following optional flags:
>>
>>> – **CV_WARP_FILL_OUTLIERS** fills all of the destination image pixels. If some of them correspond to outliers in the source image, they are set to zero
>>>
>>> – **CV_WARP_INVERSE_MAP** See below

The function `cvLogPolar` transforms the source image using the following transformation:

- Forward transformation (`CV_WARP_INVERSE_MAP` is not set):

$$dst(\varphi, \rho) = src(x, y)$$

- Inverse transformation (`CV_WARP_INVERSE_MAP` is set):

$$dst(x, y) = src(\varphi, \rho)$$

where

$$\rho = M \cdot \log \sqrt{x^2 + y^2}, \varphi = atan(y/x)$$

The function emulates the human "foveal" vision and can be used for fast scale and rotation-invariant template matching, for object tracking and so forth. The function can not operate in-place.

## remap

Applies a generic geometrical transformation to an image.

**C++:** void **remap**(InputArray **src**, OutputArray **dst**, InputArray **map1**, InputArray **map2**, int **interpolation**, int **borderMode**=BORDER_CONSTANT, const Scalar& **borderValue**=Scalar())

**Python:** cv2.**remap**(src, map1, map2, interpolation[, dst[, borderMode[, borderValue ]]]) → dst

**C:** void **cvRemap**(const CvArr* **src**, CvArr* **dst**, const CvArr* **mapx**, const CvArr* **mapy**, int **flags**=CV_INTER_LINEAR+CV_WARP_FILL_OUTLIERS, CvScalar **fillval**=cvScalarAll(0) )

---

**Python:** cv.**Remap**(src, dst, mapx, mapy, flags=CV_INNER_LINEAR+CV_WARP_FILL_OUTLIERS, fill-val=(0, 0, 0, 0)) → None

> **Parameters**

>> **src** – Source image.

>> **dst** – Destination image. It has the same size as map1 and the same type as src .

>> **map1** – The first map of either (x,y) points or just x values having the type CV_16SC2 , CV_32FC1 , or CV_32FC2 . See convertMaps() for details on converting a floating point representation to fixed-point for speed.

>> **map2** – The second map of y values having the type CV_16UC1 , CV_32FC1 , or none (empty map if map1 is (x,y) points), respectively.

>> **interpolation** – Interpolation method (see resize() ). The method INTER_AREA is not supported by this function.

>> **borderMode** – Pixel extrapolation method (see borderInterpolate() ). When borderMode=BORDER_TRANSPARENT , it means that the pixels in the destination image that corresponds to the "outliers" in the source image are not modified by the function.

>> **borderValue** – Value used in case of a constant border. By default, it is 0.

The function remap transforms the source image using the specified map:

$$dst(x, y) = src(map_x(x, y), map_y(x, y))$$

where values of pixels with non-integer coordinates are computed using one of available interpolation methods. $map_x$ and $map_y$ can be encoded as separate floating-point maps in $map_1$ and $map_2$ respectively, or interleaved floating-point maps of $(x, y)$ in $map_1$ , or fixed-point maps created by using convertMaps() . The reason you might want to convert from floating to fixed-point representations of a map is that they can yield much faster (~2x) remapping operations. In the converted case, $map_1$ contains pairs (cvFloor(x), cvFloor(y)) and $map_2$ contains indices in a table of interpolation coefficients.

This function cannot operate in-place.

## resize

Resizes an image.

**C++:** void **resize**(InputArray **src**, OutputArray **dst**, Size **dsize**, double **fx**=0, double **fy**=0, int **interpolation**=INTER_LINEAR )

**Python:** cv2.**resize**(src, dsize$\big[$, dst$\big[$, fx$\big[$, fy$\big[$, interpolation $\big]\big]\big]\big]$) → dst

**C:** void **cvResize**(const CvArr* **src**, CvArr* **dst**, int **interpolation**=CV_INTER_LINEAR )

**Python:** cv.**Resize**(src, dst, interpolation=CV_INTER_LINEAR) → None

> **Parameters**

>> **src** – input image.

>> **dst** – output image; it has the size dsize (when it is non-zero) or the size computed from src.size(), fx, and fy; the type of dst is the same as of src.

>> **dsize** – output image size; if it equals zero, it is computed as:

>>> dsize = Size(round(fx*src.cols), round(fy*src.rows))

>> Either dsize or both fx and fy must be non-zero.

**fx** – scale factor along the horizontal axis; when it equals 0, it is computed as

```
(double)dsize.width/src.cols
```

**fy** – scale factor along the vertical axis; when it equals 0, it is computed as

```
(double)dsize.height/src.rows
```

**interpolation** – interpolation method:

– **INTER_NEAREST** - a nearest-neighbor interpolation

– **INTER_LINEAR** - a bilinear interpolation (used by default)

– **INTER_AREA** - resampling using pixel area relation. It may be a preferred method for image decimation, as it gives moire'-free results. But when the image is zoomed, it is similar to the `INTER_NEAREST` method.

– **INTER_CUBIC** - a bicubic interpolation over 4x4 pixel neighborhood

– **INTER_LANCZOS4** - a Lanczos interpolation over 8x8 pixel neighborhood

The function `resize` resizes the image `src` down to or up to the specified size. Note that the initial `dst` type or size are not taken into account. Instead, the size and type are derived from the `src`,``dsize``,``fx`` , and `fy` . If you want to resize `src` so that it fits the pre-created `dst` , you may call the function as follows:

```
// explicitly specify dsize=dst.size(); fx and fy will be computed from that.
resize(src, dst, dst.size(), 0, 0, interpolation);
```

If you want to decimate the image by factor of 2 in each direction, you can call the function this way:

```
// specify fx and fy and let the function compute the destination image size.
resize(src, dst, Size(), 0.5, 0.5, interpolation);
```

To shrink an image, it will generally look best with CV_INTER_AREA interpolation, whereas to enlarge an image, it will generally look best with CV_INTER_CUBIC (slow) or CV_INTER_LINEAR (faster but still looks OK).

**See Also:**

`warpAffine()`, `warpPerspective()`, `remap()`

## warpAffine

Applies an affine transformation to an image.

**C++:** void **warpAffine**(InputArray **src**, OutputArray **dst**, InputArray **M**, Size **dsize**, int **flags**=INTER_LINEAR, int **borderMode**=BORDER_CONSTANT, const Scalar& **borderValue**=Scalar())

**Python:** cv2.**warpAffine**(src, M, dsize[, dst[, flags[, borderMode[, borderValue ] ] ] ]) → dst

**C:** void **cvWarpAffine**(const CvArr* **src**, CvArr* **dst**, const CvMat* **map_matrix**, int **flags**=CV_INTER_LINEAR+CV_WARP_FILL_OUTLIERS, CvScalar **fillval**=cvScalarAll(0) )

**Python:** cv.**WarpAffine**(src, dst, mapMatrix, flags=CV_INTER_LINEAR+CV_WARP_FILL_OUTLIERS, fillval=(0, 0, 0, 0)) → None

**C:** void **cvGetQuadrangleSubPix**(const CvArr* **src**, CvArr* **dst**, const CvMat* **map_matrix**)

**Python:** cv.**GetQuadrangleSubPix**(src, dst, mapMatrix) → None

---

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image that has the size `dsize` and the same type as `src` .
> >
> > **M** – $2 \times 3$ transformation matrix.
> >
> > **dsize** – size of the output image.
> >
> > **flags** – combination of interpolation methods (see `resize()` ) and the optional flag `WARP_INVERSE_MAP` that means that `M` is the inverse transformation ( `dst` $\rightarrow$ `src` ).
> >
> > **borderMode** – pixel extrapolation method (see `borderInterpolate()`); when `borderMode=BORDER_TRANSPARENT` , it means that the pixels in the destination image corresponding to the "outliers" in the source image are not modified by the function.
> >
> > **borderValue** – value used in case of a constant border; by default, it is 0.

The function `warpAffine` transforms the source image using the specified matrix:

$$\mathrm{dst}(x, y) = \mathrm{src}(M_{11}x + M_{12}y + M_{13}, M_{21}x + M_{22}y + M_{23})$$

when the flag `WARP_INVERSE_MAP` is set. Otherwise, the transformation is first inverted with `invertAffineTransform()` and then put in the formula above instead of `M` . The function cannot operate in-place.

**See Also:**

`warpPerspective()`, `resize()`, `remap()`, `getRectSubPix()`, `transform()`

---

**Note:** `cvGetQuadrangleSubPix` is similar to `cvWarpAffine`, but the outliers are extrapolated using replication border mode.

---

## warpPerspective

Applies a perspective transformation to an image.

**C++:** void **warpPerspective**(InputArray **src**, OutputArray **dst**, InputArray **M**, Size **dsize**, int **flags**=INTER_LINEAR, int **borderMode**=BORDER_CONSTANT, const Scalar& **borderValue**=Scalar())

**Python:** cv2.**warpPerspective**(src, M, dsize[, dst[, flags[, borderMode[, borderValue ]]]]) $\rightarrow$ dst

**C:** void **cvWarpPerspective**(const CvArr* **src**, CvArr* **dst**, const CvMat* **map_matrix**, int **flags**=CV_INTER_LINEAR+CV_WARP_FILL_OUTLIERS, CvScalar **fillval**=cvScalarAll(0) )

**Python:** cv.**WarpPerspective**(src, dst, mapMatrix, flags=CV_INNER_LINEAR+CV_WARP_FILL_OUTLIERS, fillval=(0, 0, 0, 0)) $\rightarrow$ None

> **Parameters**
>
> > **src** – input image.
> >
> > **dst** – output image that has the size `dsize` and the same type as `src` .
> >
> > **M** – $3 \times 3$ transformation matrix.
> >
> > **dsize** – size of the output image.
> >
> > **flags** – combination of interpolation methods (`INTER_LINEAR` or `INTER_NEAREST`) and the optional flag `WARP_INVERSE_MAP`, that sets `M` as the inverse transformation ( `dst` $\rightarrow$ `src` ).

> **borderMode** – pixel extrapolation method (`BORDER_CONSTANT` or `BORDER_REPLICATE`).

> **borderValue** – value used in case of a constant border; by default, it equals 0.

The function `warpPerspective` transforms the source image using the specified matrix:

$$\mathtt{dst}(x, y) = \mathtt{src}\left(\frac{M_{11}x + M_{12}y + M_{13}}{M_{31}x + M_{32}y + M_{33}}, \frac{M_{21}x + M_{22}y + M_{23}}{M_{31}x + M_{32}y + M_{33}}\right)$$

when the flag `WARP_INVERSE_MAP` is set. Otherwise, the transformation is first inverted with `invert()` and then put in the formula above instead of `M`. The function cannot operate in-place.

**See Also:**

`warpAffine()`, `resize()`, `remap()`, `getRectSubPix()`, `perspectiveTransform()`

## initUndistortRectifyMap

Computes the undistortion and rectification transformation map.

**C++:** void **initUndistortRectifyMap**(InputArray **cameraMatrix**, InputArray **distCoeffs**, InputArray **R**, InputArray **newCameraMatrix**, Size **size**, int **m1type**, OutputArray **map1**, OutputArray **map2**)

**Python:** cv2.**initUndistortRectifyMap**(cameraMatrix, distCoeffs, R, newCameraMatrix, size, m1type[, map1[, map2]]) → map1, map2

**C:** void **cvInitUndistortRectifyMap**(const CvMat* **camera_matrix**, const CvMat* **dist_coeffs**, const CvMat* **R**, const CvMat* **new_camera_matrix**, CvArr* **mapx**, CvArr* **mapy**)

**C:** void **cvInitUndistortMap**(const CvMat* **camera_matrix**, const CvMat* **distortion_coeffs**, CvArr* **mapx**, CvArr* **mapy**)

**Python:** cv.**InitUndistortRectifyMap**(cameraMatrix, distCoeffs, R, newCameraMatrix, map1, map2) → None

**Python:** cv.**InitUndistortMap**(cameraMatrix, distCoeffs, map1, map2) → None

> **Parameters**

>> **cameraMatrix** – Input camera matrix $A = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$.

>> **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.

>> **R** – Optional rectification transformation in the object space (3x3 matrix). `R1` or `R2`, computed by `stereoRectify()` can be passed here. If the matrix is empty, the identity transformation is assumed. In `cvInitUndistortMap` R assumed to be an identity matrix.

>> **newCameraMatrix** – New camera matrix $A' = \begin{bmatrix} f_x' & 0 & c_x' \\ 0 & f_y' & c_y' \\ 0 & 0 & 1 \end{bmatrix}$.

>> **size** – Undistorted image size.

>> **m1type** – Type of the first output map that can be `CV_32FC1` or `CV_16SC2`. See `convertMaps()` for details.

>> **map1** – The first output map.

>> **map2** – The second output map.

The function computes the joint undistortion and rectification transformation and represents the result in the form of maps for `remap()` . The undistorted image looks like original, as if it is captured with a camera using the camera matrix =newCameraMatrix and zero distortion. In case of a monocular camera, `newCameraMatrix` is usually equal to `cameraMatrix` , or it can be computed by `getOptimalNewCameraMatrix()` for a better control over scaling. In case of a stereo camera, `newCameraMatrix` is normally set to `P1` or `P2` computed by `stereoRectify()` .

Also, this new camera is oriented differently in the coordinate space, according to `R` . That, for example, helps to align two heads of a stereo camera so that the epipolar lines on both images become horizontal and have the same y-coordinate (in case of a horizontally aligned stereo camera).

The function actually builds the maps for the inverse mapping algorithm that is used by `remap()` . That is, for each pixel $(u, v)$ in the destination (corrected and rectified) image, the function computes the corresponding coordinates in the source image (that is, in the original image from camera). The following process is applied:

$$
\begin{aligned}
x &\leftarrow (u - c'_x)/f'_x \\
y &\leftarrow (v - c'_y)/f'_y \\
[X\,Y\,W]^T &\leftarrow R^{-1} * [x\,y\,1]^T \\
x' &\leftarrow X/W \\
y' &\leftarrow Y/W \\
x'' &\leftarrow x'(1 + k_1 r^2 + k_2 r^4 + k_3 r^6) + 2p_1 x' y' + p_2 (r^2 + 2x'^2) \\
y'' &\leftarrow y'(1 + k_1 r^2 + k_2 r^4 + k_3 r^6) + p_1 (r^2 + 2y'^2) + 2p_2 x' y' \\
map_x(u, v) &\leftarrow x'' f_x + c_x \\
map_y(u, v) &\leftarrow y'' f_y + c_y
\end{aligned}
$$

where $(k_1, k_2, p_1, p_2[, k_3])$ are the distortion coefficients.

In case of a stereo camera, this function is called twice: once for each camera head, after `stereoRectify()` , which in its turn is called after `stereoCalibrate()` . But if the stereo camera was not calibrated, it is still possible to compute the rectification transformations directly from the fundamental matrix using `stereoRectifyUncalibrated()` . For each camera, the function computes homography `H` as the rectification transformation in a pixel domain, not a rotation matrix `R` in 3D space. `R` can be computed from `H` as

$$
R = \texttt{cameraMatrix}^{-1} \cdot H \cdot \texttt{cameraMatrix}
$$

where `cameraMatrix` can be chosen arbitrarily.

## getDefaultNewCameraMatrix

Returns the default new camera matrix.

**C++:** Mat **getDefaultNewCameraMatrix**(InputArray **cameraMatrix**, Size **imgsize**=Size(), bool **centerPrincipalPoint**=false )

**Python:** cv2.**getDefaultNewCameraMatrix**(cameraMatrix[, imgsize[, centerPrincipalPoint]]) → retval

> **Parameters**
>
> > **cameraMatrix** – Input camera matrix.
> >
> > **imgsize** – Camera view image size in pixels.
> >
> > **centerPrincipalPoint** – Location of the principal point in the new camera matrix. The parameter indicates whether this location should be at the image center or not.

The function returns the camera matrix that is either an exact copy of the input `cameraMatrix` (when `centerPrinicipalPoint=false` ), or the modified one (when `centerPrincipalPoint=true`).

In the latter case, the new camera matrix will be:

$$\begin{bmatrix} f_x & 0 & (\texttt{imgSize.width} - 1) * 0.5 \\ 0 & f_y & (\texttt{imgSize.height} - 1) * 0.5 \\ 0 & 0 & 1 \end{bmatrix},$$

where $f_x$ and $f_y$ are $(0, 0)$ and $(1, 1)$ elements of `cameraMatrix`, respectively.

By default, the undistortion functions in OpenCV (see `initUndistortRectifyMap()`, `undistort()`) do not move the principal point. However, when you work with stereo, it is important to move the principal points in both views to the same y-coordinate (which is required by most of stereo correspondence algorithms), and may be to the same x-coordinate too. So, you can form the new camera matrix for each view where the principal points are located at the center.

## undistort

Transforms an image to compensate for lens distortion.

**C++:** void **undistort**(InputArray **src**, OutputArray **dst**, InputArray **cameraMatrix**, InputArray **distCoeffs**, InputArray **newCameraMatrix**=noArray() )

**Python:** cv2.**undistort**(src, cameraMatrix, distCoeffs$[$, dst$[$, newCameraMatrix $]]$) $\rightarrow$ dst

**C:** void **cvUndistort2**(const CvArr* **src**, CvArr* **dst**, const CvMat* **camera_matrix**, const CvMat* **distortion_coeffs**, const CvMat* **new_camera_matrix**=0 )

**Python:** cv.**Undistort2**(src, dst, cameraMatrix, distCoeffs) $\rightarrow$ None

> **Parameters**
>
> > **src** – Input (distorted) image.
> >
> > **dst** – Output (corrected) image that has the same size and type as `src` .
> >
> > **cameraMatrix** – Input camera matrix $A = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ .
> >
> > **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
> >
> > **newCameraMatrix** – Camera matrix of the distorted image. By default, it is the same as `cameraMatrix` but you may additionally scale and shift the result by using a different matrix.

The function transforms an image to compensate radial and tangential lens distortion.

The function is simply a combination of `initUndistortRectifyMap()` (with unity R ) and `remap()` (with bilinear interpolation). See the former function for details of the transformation being performed.

Those pixels in the destination image, for which there is no correspondent pixels in the source image, are filled with zeros (black color).

A particular subset of the source image that will be visible in the corrected image can be regulated by `newCameraMatrix` . You can use `getOptimalNewCameraMatrix()` to compute the appropriate `newCameraMatrix` depending on your requirements.

The camera matrix and the distortion parameters can be determined using `calibrateCamera()` . If the resolution of images is different from the resolution used at the calibration stage, $f_x, f_y, c_x$ and $c_y$ need to be scaled accordingly, while the distortion coefficients remain the same.

# undistortPoints

Computes the ideal point coordinates from the observed point coordinates.

**C++:** void **undistortPoints**(InputArray **src**, OutputArray **dst**, InputArray **cameraMatrix**, InputArray **distCoeffs**, InputArray **R**=noArray(), InputArray **P**=noArray())

**C:** void **cvUndistortPoints**(const CvMat* **src**, CvMat* **dst**, const CvMat* **camera_matrix**, const CvMat* **dist_coeffs**, const CvMat* **R**=0, const CvMat* **P**=0 )

**Python:** cv.**UndistortPoints**(src, dst, cameraMatrix, distCoeffs, R=None, P=None) → None

> **Parameters**
>
> > **src** – Observed point coordinates, 1xN or Nx1 2-channel (CV_32FC2 or CV_64FC2).
> >
> > **dst** – Output ideal point coordinates after undistortion and reverse perspective transformation. If matrix P is identity or omitted, dst will contain normalized point coordinates.
> >
> > **cameraMatrix** – Camera matrix $\begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ .
> >
> > **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
> >
> > **R** – Rectification transformation in the object space (3x3 matrix). R1 or R2 computed by stereoRectify() can be passed here. If the matrix is empty, the identity transformation is used.
> >
> > **P** – New camera matrix (3x3) or new projection matrix (3x4). P1 or P2 computed by stereoRectify() can be passed here. If the matrix is empty, the identity new camera matrix is used.

The function is similar to undistort() and initUndistortRectifyMap() but it operates on a sparse set of points instead of a raster image. Also the function performs a reverse transformation to projectPoints() . In case of a 3D object, it does not reconstruct its 3D coordinates, but for a planar object, it does, up to a translation vector, if the proper R is specified.

```
// (u,v) is the input point, (u', v') is the output point
// camera_matrix=[fx 0 cx; 0 fy cy; 0 0 1]
// P=[fx' 0 cx' tx; 0 fy' cy' ty; 0 0 1 tz]
x" = (u - cx)/fx
y" = (v - cy)/fy
(x',y') = undistort(x",y",dist_coeffs)
[X,Y,W]T = R*[x' y' 1]T
x = X/W, y = Y/W
// only performed if P=[fx' 0 cx' [tx]; 0 fy' cy' [ty]; 0 0 1 [tz]] is specified
u' = x*fx' + cx'
v' = y*fy' + cy',
```

where undistort() is an approximate iterative algorithm that estimates the normalized original point coordinates out of the normalized distorted point coordinates ("normalized" means that the coordinates do not depend on the camera matrix).

The function can be used for both a stereo camera head or a monocular camera (when R is empty).

## 3.3 Miscellaneous Image Transformations

### adaptiveThreshold

Applies an adaptive threshold to an array.

**C++:** void **adaptiveThreshold**(InputArray **src**, OutputArray **dst**, double **maxValue**, int **adaptiveMethod**, int **thresholdType**, int **blockSize**, double **C**)

**Python:** cv2.**adaptiveThreshold**(src, maxValue, adaptiveMethod, thresholdType, blockSize, C[, dst]) → dst

**C:** void **cvAdaptiveThreshold**(const CvArr* **src**, CvArr* **dst**, double **max_value**, int **adaptive_method**=CV_ADAPTIVE_THRESH_MEAN_C, int **threshold_type**=CV_THRESH_BINARY, int **block_size**=3, double **param1**=5 )

**Python:** cv.**AdaptiveThreshold**(src, dst, maxValue, adaptive_method=CV_ADAPTIVE_THRESH_MEAN_C, thresholdType=CV_THRESH_BINARY, blockSize=3, param1=5) → None

>        Parameters
>
>>            **src** – Source 8-bit single-channel image.
>>
>>            **dst** – Destination image of the same size and the same type as `src` .
>>
>>            **maxValue** – Non-zero value assigned to the pixels for which the condition is satisfied. See the details below.
>>
>>            **adaptiveMethod** – Adaptive thresholding algorithm to use, `ADAPTIVE_THRESH_MEAN_C` or `ADAPTIVE_THRESH_GAUSSIAN_C` . See the details below.
>>
>>            **thresholdType** – Thresholding type that must be either `THRESH_BINARY` or `THRESH_BINARY_INV` .
>>
>>            **blockSize** – Size of a pixel neighborhood that is used to calculate a threshold value for the pixel: 3, 5, 7, and so on.
>>
>>            **C** – Constant subtracted from the mean or weighted mean (see the details below). Normally, it is positive but may be zero or negative as well.

The function transforms a grayscale image to a binary image according to the formulae:

- **THRESH_BINARY**

$$dst(x, y) = \begin{cases} \texttt{maxValue} & \text{if } src(x, y) > T(x, y) \\ 0 & \text{otherwise} \end{cases}$$

- **THRESH_BINARY_INV**

$$dst(x, y) = \begin{cases} 0 & \text{if } src(x, y) > T(x, y) \\ \texttt{maxValue} & \text{otherwise} \end{cases}$$

where $T(x, y)$ is a threshold calculated individually for each pixel.

- For the method `ADAPTIVE_THRESH_MEAN_C` , the threshold value $T(x, y)$ is a mean of the `blockSize` × `blockSize` neighborhood of $(x, y)$ minus `C` .

- For the method `ADAPTIVE_THRESH_GAUSSIAN_C` , the threshold value $T(x, y)$ is a weighted sum (cross-correlation with a Gaussian window) of the `blockSize` $\times$ `blockSize` neighborhood of $(x, y)$ minus `C` . The default sigma (standard deviation) is used for the specified `blockSize` . See `getGaussianKernel()` .

The function can process the image in-place.

**See Also:**

`threshold()` , `blur()` , `GaussianBlur()`

## cvtColor

Converts an image from one color space to another.

**C++:** void **cvtColor**(InputArray **src**, OutputArray **dst**, int **code**, int **dstCn**=0 )

**Python:** cv2.**cvtColor**(src, code[, dst[, dstCn ] ]) $\rightarrow$ dst

**C:** void **cvCvtColor**(const CvArr* **src**, CvArr* **dst**, int **code**)

**Python:** cv.**CvtColor**(src, dst, code) $\rightarrow$ None

> **Parameters**
>
> > **src** – input image: 8-bit unsigned, 16-bit unsigned ( `CV_16UC...` ), or single-precision floating-point.
> >
> > **dst** – output image of the same size and depth as `src` .
> >
> > **code** – color space conversion code (see the description below).
> >
> > **dstCn** – number of channels in the destination image; if the parameter is 0, the number of the channels is derived automatically from `src` and `code` .

The function converts an input image from one color space to another. In case of a transformation to-from RGB color space, the order of the channels should be specified explicitly (RGB or BGR). Note that the default color format in OpenCV is often referred to as RGB but it is actually BGR (the bytes are reversed). So the first byte in a standard (24-bit) color image will be an 8-bit Blue component, the second byte will be Green, and the third byte will be Red. The fourth, fifth, and sixth bytes would then be the second pixel (Blue, then Green, then Red), and so on.

The conventional ranges for R, G, and B channel values are:

- 0 to 255 for `CV_8U` images
- 0 to 65535 for `CV_16U` images
- 0 to 1 for `CV_32F` images

In case of linear transformations, the range does not matter. But in case of a non-linear transformation, an input RGB image should be normalized to the proper value range to get the correct results, for example, for RGB $\rightarrow$ L*u*v* transformation. For example, if you have a 32-bit floating-point image directly converted from an 8-bit image without any scaling, then it will have the 0..255 value range instead of 0..1 assumed by the function. So, before calling `cvtColor` , you need first to scale the image down:

```
img *= 1./255;
cvtColor(img, img, CV_BGR2Luv);
```

If you use `cvtColor` with 8-bit images, the conversion will have some information lost. For many applications, this will not be noticeable but it is recommended to use 32-bit images in applications that need the full range of colors or that convert an image before an operation and then convert back.

The function can do the following transformations:

- Transformations within RGB space like adding/removing the alpha channel, reversing the channel order, conversion to/from 16-bit RGB color (R5:G6:B5 or R5:G5:B5), as well as conversion to/from grayscale using:

$$\text{RGB[A] to Gray:} \quad Y \leftarrow 0.299 \cdot R + 0.587 \cdot G + 0.114 \cdot B$$

and

$$\text{Gray to RGB[A]:} \quad R \leftarrow Y, G \leftarrow Y, B \leftarrow Y, A \leftarrow 0$$

The conversion from a RGB image to gray is done with:

```
cvtColor(src, bwsrc, CV_RGB2GRAY);
```

More advanced channel reordering can also be done with `mixChannels()`.

- RGB $\leftrightarrow$ CIE XYZ.Rec 709 with D65 white point ( `CV_BGR2XYZ`, `CV_RGB2XYZ`, `CV_XYZ2BGR`, `CV_XYZ2RGB` ):

$$\begin{bmatrix} X \\ Y \\ Z \end{bmatrix} \leftarrow \begin{bmatrix} 0.412453 & 0.357580 & 0.180423 \\ 0.212671 & 0.715160 & 0.072169 \\ 0.019334 & 0.119193 & 0.950227 \end{bmatrix} \cdot \begin{bmatrix} R \\ G \\ B \end{bmatrix}$$

$$\begin{bmatrix} R \\ G \\ B \end{bmatrix} \leftarrow \begin{bmatrix} 3.240479 & -1.53715 & -0.498535 \\ -0.969256 & 1.875991 & 0.041556 \\ 0.055648 & -0.204043 & 1.057311 \end{bmatrix} \cdot \begin{bmatrix} X \\ Y \\ Z \end{bmatrix}$$

X, Y and Z cover the whole value range (in case of floating-point images, Z may exceed 1).

- RGB $\leftrightarrow$ YCrCb JPEG (or YCC) ( `CV_BGR2YCrCb`, `CV_RGB2YCrCb`, `CV_YCrCb2BGR`, `CV_YCrCb2RGB` )

$$Y \leftarrow 0.299 \cdot R + 0.587 \cdot G + 0.114 \cdot B$$

$$Cr \leftarrow (R - Y) \cdot 0.713 + \text{delta}$$

$$Cb \leftarrow (B - Y) \cdot 0.564 + \text{delta}$$

$$R \leftarrow Y + 1.403 \cdot (Cr - \text{delta})$$

$$G \leftarrow Y - 0.344 \cdot (Cr - \text{delta}) - 0.714 \cdot (Cb - \text{delta})$$

$$B \leftarrow Y + 1.773 \cdot (Cb - \text{delta})$$

where

$$\text{delta} = \begin{cases} 128 & \text{for 8-bit images} \\ 32768 & \text{for 16-bit images} \\ 0.5 & \text{for floating-point images} \end{cases}$$

Y, Cr, and Cb cover the whole value range.

- **RGB $\leftrightarrow$ HSV** ( `CV_BGR2HSV`, `CV_RGB2HSV`, `CV_HSV2BGR`, `CV_HSV2RGB` ) In case of 8-bit and 16-bit images, R, G, and B are converted to the floating-point format and scaled to fit the 0 to 1 range.

$$V \leftarrow \max(R, G, B)$$

$$S \leftarrow \begin{cases} \frac{V - \min(R,G,B)}{V} & \text{if } V \neq 0 \\ 0 & \text{otherwise} \end{cases}$$

$$H \leftarrow \begin{cases} 60(G - B)/(V - \min(R, G, B)) & \text{if } V = R \\ 120 + 60(B - R)/(V - \min(R, G, B)) & \text{if } V = G \\ 240 + 60(R - G)/(V - \min(R, G, B)) & \text{if } V = B \end{cases}$$

If $H < 0$ then $H \leftarrow H + 360$ . On output $0 \leq V \leq 1, 0 \leq S \leq 1, 0 \leq H \leq 360$ .

The values are then converted to the destination data type:

– 8-bit images

$$V \leftarrow 255V, S \leftarrow 255S, H \leftarrow H/2 \text{(to fit to 0 to 255)}$$

– 16-bit images (currently not supported)

$$V < -65535V, S < -65535S, H < -H$$

– **32-bit images** H, S, and V are left as is

- **RGB** $\leftrightarrow$ **HLS** ( `CV_BGR2HLS, CV_RGB2HLS, CV_HLS2BGR, CV_HLS2RGB` ). In case of 8-bit and 16-bit images, R, G, and B are converted to the floating-point format and scaled to fit the 0 to 1 range.

$$V_{max} \leftarrow \max(R, G, B)$$

$$V_{min} \leftarrow \min(R, G, B)$$

$$L \leftarrow \frac{V_{max} + V_{min}}{2}$$

$$S \leftarrow \begin{cases} \frac{V_{max} - V_{min}}{V_{max} + V_{min}} & \text{if } L < 0.5 \\ \frac{V_{max} - V_{min}}{2 - (V_{max} + V_{min})} & \text{if } L \geq 0.5 \end{cases}$$

$$H \leftarrow \begin{cases} 60(G - B)/S & \text{if } V_{max} = R \\ 120 + 60(B - R)/S & \text{if } V_{max} = G \\ 240 + 60(R - G)/S & \text{if } V_{max} = B \end{cases}$$

If $H < 0$ then $H \leftarrow H + 360$ . On output $0 \leq L \leq 1, 0 \leq S \leq 1, 0 \leq H \leq 360$ .

The values are then converted to the destination data type:

– 8-bit images

$$V \leftarrow 255 \cdot V, S \leftarrow 255 \cdot S, H \leftarrow H/2 \text{ (to fit to 0 to 255)}$$

– 16-bit images (currently not supported)

$$V < -65535 \cdot V, S < -65535 \cdot S, H < -H$$

  – **32-bit images** H, S, V are left as is

- **RGB** $\leftrightarrow$ **CIE L\*a\*b\* (** `CV_BGR2Lab, CV_RGB2Lab, CV_Lab2BGR, CV_Lab2RGB` **).** In case of 8-bit and 16-bit images, R, G, and B are converted to the floating-point format and scaled to fit the 0 to 1 range.

$$\begin{bmatrix} X \\ Y \\ Z \end{bmatrix} \leftarrow \begin{bmatrix} 0.412453 & 0.357580 & 0.180423 \\ 0.212671 & 0.715160 & 0.072169 \\ 0.019334 & 0.119193 & 0.950227 \end{bmatrix} \cdot \begin{bmatrix} R \\ G \\ B \end{bmatrix}$$

$$X \leftarrow X/X_n, \text{where} X_n = 0.950456$$

$$Z \leftarrow Z/Z_n, \text{where} Z_n = 1.088754$$

$$L \leftarrow \begin{cases} 116 * Y^{1/3} - 16 & \text{for } Y > 0.008856 \\ 903.3 * Y & \text{for } Y \leq 0.008856 \end{cases}$$

$$a \leftarrow 500(f(X) - f(Y)) + delta$$

$$b \leftarrow 200(f(Y) - f(Z)) + delta$$

where

$$f(t) = \begin{cases} t^{1/3} & \text{for } t > 0.008856 \\ 7.787t + 16/116 & \text{for } t \leq 0.008856 \end{cases}$$

and

$$delta = \begin{cases} 128 & \text{for 8-bit images} \\ 0 & \text{for floating-point images} \end{cases}$$

This outputs $0 \leq L \leq 100, -127 \leq a \leq 127, -127 \leq b \leq 127$ . The values are then converted to the destination data type:

  – 8-bit images

$$L \leftarrow L * 255/100, \ a \leftarrow a + 128, \ b \leftarrow b + 128$$

  – **16-bit images** (currently not supported)

  – **32-bit images** L, a, and b are left as is

- **RGB** $\leftrightarrow$ **CIE L\*u\*v\* (** `CV_BGR2Luv, CV_RGB2Luv, CV_Luv2BGR, CV_Luv2RGB` **).** In case of 8-bit and 16-bit images, R, G, and B are converted to the floating-point format and scaled to fit 0 to 1 range.

$$\begin{bmatrix} X \\ Y \\ Z \end{bmatrix} \leftarrow \begin{bmatrix} 0.412453 & 0.357580 & 0.180423 \\ 0.212671 & 0.715160 & 0.072169 \\ 0.019334 & 0.119193 & 0.950227 \end{bmatrix} \cdot \begin{bmatrix} R \\ G \\ B \end{bmatrix}$$

$$L \leftarrow \begin{cases} 116Y^{1/3} & \text{for } Y > 0.008856 \\ 903.3Y & \text{for } Y \leq 0.008856 \end{cases}$$

$$u' \leftarrow 4 * X/(X + 15 * Y + 3Z)$$

$$v' \leftarrow 9 * Y/(X + 15 * Y + 3Z)$$

$$u \leftarrow 13 * L * (u' - u_n) \quad \text{where} \quad u_n = 0.19793943$$

$$v \leftarrow 13 * L * (v' - v_n) \quad \text{where} \quad v_n = 0.46831096$$

This outputs $0 \leq L \leq 100, -134 \leq u \leq 220, -140 \leq v \leq 122$ .

The values are then converted to the destination data type:
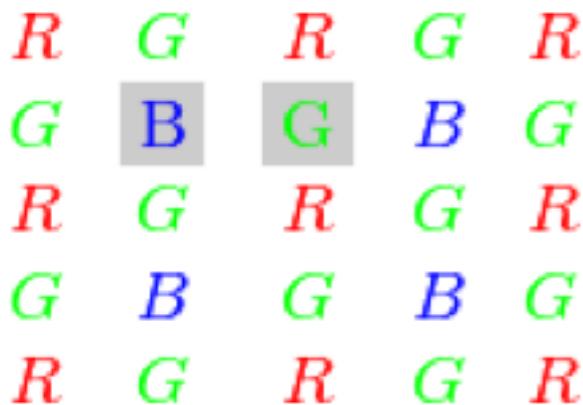
– 8-bit images

$$L \leftarrow 255/100L, \ u \leftarrow 255/354(u + 134), \ v \leftarrow 255/256(v + 140)$$

– **16-bit images** (currently not supported)

– **32-bit images** L, u, and v are left as is

The above formulae for converting RGB to/from various color spaces have been taken from multiple sources on the web, primarily from the Charles Poynton site http://www.poynton.com/ColorFAQ.html

- Bayer → RGB ( `CV_BayerBG2BGR, CV_BayerGB2BGR, CV_BayerRG2BGR, CV_BayerGR2BGR,` `CV_BayerBG2RGB, CV_BayerGB2RGB, CV_BayerRG2RGB, CV_BayerGR2RGB` ). The Bayer pattern is widely used in CCD and CMOS cameras. It enables you to get color pictures from a single plane where R,G, and B pixels (sensors of a particular component) are interleaved as follows:



The output RGB components of a pixel are interpolated from 1, 2, or 4 neighbors of the pixel having the same color. There are several modifications of the above pattern that can be achieved by shifting the pattern one pixel left and/or one pixel up. The two letters $C_1$ and $C_2$ in the conversion constants `CV_Bayer` $C_1 C_2$ `2BGR` and `CV_Bayer` $C_1 C_2$ `2RGB` indicate the particular pattern type. These are components from the second row, second and third columns, respectively. For example, the above pattern has a very popular "BG" type.

# distanceTransform

Calculates the distance to the closest zero pixel for each pixel of the source image.

**C++:** void **distanceTransform**(InputArray **src**, OutputArray **dst**, int **distanceType**, int **maskSize**)

**C++:** void **distanceTransform**(InputArray **src**, OutputArray **dst**, OutputArray **labels**, int **distanceType**, int **maskSize**, int **labelType**=DIST_LABEL_CCOMP )

**Python:** cv2.**distanceTransform**(src, distanceType, maskSize[, dst ]) → dst

**C:** void **cvDistTransform**(const CvArr* **src**, CvArr* **dst**, int **distance_type**=CV_DIST_L2, int **mask_size**=3, const float* **mask**=NULL, CvArr* **labels**=NULL, int **labelType**=CV_DIST_LABEL_CCOMP )

**Python:** cv.**DistTransform**(src, dst, distance_type=CV_DIST_L2, mask_size=3, mask=None, labels=None) → None

> **Parameters**
>
> > **src** – 8-bit, single-channel (binary) source image.
> >
> > **dst** – Output image with calculated distances. It is a 32-bit floating-point, single-channel image of the same size as src .
> >
> > **distanceType** – Type of distance. It can be CV_DIST_L1, CV_DIST_L2 , or CV_DIST_C .
> >
> > **maskSize** – Size of the distance transform mask. It can be 3, 5, or CV_DIST_MASK_PRECISE (the latter option is only supported by the first function). In case of the CV_DIST_L1 or CV_DIST_C distance type, the parameter is forced to 3 because a $3 \times 3$ mask gives the same result as $5 \times 5$ or any larger aperture.
> >
> > **labels** – Optional output 2D array of labels (the discrete Voronoi diagram). It has the type CV_32SC1 and the same size as src . See the details below.
> >
> > **labelType** – Type of the label array to build. If labelType==DIST_LABEL_CCOMP then each connected component of zeros in src (as well as all the non-zero pixels closest to the connected component) will be assigned the same label. If labelType==DIST_LABEL_PIXEL then each zero pixel (and all the non-zero pixels closest to it) gets its own label.

The functions distanceTransform calculate the approximate or precise distance from every binary image pixel to the nearest zero pixel. For zero image pixels, the distance will obviously be zero.

When maskSize == CV_DIST_MASK_PRECISE and distanceType == CV_DIST_L2 , the function runs the algorithm described in [Felzenszwalb04]. This algorithm is parallelized with the TBB library.

In other cases, the algorithm [Borgefors86] is used. This means that for a pixel the function finds the shortest path to the nearest zero pixel consisting of basic shifts: horizontal, vertical, diagonal, or knight's move (the latest is available for a $5 \times 5$ mask). The overall distance is calculated as a sum of these basic distances. Since the distance function should be symmetric, all of the horizontal and vertical shifts must have the same cost (denoted as a ), all the diagonal shifts must have the same cost (denoted as b ), and all knight's moves must have the same cost (denoted as c ). For the CV_DIST_C and CV_DIST_L1 types, the distance is calculated precisely, whereas for CV_DIST_L2 (Euclidean distance) the distance can be calculated only with a relative error (a $5 \times 5$ mask gives more accurate results). For a,``b`` , and c , OpenCV uses the values suggested in the original paper:

| **CV_DIST_C** | $(3 \times 3)$ | **a = 1, b = 1** |
|---|---|---|
| CV_DIST_L1 | $(3 \times 3)$ | a = 1, b = 2 |
| CV_DIST_L2 | $(3 \times 3)$ | a=0.955, b=1.3693 |
| CV_DIST_L2 | $(5 \times 5)$ | a=1, b=1.4, c=2.1969 |

Typically, for a fast, coarse distance estimation CV_DIST_L2, a $3 \times 3$ mask is used. For a more accurate distance estimation CV_DIST_L2 , a $5 \times 5$ mask or the precise algorithm is used. Note that both the precise and the approximate algorithms are linear on the number of pixels.

The second variant of the function does not only compute the minimum distance for each pixel $(x, y)$ but also identifies the nearest connected component consisting of zero pixels (labelType==DIST_LABEL_CCOMP) or the nearest zero pixel (labelType==DIST_LABEL_PIXEL). Index of the component/pixel is stored in $labels(x, y)$. When labelType==DIST_LABEL_CCOMP, the function automatically finds connected components of zero pixels in the input image and marks them with distinct labels. When labelType==DIST_LABEL_CCOMP, the function scans through the input image and marks all the zero pixels with distinct labels.

In this mode, the complexity is still linear. That is, the function provides a very fast way to compute the Voronoi diagram for a binary image. Currently, the second variant can use only the approximate distance transform algorithm, i.e. maskSize=CV_DIST_MASK_PRECISE is not supported yet.

## floodFill

Fills a connected component with the given color.

**C++:** int **floodFill**(InputOutputArray **image**, Point **seedPoint**, Scalar **newVal**, Rect* **rect**=0, Scalar **loDiff**=Scalar(), Scalar **upDiff**=Scalar(), int **flags**=4 )

**C++:** int **floodFill**(InputOutputArray **image**, InputOutputArray **mask**, Point **seedPoint**, Scalar **newVal**, Rect* **rect**=0, Scalar **loDiff**=Scalar(), Scalar **upDiff**=Scalar(), int **flags**=4 )

**Python:** cv2.**floodFill**(image, mask, seedPoint, newVal[, loDiff[, upDiff[, flags ] ] ]) → retval, rect

**C:** void **cvFloodFill**(CvArr* **image**, CvPoint **seed_point**, CvScalar **new_val**, CvScalar **lo_diff**=cvScalarAll(0), CvScalar **up_diff**=cvScalarAll(0), CvConnectedComp* **comp**=NULL, int **flags**=4, CvArr* **mask**=NULL )

**Python:** cv.**FloodFill**(image, seed_point, new_val, lo_diff=(0, 0, 0, 0), up_diff=(0, 0, 0, 0), flags=4, mask=None) → comp

> **Parameters**
>
> > **image** – Input/output 1- or 3-channel, 8-bit, or floating-point image. It is modified by the function unless the FLOODFILL_MASK_ONLY flag is set in the second variant of the function. See the details below.
> >
> > **mask** – (For the second function only) Operation mask that should be a single-channel 8-bit image, 2 pixels wider and 2 pixels taller. The function uses and updates the mask, so you take responsibility of initializing the mask content. Flood-filling cannot go across non-zero pixels in the mask. For example, an edge detector output can be used as a mask to stop filling at edges. It is possible to use the same mask in multiple calls to the function to make sure the filled area does not overlap.
> >
> > ---
> >
> > **Note:** Since the mask is larger than the filled image, a pixel $(x, y)$ in image corresponds to the pixel $(x + 1, y + 1)$ in the mask .
> >
> > ---
> >
> > **seedPoint** – Starting point.
> >
> > **newVal** – New value of the repainted domain pixels.
> >
> > **loDiff** – Maximal lower brightness/color difference between the currently observed pixel and one of its neighbors belonging to the component, or a seed pixel being added to the component.
> >
> > **upDiff** – Maximal upper brightness/color difference between the currently observed pixel and one of its neighbors belonging to the component, or a seed pixel being added to the component.

**rect** – Optional output parameter set by the function to the minimum bounding rectangle of the repainted domain.

**flags** – Operation flags. Lower bits contain a connectivity value, 4 (default) or 8, used within the function. Connectivity determines which neighbors of a pixel are considered. Upper bits can be 0 or a combination of the following flags:

– **FLOODFILL_FIXED_RANGE** If set, the difference between the current pixel and seed pixel is considered. Otherwise, the difference between neighbor pixels is considered (that is, the range is floating).

– **FLOODFILL_MASK_ONLY** If set, the function does not change the image ( `newVal` is ignored), but fills the mask. The flag can be used for the second variant only.

The functions `floodFill` fill a connected component starting from the seed point with the specified color. The connectivity is determined by the color/brightness closeness of the neighbor pixels. The pixel at $(x, y)$ is considered to belong to the repainted domain if:

•

$$\texttt{src}(x', y') - \texttt{loDiff} \leq \texttt{src}(x, y) \leq \texttt{src}(x', y') + \texttt{upDiff}$$

in case of a grayscale image and floating range

•

$$\texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y}) - \texttt{loDiff} \leq \texttt{src}(x, y) \leq \texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y}) + \texttt{upDiff}$$

in case of a grayscale image and fixed range

•

$$\texttt{src}(x', y')_r - \texttt{loDiff}_r \leq \texttt{src}(x, y)_r \leq \texttt{src}(x', y')_r + \texttt{upDiff}_r,$$

$$\texttt{src}(x', y')_g - \texttt{loDiff}_g \leq \texttt{src}(x, y)_g \leq \texttt{src}(x', y')_g + \texttt{upDiff}_g$$

and

$$\texttt{src}(x', y')_b - \texttt{loDiff}_b \leq \texttt{src}(x, y)_b \leq \texttt{src}(x', y')_b + \texttt{upDiff}_b$$

in case of a color image and floating range

•

$$\texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_r - \texttt{loDiff}_r \leq \texttt{src}(x, y)_r \leq \texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_r + \texttt{upDiff}_r,$$

$$\texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_g - \texttt{loDiff}_g \leq \texttt{src}(x, y)_g \leq \texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_g + \texttt{upDiff}_g$$

and

$$\texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_b - \texttt{loDiff}_b \leq \texttt{src}(x, y)_b \leq \texttt{src}(\texttt{seedPoint.x}, \texttt{seedPoint.y})_b + \texttt{upDiff}_b$$

in case of a color image and fixed range

where $\texttt{src}(x', y')$ is the value of one of pixel neighbors that is already known to belong to the component. That is, to be added to the connected component, a color/brightness of the pixel should be close enough to:

• Color/brightness of one of its neighbors that already belong to the connected component in case of a floating range.

• Color/brightness of the seed point in case of a fixed range.

---

Use these functions to either mark a connected component with the specified color in-place, or build a mask and then extract the contour, or copy the region to another image, and so on. Various modes of the function are demonstrated in the `floodfill.cpp` sample.

**See Also:**

`findContours()`

## integral

Calculates the integral of an image.

**C++:** void **integral**(InputArray **src**, OutputArray **sum**, int **sdepth**=-1 )

**C++:** void **integral**(InputArray **src**, OutputArray **sum**, OutputArray **sqsum**, int **sdepth**=-1 )

**C++:** void **integral**(InputArray **src**, OutputArray **sum**, OutputArray **sqsum**, OutputArray **tilted**, int **sdepth**=-1 )

**Python:** cv2.**integral**(src[, sum[, sdepth ] ]) → sum

**Python:** cv2.**integral2**(src[, sum[, sqsum[, sdepth ] ] ]) → sum, sqsum

**Python:** cv2.**integral3**(src[, sum[, sqsum[, tilted[, sdepth ] ] ] ]) → sum, sqsum, tilted

**C:** void **cvIntegral**(const CvArr* **image**, CvArr* **sum**, CvArr* **sqsum**=NULL, CvArr* **tilted_sum**=NULL )

**Python:** cv.**Integral**(image, sum, sqsum=None, tiltedSum=None) → None

> **Parameters**
>
>> **image** – input image as $W \times H$, 8-bit or floating-point (32f or 64f).
>>
>> **sum** – integral image as $(W + 1) \times (H + 1)$ , 32-bit integer or floating-point (32f or 64f).
>>
>> **sqsum** – integral image for squared pixel values; it is $(W + 1) \times (H + 1)$, double-precision floating-point (64f) array.
>>
>> **tilted** – integral for the image rotated by 45 degrees; it is $(W + 1) \times (H + 1)$ array with the same data type as `sum`.
>>
>> **sdepth** – desired depth of the integral and the tilted integral images, `CV_32S`, `CV_32F`, or `CV_64F`.

The functions calculate one or more integral images for the source image as follows:

$$\texttt{sum}(X, Y) = \sum_{x<X, y<Y} \texttt{image}(x, y)$$

$$\texttt{sqsum}(X, Y) = \sum_{x<X, y<Y} \texttt{image}(x, y)^2$$
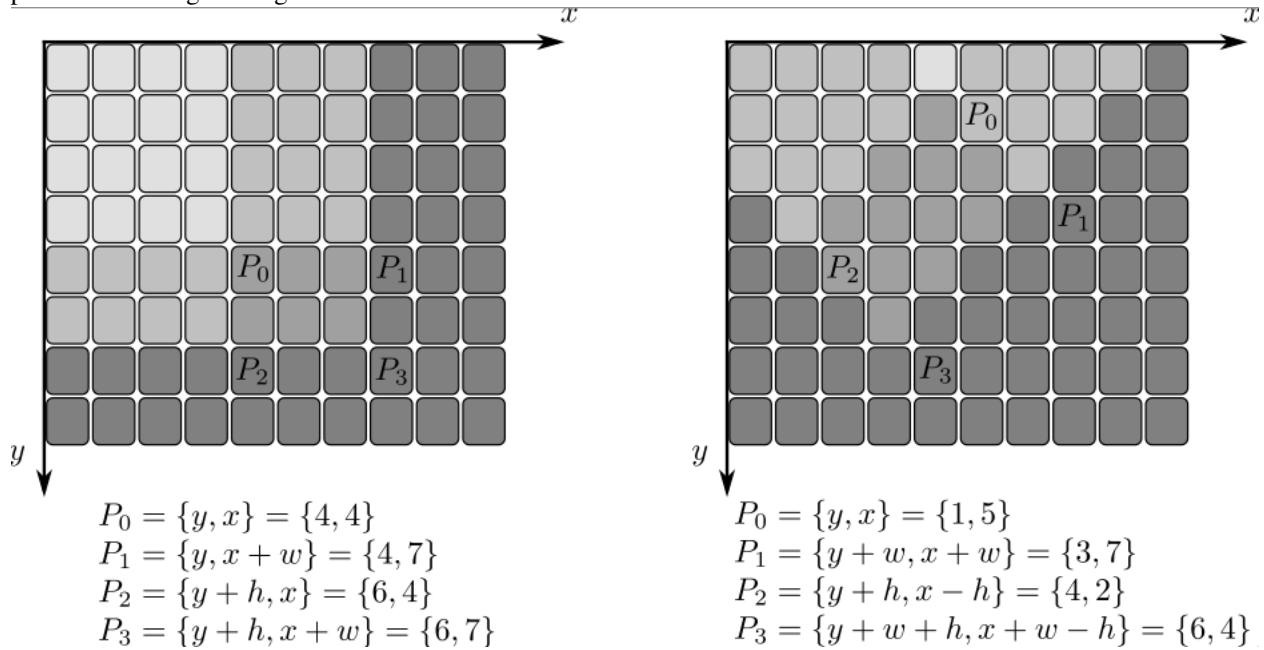
$$\texttt{tilted}(X, Y) = \sum_{y<Y, abs(x-X+1)\leq Y-y-1} \texttt{image}(x, y)$$

Using these integral images, you can calculate sa um, mean, and standard deviation over a specific up-right or rotated rectangular region of the image in a constant time, for example:

$$\sum_{x_1 \leq x < x_2, y_1 \leq y < y_2} \texttt{image}(x, y) = \texttt{sum}(x_2, y_2) - \texttt{sum}(x_1, y_2) - \texttt{sum}(x_2, y_1) + \texttt{sum}(x_1, y_1)$$

It makes possible to do a fast blurring or fast block correlation with a variable window size, for example. In case of multi-channel images, sums for each channel are accumulated independently.

As a practical example, the next figure shows the calculation of the integral of a straight rectangle Rect(3,3,3,2) and of a tilted rectangle Rect(5,1,2,3) . The selected pixels in the original image are shown, as well as the relative pixels in the integral images sum and tilted .



$$P_0 = \{y, x\} = \{4, 4\}$$
$$P_1 = \{y, x + w\} = \{4, 7\}$$
$$P_2 = \{y + h, x\} = \{6, 4\}$$
$$P_3 = \{y + h, x + w\} = \{6, 7\}$$

$$P_0 = \{y, x\} = \{1, 5\}$$
$$P_1 = \{y + w, x + w\} = \{3, 7\}$$
$$P_2 = \{y + h, x - h\} = \{4, 2\}$$
$$P_3 = \{y + w + h, x + w - h\} = \{6, 4\}$$

## threshold

Applies a fixed-level threshold to each array element.

**C++:** double **threshold**(InputArray **src**, OutputArray **dst**, double **thresh**, double **maxval**, int **type**)

**Python:** cv2.**threshold**(src, thresh, maxval, type[, dst ]) → retval, dst

**C:** double **cvThreshold**(const CvArr* **src**, CvArr* **dst**, double **threshold**, double **max_value**, int **threshold_type**)

**Python:** cv.**Threshold**(src, dst, threshold, maxValue, thresholdType) → None

> **Parameters**
>
> > **src** – input array (single-channel, 8-bit or 32-bit floating point).
> >
> > **dst** – output array of the same size and type as src.
> >
> > **thresh** – treshold value.
> >
> > **maxval** – maximum value to use with the THRESH_BINARY and THRESH_BINARY_INV thresholding types.
> >
> > **type** – thresholding type (see the details below).

The function applies fixed-level thresholding to a single-channel array. The function is typically used to get a bi-level (binary) image out of a grayscale image ( compare() could be also used for this purpose) or for removing a noise, that is, filtering out pixels with too small or too large values. There are several types of thresholding supported by the function. They are determined by type :

- **THRESH_BINARY**

$$\texttt{dst}(x, y) = \begin{cases} \texttt{maxval} & \text{if } \texttt{src}(x, y) > \texttt{thresh} \\ 0 & \text{otherwise} \end{cases}$$

- **THRESH_BINARY_INV**

$$\texttt{dst}(x, y) = \begin{cases} 0 & \text{if } \texttt{src}(x, y) > \texttt{thresh} \\ \texttt{maxval} & \text{otherwise} \end{cases}$$

- **THRESH_TRUNC**

$$\texttt{dst}(x, y) = \begin{cases} \texttt{threshold} & \text{if } \texttt{src}(x, y) > \texttt{thresh} \\ \texttt{src}(x, y) & \text{otherwise} \end{cases}$$
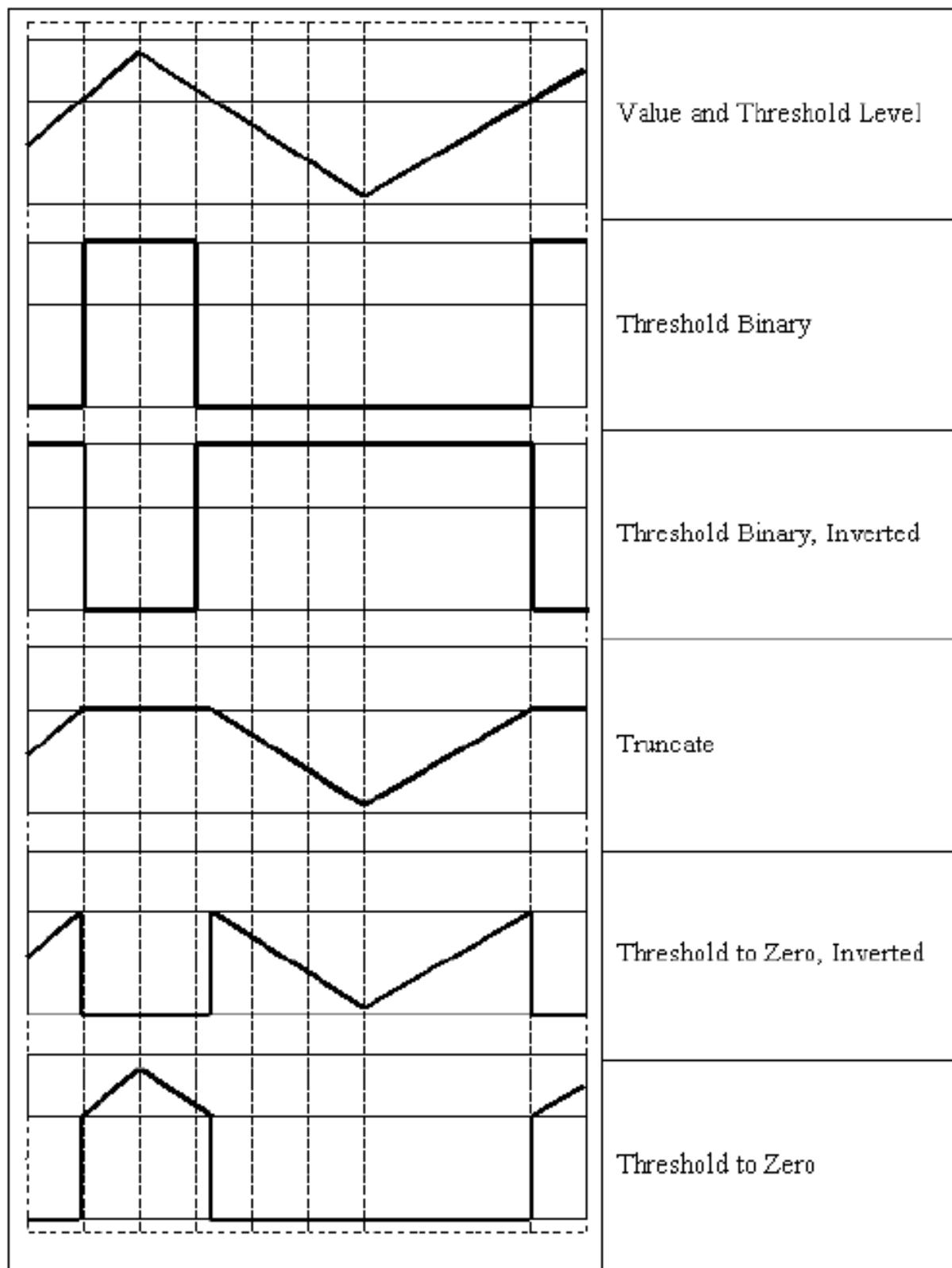
- **THRESH_TOZERO**

$$\texttt{dst}(x, y) = \begin{cases} \texttt{src}(x, y) & \text{if } \texttt{src}(x, y) > \texttt{thresh} \\ 0 & \text{otherwise} \end{cases}$$

- **THRESH_TOZERO_INV**

$$\texttt{dst}(x, y) = \begin{cases} 0 & \text{if } \texttt{src}(x, y) > \texttt{thresh} \\ \texttt{src}(x, y) & \text{otherwise} \end{cases}$$

Also, the special value `THRESH_OTSU` may be combined with one of the above values. In this case, the function determines the optimal threshold value using the Otsu's algorithm and uses it instead of the specified `thresh`. The function returns the computed threshold value. Currently, the Otsu's method is implemented only for 8-bit images.

**See Also:**

adaptiveThreshold(), findContours(), compare(), min(), max()

## watershed

Performs a marker-based image segmentation using the watershed algorithm.

**C++:** void **watershed**(InputArray **image**, InputOutputArray **markers**)

**C:** void **cvWatershed**(const CvArr* **image**, CvArr* **markers**)

**Python:** cv2.**watershed**(image, markers) → None

> **Parameters**
>
>> **image** – Input 8-bit 3-channel image.
>>
>> **markers** – Input/output 32-bit single-channel image (map) of markers. It should have the same size as image .

The function implements one of the variants of watershed, non-parametric marker-based segmentation algorithm, described in [Meyer92].

Before passing the image to the function, you have to roughly outline the desired regions in the image markers with positive (>0) indices. So, every region is represented as one or more connected components with the pixel values 1, 2, 3, and so on. Such markers can be retrieved from a binary mask using findContours() and drawContours() (see the watershed.cpp demo). The markers are "seeds" of the future image regions. All the other pixels in markers , whose relation to the outlined regions is not known and should be defined by the algorithm, should be set to 0's. In the function output, each pixel in markers is set to a value of the "seed" components or to -1 at boundaries between the regions.

Visual demonstration and usage example of the function can be found in the OpenCV samples directory (see the watershed.cpp demo).

---

**Note:** Any two neighbor connected components are not necessarily separated by a watershed boundary (-1's pixels); for example, they can touch each other in the initial marker image passed to the function.

---

**See Also:**

findContours()

## grabCut

Runs the GrabCut algorithm.

**C++:** void **grabCut**(InputArray **img**, InputOutputArray **mask**, Rect **rect**, InputOutputArray **bgdModel**, InputOutputArray **fgdModel**, int **iterCount**, int **mode**=GC_EVAL )

**Python:** cv2.**grabCut**(img, mask, rect, bgdModel, fgdModel, iterCount[, mode ]) → None

> **Parameters**
>
>> **img** – Input 8-bit 3-channel image.
>>
>> **mask** – Input/output 8-bit single-channel mask. The mask is initialized by the function when mode is set to GC_INIT_WITH_RECT. Its elements may have one of following values:
>>
>> – **GC_BGD** defines an obvious background pixels.
>>
>> – **GC_FGD** defines an obvious foreground (object) pixel.
>>
>> – **GC_PR_BGD** defines a possible background pixel.
>>
>> – **GC_PR_BGD** defines a possible foreground pixel.

> > **rect** – ROI containing a segmented object. The pixels outside of the ROI are marked as "obvious background". The parameter is only used when `mode==GC_INIT_WITH_RECT` .
>
> > **bgdModel** – Temporary array for the background model. Do not modify it while you are processing the same image.
>
> > **fgdModel** – Temporary arrays for the foreground model. Do not modify it while you are processing the same image.
>
> > **iterCount** – Number of iterations the algorithm should make before returning the result. Note that the result can be refined with further calls with `mode==GC_INIT_WITH_MASK` or `mode==GC_EVAL` .
>
> > **mode** – Operation mode that could be one of the following:
> >
> > – **GC_INIT_WITH_RECT** The function initializes the state and the mask using the provided rectangle. After that it runs `iterCount` iterations of the algorithm.
> >
> > – **GC_INIT_WITH_MASK** The function initializes the state using the provided mask. Note that `GC_INIT_WITH_RECT` and `GC_INIT_WITH_MASK` can be combined. Then, all the pixels outside of the ROI are automatically initialized with `GC_BGD` .
> >
> > – **GC_EVAL** The value means that the algorithm should just resume.

The function implements the GrabCut image segmentation algorithm. See the sample `grabcut.cpp` to learn how to use the function.

# 3.4 Histograms

## calcHist

Calculates a histogram of a set of arrays.

**C++:** void **calcHist**(const Mat* **images**, int **nimages**, const int* **channels**, InputArray **mask**, OutputArray **hist**, int **dims**, const int* **histSize**, const float** **ranges**, bool **uniform**=true, bool **accumulate**=false )

**C++:** void **calcHist**(const Mat* **images**, int **nimages**, const int* **channels**, InputArray **mask**, SparseMat& **hist**, int **dims**, const int* **histSize**, const float** **ranges**, bool **uniform**=true, bool **accumulate**=false )

**Python:** cv2.**calcHist**(images, channels, mask, histSize, ranges[, hist[, accumulate ]]) → hist

**C:** void **cvCalcHist**(IplImage** **image**, CvHistogram* **hist**, int **accumulate**=0, const CvArr* **mask**=NULL )

**Python:** cv.**CalcHist**(image, hist, accumulate=0, mask=None) → None

> **Parameters**
>
> > **images** – Source arrays. They all should have the same depth, `CV_8U` or `CV_32F` , and the same size. Each of them can have an arbitrary number of channels.
> >
> > **nimages** – Number of source images.
> >
> > **channels** – List of the `dims` channels used to compute the histogram. The first array channels are numerated from 0 to `images[0].channels()-1` , the second array channels are counted from `images[0].channels()` to `images[0].channels() + images[1].channels()-1`, and so on.

> **mask** – Optional mask. If the matrix is not empty, it must be an 8-bit array of the same size as `images[i]` . The non-zero mask elements mark the array elements counted in the histogram.

> **hist** – Output histogram, which is a dense or sparse `dims` -dimensional array.

> **dims** – Histogram dimensionality that must be positive and not greater than `CV_MAX_DIMS` (equal to 32 in the current OpenCV version).

> **histSize** – Array of histogram sizes in each dimension.

> **ranges** – Array of the `dims` arrays of the histogram bin boundaries in each dimension. When the histogram is uniform ( `uniform` =true), then for each dimension `i` it is enough to specify the lower (inclusive) boundary $L_0$ of the 0-th histogram bin and the upper (exclusive) boundary $U_{histSize[i]-1}$ for the last histogram bin `histSize[i]-1` . That is, in case of a uniform histogram each of `ranges[i]` is an array of 2 elements. When the histogram is not uniform ( `uniform=false` ), then each of `ranges[i]` contains `histSize[i]+1` elements: $L_0, U_0 = L_1, U_1 = L_2, ..., U_{histSize[i]-2} = L_{histSize[i]-1}, U_{histSize[i]-1}$ . The array elements, that are not between $L_0$ and $U_{histSize[i]-1}$ , are not counted in the histogram.

> **uniform** – Flag indicating whether the histogram is uniform or not (see above).

> **accumulate** – Accumulation flag. If it is set, the histogram is not cleared in the beginning when it is allocated. This feature enables you to compute a single histogram from several sets of arrays, or to update the histogram in time.

The functions `calcHist` calculate the histogram of one or more arrays. The elements of a tuple used to increment a histogram bin are taken from the corresponding input arrays at the same location. The sample below shows how to compute a 2D Hue-Saturation histogram for a color image.

```cpp
#include <cv.h>
#include <highgui.h>

using namespace cv;

int main( int argc, char** argv )
{
    Mat src, hsv;
    if( argc != 2 || !(src=imread(argv[1], 1)).data )
        return -1;

    cvtColor(src, hsv, CV_BGR2HSV);

    // Quantize the hue to 30 levels
    // and the saturation to 32 levels
    int hbins = 30, sbins = 32;
    int histSize[] = {hbins, sbins};
    // hue varies from 0 to 179, see cvtColor
    float hranges[] = { 0, 180 };
    // saturation varies from 0 (black-gray-white) to
    // 255 (pure spectrum color)
    float sranges[] = { 0, 256 };
    const float* ranges[] = { hranges, sranges };
    MatND hist;
    // we compute the histogram from the 0-th and 1-st channels
    int channels[] = {0, 1};

    calcHist( &hsv, 1, channels, Mat(), // do not use mask
             hist, 2, histSize, ranges,
             true, // the histogram is uniform
```

```
            false );
double maxVal=0;
minMaxLoc(hist, 0, &maxVal, 0, 0);

int scale = 10;
Mat histImg = Mat::zeros(sbins*scale, hbins*10, CV_8UC3);

for( int h = 0; h < hbins; h++ )
    for( int s = 0; s < sbins; s++ )
    {
        float binVal = hist.at<float>(h, s);
        int intensity = cvRound(binVal*255/maxVal);
        rectangle( histImg, Point(h*scale, s*scale),
                    Point( (h+1)*scale - 1, (s+1)*scale - 1),
                    Scalar::all(intensity),
                    CV_FILLED );
    }

namedWindow( "Source", 1 );
imshow( "Source", src );

namedWindow( "H-S Histogram", 1 );
imshow( "H-S Histogram", histImg );
waitKey();
}
```

## calcBackProject

Calculates the back projection of a histogram.

**C++:** void **calcBackProject**(const Mat* **images**, int **nimages**, const int* **channels**, InputArray **hist**, OutputArray **backProject**, const float** **ranges**, double **scale**=1, bool **uniform**=true )

**C++:** void **calcBackProject**(const Mat* **images**, int **nimages**, const int* **channels**, const SparseMat& **hist**, OutputArray **backProject**, const float** **ranges**, double **scale**=1, bool **uniform**=true )

**Python:** cv2.**calcBackProject**(images, channels, hist, ranges, scale[, dst ]) → dst

**C:** void **cvCalcBackProject**(IplImage** **image**, CvArr* **backProject**, const CvHistogram* **hist**)

**Python:** cv.**CalcBackProject**(image, back_project, hist) → None

>    **Parameters**

>    >   **images** – Source arrays. They all should have the same depth, CV_8U or CV_32F , and the same size. Each of them can have an arbitrary number of channels.

>    >   **nimages** – Number of source images.

>    >   **channels** – The list of channels used to compute the back projection. The number of channels must match the histogram dimensionality. The first array channels are numerated from 0 to images[0].channels()-1 , the second array channels are counted from images[0].channels() to images[0].channels() + images[1].channels()-1, and so on.

>    >   **hist** – Input histogram that can be dense or sparse.

> **backProject** – Destination back projection array that is a single-channel array of the same size and depth as `images[0]` .
>
> **ranges** – Array of arrays of the histogram bin boundaries in each dimension. See `calcHist()` .
>
> **scale** – Optional scale factor for the output back projection.
>
> **uniform** – Flag indicating whether the histogram is uniform or not (see above).

The functions `calcBackProject` calculate the back project of the histogram. That is, similarly to `calcHist` , at each location `(x, y)` the function collects the values from the selected channels in the input images and finds the corresponding histogram bin. But instead of incrementing it, the function reads the bin value, scales it by `scale` , and stores in `backProject(x,y)` . In terms of statistics, the function computes probability of each element value in respect with the empirical probability distribution represented by the histogram. See how, for example, you can find and track a bright-colored object in a scene:

1. Before tracking, show the object to the camera so that it covers almost the whole frame. Calculate a hue histogram. The histogram may have strong maximums, corresponding to the dominant colors in the object.

2. When tracking, calculate a back projection of a hue plane of each input video frame using that pre-computed histogram. Threshold the back projection to suppress weak colors. It may also make sense to suppress pixels with non-sufficient color saturation and too dark or too bright pixels.

3. Find connected components in the resulting picture and choose, for example, the largest component.

This is an approximate algorithm of the `CamShift()` color object tracker.

**See Also:**

`calcHist()`

## compareHist

Compares two histograms.

**C++:** double **compareHist**(InputArray **H1**, InputArray **H2**, int **method**)

**C++:** double **compareHist**(const SparseMat& **H1**, const SparseMat& **H2**, int **method**)

**Python:** cv2.**compareHist**(H1, H2, method) → retval

**C:** double **cvCompareHist**(const CvHistogram* **hist1**, const CvHistogram* **hist2**, int **method**)

**Python:** cv.**CompareHist**(hist1, hist2, method) → float

> **Parameters**
>
> > **H1** – First compared histogram.
> >
> > **H2** – Second compared histogram of the same size as `H1` .
> >
> > **method** – Comparison method that could be one of the following:
> >
> > – **CV_COMP_CORREL** Correlation
> >
> > – **CV_COMP_CHISQR** Chi-Square
> >
> > – **CV_COMP_INTERSECT** Intersection
> >
> > – **CV_COMP_BHATTACHARYYA** Bhattacharyya distance
> >
> > – **CV_COMP_HELLINGER** Synonym for `CV_COMP_BHATTACHARYYA`

The functions `compareHist` compare two dense or two sparse histograms using the specified method:

- Correlation (`method=CV_COMP_CORREL`)

$$d(H_1, H_2) = \frac{\sum_I (H_1(I) - \bar{H_1})(H_2(I) - \bar{H_2})}{\sqrt{\sum_I (H_1(I) - \bar{H_1})^2 \sum_I (H_2(I) - \bar{H_2})^2}}$$

where

$$\bar{H_k} = \frac{1}{N} \sum_J H_k(J)$$

and $N$ is a total number of histogram bins.

- Chi-Square (`method=CV_COMP_CHISQR`)

$$d(H_1, H_2) = \sum_I \frac{(H_1(I) - H_2(I))^2}{H_1(I)}$$

- Intersection (`method=CV_COMP_INTERSECT`)

$$d(H_1, H_2) = \sum_I \min(H_1(I), H_2(I))$$

- Bhattacharyya distance (`method=CV_COMP_BHATTACHARYYA` or `method=CV_COMP_HELLINGER`). In fact, OpenCV computes Hellinger distance, which is related to Bhattacharyya coefficient.

$$d(H_1, H_2) = \sqrt{1 - \frac{1}{\sqrt{\bar{H_1}\bar{H_2}N^2}} \sum_I \sqrt{H_1(I) \cdot H_2(I)}}$$

The function returns $d(H_1, H_2)$ .

While the function works well with 1-, 2-, 3-dimensional dense histograms, it may not be suitable for high-dimensional sparse histograms. In such histograms, because of aliasing and sampling problems, the coordinates of non-zero histogram bins can slightly shift. To compare such histograms or more general sparse configurations of weighted points, consider using the `EMD()` function.

## EMD

Computes the "minimal work" distance between two weighted point configurations.

**C++:** float **EMD**(InputArray **signature1**, InputArray **signature2**, int **distType**, InputArray **cost**=noArray(), float* **lowerBound**=0, OutputArray **flow**=noArray() )

**C:** float **cvCalcEMD2**(const CvArr* **signature1**, const CvArr* **signature2**, int **distance_type**, CvDistanceFunction **distance_func**=NULL, const CvArr* **cost_matrix**=NULL, CvArr* **flow**=NULL, float* **lower_bound**=NULL, void* **userdata**=NULL )

**Python:** cv.**CalcEMD2**(signature1, signature2, distance_type, distance_func=None, cost_matrix=None, flow=None, lower_bound=None, userdata=None) → float

       **Parameters**

> **signature1** – First signature, a `size1` × `dims` + 1 floating-point matrix. Each row stores the point weight followed by the point coordinates. The matrix is allowed to have a single column (weights only) if the user-defined cost matrix is used.
>
> **signature2** – Second signature of the same format as `signature1` , though the number of rows may be different. The total weights may be different. In this case an extra "dummy" point is added to either `signature1` or `signature2` .
>
> **distType** – Used metric. `CV_DIST_L1`, `CV_DIST_L2` , and `CV_DIST_C` stand for one of the standard metrics. `CV_DIST_USER` means that a pre-calculated cost matrix `cost` is used.
>
> **distance_func** – Custom distance function supported by the old interface. `CvDistanceFunction` is defined as:
>
> ```
> typedef float (CV_CDECL * CvDistanceFunction)( const float* a,
>                     const float* b, void* userdata );
> ```
>
> where `a` and `b` are point coordinates and `userdata` is the same as the last parameter.
>
> **cost** – User-defined `size1` × `size2` cost matrix. Also, if a cost matrix is used, lower boundary `lowerBound` cannot be calculated because it needs a metric function.
>
> **lowerBound** – Optional input/output parameter: lower boundary of a distance between the two signatures that is a distance between mass centers. The lower boundary may not be calculated if the user-defined cost matrix is used, the total weights of point configurations are not equal, or if the signatures consist of weights only (the signature matrices have a single column). You **must** initialize `*lowerBound` . If the calculated distance between mass centers is greater or equal to `*lowerBound` (it means that the signatures are far enough), the function does not calculate EMD. In any case `*lowerBound` is set to the calculated distance between mass centers on return. Thus, if you want to calculate both distance between mass centers and EMD, `*lowerBound` should be set to 0.
>
> **flow** – Resultant `size1` × `size2` flow matrix: $flow_{i,j}$ is a flow from `i` -th point of `signature1` to `j` -th point of `signature2` .
>
> **userdata** – Optional pointer directly passed to the custom distance function.

The function computes the earth mover distance and/or a lower boundary of the distance between the two weighted point configurations. One of the applications described in [RubnerSept98] is multi-dimensional histogram comparison for image retrieval. EMD is a transportation problem that is solved using some modification of a simplex algorithm, thus the complexity is exponential in the worst case, though, on average it is much faster. In the case of a real metric the lower boundary can be calculated even faster (using linear-time algorithm) and it can be used to determine roughly whether the two signatures are far enough so that they cannot relate to the same object.

## equalizeHist

Equalizes the histogram of a grayscale image.

**C++:** void **equalizeHist**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**equalizeHist**(src[, dst]) → dst

**C:** void **cvEqualizeHist**(const CvArr* **src**, CvArr* **dst**)

> **Parameters**
>
> > **src** – Source 8-bit single channel image.
> >
> > **dst** – Destination image of the same size and type as `src` .

The function equalizes the histogram of the input image using the following algorithm:

1. Calculate the histogram H for `src` .

2. Normalize the histogram so that the sum of histogram bins is 255.

3. Compute the integral of the histogram:

$$H'_i = \sum_{0 \le j < i} H(j)$$

4. Transform the image using $H'$ as a look-up table: $dst(x,y) = H'(src(x,y))$

The algorithm normalizes the brightness and increases the contrast of the image.


## Extra Histogram Functions (C API)

The rest of the section describes additional C functions operating on `CvHistogram`.


## CalcBackProjectPatch

Locates a template within an image by using a histogram comparison.

**C:** void **cvCalcBackProjectPatch**(IplImage** **images**, CvArr* **dst**, CvSize **patch_size**, CvHistogram* **hist**, int **method**, double **factor**)

**Python:** cv.**CalcBackProjectPatch**(images, dst, patch_size, hist, method, factor) → None

> **Parameters**
>
>> **images** – Source images (though, you may pass CvMat** as well).
>>
>> **dst** – Destination image.
>>
>> **patch_size** – Size of the patch slid though the source image.
>>
>> **hist** – Histogram.
>>
>> **method** – Comparison method passed to `CompareHist()` (see the function description).
>>
>> **factor** – Normalization factor for histograms that affects the normalization scale of the destination image. Pass 1 if not sure.
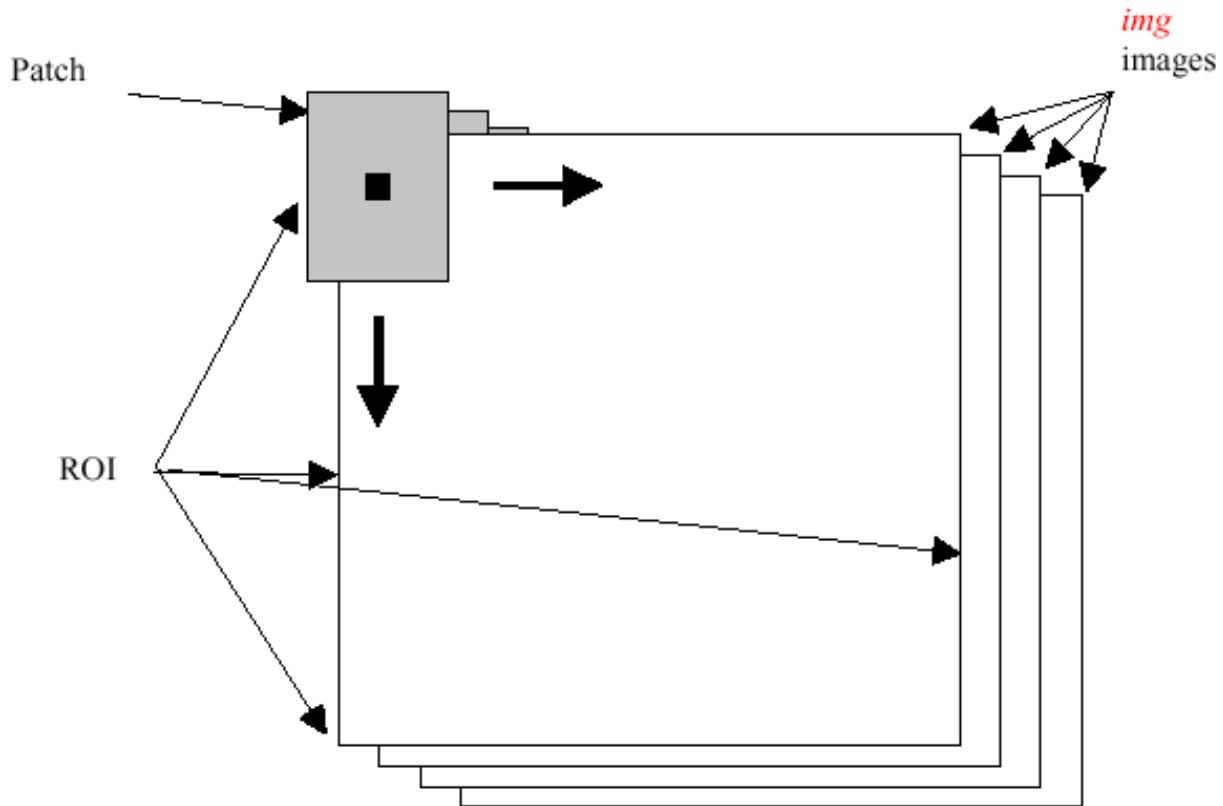
The function calculates the back projection by comparing histograms of the source image patches with the given histogram. The function is similar to `matchTemplate()`, but instead of comparing the raster patch with all its possible positions within the search window, the function `CalcBackProjectPatch` compares histograms. See the algorithm diagram below:

## CalcProbDensity

Divides one histogram by another.

**C:** void **cvCalcProbDensity**(const CvHistogram* **hist1**, const CvHistogram* **hist2**, CvHistogram* **dst_hist**, double **scale**=255 )

**Python:** cv.**CalcProbDensity**(hist1, hist2, dst_hist, scale=255) → None

>   **Parameters**

>>   **hist1** – First histogram (the divisor).

>>   **hist2** – Second histogram.

>>   **dst_hist** – Destination histogram.

>>   **scale** – Scale factor for the destination histogram.

The function calculates the object probability density from two histograms as:

$$\texttt{disthist}(I) = \begin{cases} 0 & \text{if } \texttt{hist1}(I) = 0 \\ \texttt{scale} & \text{if } \texttt{hist1}(I) \neq 0 \text{ and } \texttt{hist2}(I) > \texttt{hist1}(I) \\ \frac{\texttt{hist2}(I) \cdot \texttt{scale}}{\texttt{hist1}(I)} & \text{if } \texttt{hist1}(I) \neq 0 \text{ and } \texttt{hist2}(I) \leq \texttt{hist1}(I) \end{cases}$$

## ClearHist

Clears the histogram.

**C:** void **cvClearHist**(CvHistogram* **hist**)

**Python:** cv.**ClearHist**(hist) → None

> **Parameters**
>
> > **hist** – Histogram.

The function sets all of the histogram bins to 0 in case of a dense histogram and removes all histogram bins in case of a sparse array.

## CopyHist

Copies a histogram.

**C:** void **cvCopyHist**(const CvHistogram* **src**, CvHistogram** **dst**)

> **Parameters**
>
> > **src** – Source histogram.
> >
> > **dst** – Pointer to the destination histogram.

The function makes a copy of the histogram. If the second histogram pointer *dst is NULL, a new histogram of the same size as src is created. Otherwise, both histograms must have equal types and sizes. Then the function copies the bin values of the source histogram to the destination histogram and sets the same bin value ranges as in src.

## CreateHist

Creates a histogram.

**C:** CvHistogram* **cvCreateHist**(int **dims**, int* **sizes**, int **type**, float** **ranges**=NULL, int **uniform**=1 )

**Python:** cv.**CreateHist**(dims, type, ranges=None, uniform=1) → hist

> **Parameters**
>
> > **dims** – Number of histogram dimensions.
> >
> > **sizes** – Array of the histogram dimension sizes.
> >
> > **type** – Histogram representation format. CV_HIST_ARRAY means that the histogram data is represented as a multi-dimensional dense array CvMatND. CV_HIST_SPARSE means that histogram data is represented as a multi-dimensional sparse array CvSparseMat.
> >
> > **ranges** – Array of ranges for the histogram bins. Its meaning depends on the uniform parameter value. The ranges are used when the histogram is calculated or backprojected to determine which histogram bin corresponds to which value/tuple of values from the input image(s).
> >
> > **uniform** – Uniformity flag. If not zero, the histogram has evenly spaced bins and for every $0 <= i < cDims$ ranges[i] is an array of two numbers: lower and upper boundaries for the i-th histogram dimension. The whole range [lower,upper] is then split into dims[i] equal parts to determine the i-th input tuple value ranges for every histogram bin. And if uniform=0 , then the i-th element of the ranges array contains dims[i]+1 elements: $lower_0, upper_0, lower_1, upper_1 = lower_2, ...upper_{dims[i]-1}$ where $lower_j$ and $upper_j$ are lower and upper boundaries of the i-th input tuple value for the j-th bin, respectively. In either case, the input values that are beyond the specified range for a histogram bin are not counted by CalcHist() and filled with 0 by CalcBackProject().

The function creates a histogram of the specified size and returns a pointer to the created histogram. If the array ranges is 0, the histogram bin ranges must be specified later via the function SetHistBinRanges(). Though CalcHist()

and `CalcBackProject()` may process 8-bit images without setting bin ranges, they assume they are equally spaced in 0 to 255 bins.

## GetMinMaxHistValue

Finds the minimum and maximum histogram bins.

**C:** void **cvGetMinMaxHistValue**(const CvHistogram* **hist**, float* **min_value**, float* **max_value**, int* **min_idx**=NULL, int* **max_idx**=NULL )

**Python:** cv.**GetMinMaxHistValue**(hist)-> (min_value, max_value, min_idx, max_idx)

> **Parameters**
>
>> **hist** – Histogram.
>>
>> **min_value** – Pointer to the minimum value of the histogram.
>>
>> **max_value** – Pointer to the maximum value of the histogram.
>>
>> **min_idx** – Pointer to the array of coordinates for the minimum.
>>
>> **max_idx** – Pointer to the array of coordinates for the maximum.

The function finds the minimum and maximum histogram bins and their positions. All of output arguments are optional. Among several extremas with the same value the ones with the minimum index (in the lexicographical order) are returned. In case of several maximums or minimums, the earliest in the lexicographical order (extrema locations) is returned.

## MakeHistHeaderForArray

Makes a histogram out of an array.

**C:** CvHistogram* **cvMakeHistHeaderForArray**(int **dims**, int* **sizes**, CvHistogram* **hist**, float* **data**, float** **ranges**=NULL, int **uniform**=1 )

> **Parameters**
>
>> **dims** – Number of the histogram dimensions.
>>
>> **sizes** – Array of the histogram dimension sizes.
>>
>> **hist** – Histogram header initialized by the function.
>>
>> **data** – Array used to store histogram bins.
>>
>> **ranges** – Histogram bin ranges. See `CreateHist()` for details.
>>
>> **uniform** – Uniformity flag. See `CreateHist()` for details.

The function initializes the histogram, whose header and bins are allocated by the user. `ReleaseHist()` does not need to be called afterwards. Only dense histograms can be initialized this way. The function returns `hist`.

## NormalizeHist

Normalizes the histogram.

**C:** void **cvNormalizeHist**(CvHistogram* **hist**, double **factor**)

**Python:** cv.**NormalizeHist**(hist, factor) → None

> **Parameters**

> **hist** – Pointer to the histogram.

> **factor** – Normalization factor.

The function normalizes the histogram bins by scaling them so that the sum of the bins becomes equal to `factor`.

## ReleaseHist

Releases the histogram.

**C:** void **cvReleaseHist**(CvHistogram** **hist**)

> **Parameters**

>> **hist** – Double pointer to the released histogram.

The function releases the histogram (header and the data). The pointer to the histogram is cleared by the function. If `*hist` pointer is already NULL, the function does nothing.

## SetHistBinRanges

Sets the bounds of the histogram bins.

**C:** void **cvSetHistBinRanges**(CvHistogram* **hist**, float** **ranges**, int **uniform**=1 )

> **Parameters**

>> **hist** – Histogram.

>> **ranges** – Array of bin ranges arrays. See `CreateHist()` for details.

>> **uniform** – Uniformity flag. See `CreateHist()` for details.

This is a standalone function for setting bin ranges in the histogram. For a more detailed description of the parameters `ranges` and `uniform`, see the `CalcHist()` function that can initialize the ranges as well. Ranges for the histogram bins must be set before the histogram is calculated or the backproject of the histogram is calculated.

## ThreshHist

Thresholds the histogram.

**C:** void **cvThreshHist**(CvHistogram* **hist**, double **threshold**)

**Python:** cv.**ThreshHist**(hist, threshold) → None

> **Parameters**

>> **hist** – Pointer to the histogram.

>> **threshold** – Threshold level.

The function clears histogram bins that are below the specified threshold.

# 3.5 Structural Analysis and Shape Descriptors

## moments

Calculates all of the moments up to the third order of a polygon or rasterized shape.

---

**C++:** Moments **moments** (InputArray **array**, bool **binaryImage**=false )

**Python:** cv2.**moments**(array[, binaryImage ]) → retval

**C:** void **cvMoments** (const CvArr* **arr**, CvMoments* **moments**, int **binary**=0 )

**Python:** cv.**Moments**(arr, binary=0) → moments

> **Parameters**

>> **array** – Raster image (single-channel, 8-bit or floating-point 2D array) or an array ( $1 \times N$ or $N \times 1$ ) of 2D points (Point or Point2f ).

>> **binaryImage** – If it is true, all non-zero image pixels are treated as 1's. The parameter is used for images only.

>> **moments** – Output moments.

The function computes moments, up to the 3rd order, of a vector shape or a rasterized shape. The results are returned in the structure Moments defined as:

```cpp
class Moments
{
public:
    Moments();
    Moments(double m00, double m10, double m01, double m20, double m11,
            double m02, double m30, double m21, double m12, double m03 );
    Moments( const CvMoments& moments );
    operator CvMoments() const;

    // spatial moments
    double  m00, m10, m01, m20, m11, m02, m30, m21, m12, m03;
    // central moments
    double  mu20, mu11, mu02, mu30, mu21, mu12, mu03;
    // central normalized moments
    double  nu20, nu11, nu02, nu30, nu21, nu12, nu03;
}
```

In case of a raster image, the spatial moments Moments::$m_{ji}$ are computed as:

$$m_{ji} = \sum_{x,y} \left( \texttt{array}(x, y) \cdot x^j \cdot y^i \right)$$

The central moments Moments::$mu_{ji}$ are computed as:

$$mu_{ji} = \sum_{x,y} \left( \texttt{array}(x, y) \cdot (x - \bar{x})^j \cdot (y - \bar{y})^i \right)$$

where $(\bar{x}, \bar{y})$ is the mass center:

$$\bar{x} = \frac{m_{10}}{m_{00}}, \ \bar{y} = \frac{m_{01}}{m_{00}}$$

The normalized central moments Moments::$nu_{ij}$ are computed as:

$$nu_{ji} = \frac{mu_{ji}}{m_{00}^{(i+j)/2+1}}.$$

**Note:** $mu_{00} = m_{00}$, $nu_{00} = 1$ $nu_{10} = mu_{10} = mu_{01} = mu_{10} = 0$ , hence the values are not stored.

The moments of a contour are defined in the same way but computed using the Green's formula (see http://en.wikipedia.org/wiki/Green_theorem). So, due to a limited raster resolution, the moments computed for a contour are slightly different from the moments computed for the same rasterized contour.

---

**Note:** Since the contour moments are computed using Green formula, you may get seemingly odd results for contours with self-intersections, e.g. a zero area (`m00`) for butterfly-shaped contours.

---

**See Also:**

`contourArea()`, `arcLength()`

## HuMoments

Calculates seven Hu invariants.

**C++:** void **HuMoments**(const Moments& **m**, OutputArray **hu**)

**C++:** void **HuMoments**(const Moments& **moments**, double **hu**[7])

**Python:** cv2.**HuMoments**(m[, hu]) → hu

**C:** void **cvGetHuMoments**(CvMoments* **moments**, CvHuMoments* **hu_moments**)

**Python:** cv.**GetHuMoments**(moments) → hu

      **Parameters**

            **moments** – Input moments computed with `moments()` .

            **hu** – Output Hu invariants.

The function calculates seven Hu invariants (introduced in [Hu62]; see also http://en.wikipedia.org/wiki/Image_moment) defined as:

$$
\begin{aligned}
hu[0] &= \eta_{20} + \eta_{02} \\
hu[1] &= (\eta_{20} - \eta_{02})^2 + 4\eta_{11}^2 \\
hu[2] &= (\eta_{30} - 3\eta_{12})^2 + (3\eta_{21} - \eta_{03})^2 \\
hu[3] &= (\eta_{30} + \eta_{12})^2 + (\eta_{21} + \eta_{03})^2 \\
hu[4] &= (\eta_{30} - 3\eta_{12})(\eta_{30} + \eta_{12})[(\eta_{30} + \eta_{12})^2 - 3(\eta_{21} + \eta_{03})^2] + (3\eta_{21} - \eta_{03})(\eta_{21} + \eta_{03})[3(\eta_{30} + \eta_{12})^2 - (\eta_{21} + \eta_{03})^2] \\
hu[5] &= (\eta_{20} - \eta_{02})[(\eta_{30} + \eta_{12})^2 - (\eta_{21} + \eta_{03})^2] + 4\eta_{11}(\eta_{30} + \eta_{12})(\eta_{21} + \eta_{03}) \\
hu[6] &= (3\eta_{21} - \eta_{03})(\eta_{21} + \eta_{03})[3(\eta_{30} + \eta_{12})^2 - (\eta_{21} + \eta_{03})^2] - (\eta_{30} - 3\eta_{12})(\eta_{21} + \eta_{03})[3(\eta_{30} + \eta_{12})^2 - (\eta_{21} + \eta_{03})^2]
\end{aligned}
$$

where $\eta_{ji}$ stands for `Moments::nu`$_{ji}$ .

These values are proved to be invariants to the image scale, rotation, and reflection except the seventh one, whose sign is changed by reflection. This invariance is proved with the assumption of infinite image resolution. In case of raster images, the computed Hu invariants for the original and transformed images are a bit different.

**See Also:**

`matchShapes()`

## findContours

Finds contours in a binary image.

**C++:** void **findContours**(InputOutputArray **image**, OutputArrayOfArrays **contours**, OutputArray **hierarchy**, int **mode**, int **method**, Point **offset**=Point())

**C++:** void **findContours**(InputOutputArray **image**, OutputArrayOfArrays **contours**, int **mode**, int **method**, Point **offset**=Point())

**Python:** cv2.**findContours**(image, mode, method[, contours[, hierarchy[, offset ]]]) → contours, hierarchy

**C:** int **cvFindContours**(CvArr* **image**, CvMemStorage* **storage**, CvSeq** **first_contour**, int **header_size**=sizeof(CvContour), int **mode**=CV_RETR_LIST, int **method**=CV_CHAIN_APPROX_SIMPLE, CvPoint **offset**=cvPoint(0,0) )

**Python:** cv.**FindContours**(image, storage, mode=CV_RETR_LIST, method=CV_CHAIN_APPROX_SIMPLE, offset=(0, 0)) → contours

> Parameters
>
>> **image** – Source, an 8-bit single-channel image. Non-zero pixels are treated as 1's. Zero pixels remain 0's, so the image is treated as `binary` . You can use `compare()` , `inRange()` , `threshold()` , `adaptiveThreshold()` , `Canny()` , and others to create a binary image out of a grayscale or color one. The function modifies the `image` while extracting the contours.
>>
>> **contours** – Detected contours. Each contour is stored as a vector of points.
>>
>> **hierarchy** – Optional output vector containing information about the image topology. It has as many elements as the number of contours. For each contour `contours[i]` , the elements `hierarchy[i][0]` , `hierarchy[i][1]` , `hierarchy[i][2]` , and `hierarchy[i][3]` are set to 0-based indices in `contours` of the next and previous contours at the same hierarchical level: the first child contour and the parent contour, respectively. If for a contour `i` there are no next, previous, parent, or nested contours, the corresponding elements of `hierarchy[i]` will be negative.
>>
>> **mode** – Contour retrieval mode (if you use Python see also a note below).
>>
>>> – **CV_RETR_EXTERNAL** retrieves only the extreme outer contours. It sets `hierarchy[i][2]=hierarchy[i][3]=-1` for all the contours.
>>>
>>> – **CV_RETR_LIST** retrieves all of the contours without establishing any hierarchical relationships.
>>>
>>> – **CV_RETR_CCOMP** retrieves all of the contours and organizes them into a two-level hierarchy. At the top level, there are external boundaries of the components. At the second level, there are boundaries of the holes. If there is another contour inside a hole of a connected component, it is still put at the top level.
>>>
>>> – **CV_RETR_TREE** retrieves all of the contours and reconstructs a full hierarchy of nested contours. This full hierarchy is built and shown in the OpenCV `contours.c` demo.
>>
>> **method** – Contour approximation method (if you use Python see also a note below).
>>
>>> – **CV_CHAIN_APPROX_NONE** stores absolutely all the contour points. That is, any 2 subsequent points `(x1,y1)` and `(x2,y2)` of the contour will be either horizontal, vertical or diagonal neighbors, that is, `max(abs(x1-x2),abs(y2-y1))==1`.
>>>
>>> – **CV_CHAIN_APPROX_SIMPLE** compresses horizontal, vertical, and diagonal segments and leaves only their end points. For example, an up-right rectangular contour is encoded with 4 points.
>>>
>>> – **CV_CHAIN_APPROX_TC89_L1,CV_CHAIN_APPROX_TC89_KCOS** applies one of the flavors of the Teh-Chin chain approximation algorithm. See [TehChin89] for details.
>>
>> **offset** – Optional offset by which every contour point is shifted. This is useful if the contours are extracted from the image ROI and then they should be analyzed in the whole image context.

The function retrieves contours from the binary image using the algorithm [Suzuki85]. The contours are a useful tool for shape analysis and object detection and recognition. See `squares.c` in the OpenCV sample directory.

---

**Note:** Source `image` is modified by this function. Also, the function does not take into account 1-pixel border of the image (it's filled with 0's and used for neighbor analysis in the algorithm), therefore the contours touching the image border will be clipped.

---

**Note:** If you use the new Python interface then the `CV_` prefix has to be omitted in contour retrieval mode and contour approximation method parameters (for example, use `cv2.RETR_LIST` and `cv2.CHAIN_APPROX_NONE` parameters). If you use the old Python interface then these parameters have the `CV_` prefix (for example, use `cv.CV_RETR_LIST` and `cv.CV_CHAIN_APPROX_NONE`).

---

## drawContours

Draws contours outlines or filled contours.

**C++:** void **drawContours**(InputOutputArray **image**, InputArrayOfArrays **contours**, int **contourIdx**, const Scalar& **color**, int **thickness**=1, int **lineType**=8, InputArray **hierarchy**=noArray(), int **maxLevel**=INT_MAX, Point **offset**=Point() )

**Python:** cv2.**drawContours**(image, contours, contourIdx, color[, thickness[, lineType[, hierarchy[, maxLevel[, offset ]]]]]) → None

**C:** void **cvDrawContours**(CvArr* **img**, CvSeq* **contour**, CvScalar **externalColor**, CvScalar **holeColor**, int **maxLevel**, int **thickness**=1, int **lineType**=8 )

**Python:** cv.**DrawContours**(img, contour, external_color, hole_color, max_level, thickness=1, lineType=8, offset=(0, 0)) → None

>    **Parameters**

>    >    **image** – Destination image.

>    >    **contours** – All the input contours. Each contour is stored as a point vector.

>    >    **contourIdx** – Parameter indicating a contour to draw. If it is negative, all the contours are drawn.

>    >    **color** – Color of the contours.

>    >    **thickness** – Thickness of lines the contours are drawn with. If it is negative (for example, `thickness=CV_FILLED` ), the contour interiors are drawn.

>    >    **lineType** – Line connectivity. See `line()` for details.

>    >    **hierarchy** – Optional information about hierarchy. It is only needed if you want to draw only some of the contours (see `maxLevel` ).

>    >    **maxLevel** – Maximal level for drawn contours. If it is 0, only the specified contour is drawn. If it is 1, the function draws the contour(s) and all the nested contours. If it is 2, the function draws the contours, all the nested contours, all the nested-to-nested contours, and so on. This parameter is only taken into account when there is `hierarchy` available.

>    >    **offset** – Optional contour shift parameter. Shift all the drawn contours by the specified $offset = (dx, dy)$ .

>    >    **contour** – Pointer to the first contour.

>    >    **externalColor** – Color of external contours.

>> **holeColor** – Color of internal contours (holes).

The function draws contour outlines in the image if `thickness` $\geq 0$ or fills the area bounded by the contours if `thickness` $< 0$. The example below shows how to retrieve connected components from the binary image and label them:

```cpp
#include "cv.h"
#include "highgui.h"

using namespace cv;

int main( int argc, char** argv )
{
    Mat src;
    // the first command-line parameter must be a filename of the binary
    // (black-n-white) image
    if( argc != 2 || !(src=imread(argv[1], 0)).data)
        return -1;

    Mat dst = Mat::zeros(src.rows, src.cols, CV_8UC3);

    src = src > 1;
    namedWindow( "Source", 1 );
    imshow( "Source", src );

    vector<vector<Point> > contours;
    vector<Vec4i> hierarchy;

    findContours( src, contours, hierarchy,
        CV_RETR_CCOMP, CV_CHAIN_APPROX_SIMPLE );

    // iterate through all the top-level contours,
    // draw each connected component with its own random color
    int idx = 0;
    for( ; idx >= 0; idx = hierarchy[idx][0] )
    {
        Scalar color( rand()&255, rand()&255, rand()&255 );
        drawContours( dst, contours, idx, color, CV_FILLED, 8, hierarchy );
    }

    namedWindow( "Components", 1 );
    imshow( "Components", dst );
    waitKey(0);
}
```

## approxPolyDP

Approximates a polygonal curve(s) with the specified precision.

**C++:** void **approxPolyDP**(InputArray **curve**, OutputArray **approxCurve**, double **epsilon**, bool **closed**)

**Python:** cv2.**approxPolyDP**(curve, epsilon, closed[, approxCurve]) → approxCurve

**C:** CvSeq* **cvApproxPoly**(const void* **src_seq**, int **header_size**, CvMemStorage* **storage**, int **method**, double **eps**, int **recursive**=0 )

>> **Parameters**

>>> **curve** – Input vector of a 2D point stored in:

---

- – `std::vector` or `Mat` (C++ interface)

- – `Nx2` numpy array (Python interface)

- – `CvSeq` or `` `CvMat` (C interface)

**approxCurve** – Result of the approximation. The type should match the type of the input curve. In case of C interface the approximated curve is stored in the memory storage and pointer to it is returned.

**epsilon** – Parameter specifying the approximation accuracy. This is the maximum distance between the original curve and its approximation.

**closed** – If true, the approximated curve is closed (its first and last vertices are connected). Otherwise, it is not closed.

**header_size** – Header size of the approximated curve. Normally, `sizeof(CvContour)` is used.

**storage** – Memory storage where the approximated curve is stored.

**method** – Contour approximation algorithm. Only `CV_POLY_APPROX_DP` is supported.

**recursive** – Recursion flag. If it is non-zero and `curve` is `CvSeq*`, the function `cvApproxPoly` approximates all the contours accessible from `curve` by `h_next` and `v_next` links.

The functions `approxPolyDP` approximate a curve or a polygon with another curve/polygon with less vertices so that the distance between them is less or equal to the specified precision. It uses the Douglas-Peucker algorithm http://en.wikipedia.org/wiki/Ramer-Douglas-Peucker_algorithm

See http://code.opencv.org/projects/opencv/repository/revisions/master/entry/samples/cpp/contours.cpp for the function usage model.

## ApproxChains

Approximates Freeman chain(s) with a polygonal curve.

**C:** CvSeq* **cvApproxChains**(CvSeq* **src_seq**, CvMemStorage* **storage**, int **method**=CV_CHAIN_APPROX_SIMPLE, double **parameter**=0, int **minimal_perimeter**=0, int **recursive**=0 )

**Python:** cv.**ApproxChains**(src_seq, storage, method=CV_CHAIN_APPROX_SIMPLE, parameter=0, minimal_perimeter=0, recursive=0) → contours

> **Parameters**

> > **src_seq** – Pointer to the approximated Freeman chain that can refer to other chains.

> > **storage** – Storage location for the resulting polylines.

> > **method** – Approximation method (see the description of the function `FindContours()` ).

> > **parameter** – Method parameter (not used now).

> > **minimal_perimeter** – Approximates only those contours whose perimeters are not less than `minimal_perimeter` . Other chains are removed from the resulting structure.

> > **recursive** – Recursion flag. If it is non-zero, the function approximates all chains that can be obtained from `chain` by using the `h_next` or `v_next` links. Otherwise, the single input chain is approximated.

This is a standalone contour approximation routine, not represented in the new interface. When `FindContours()` retrieves contours as Freeman chains, it calls the function to get approximated contours, represented as polygons.

## arcLength

Calculates a contour perimeter or a curve length.

**C++:** double **arcLength**(InputArray **curve**, bool **closed**)

**Python:** cv2.**arcLength**(curve, closed) → retval

**C:** double **cvArcLength**(const void* **curve**, CvSlice **slice**=CV_WHOLE_SEQ, int **is_closed**=-1 )

**Python:** cv.**ArcLength**(curve, slice=CV_WHOLE_SEQ, isClosed=-1) → float

> **Parameters**

>> **curve** – Input vector of 2D points, stored in `std::vector` or `Mat`.

>> **closed** – Flag indicating whether the curve is closed or not.

The function computes a curve length or a closed contour perimeter.

## boundingRect

Calculates the up-right bounding rectangle of a point set.

**C++:** Rect **boundingRect**(InputArray **points**)

**Python:** cv2.**boundingRect**(points) → retval

**C:** CvRect **cvBoundingRect**(CvArr* **points**, int **update**=0 )

**Python:** cv.**BoundingRect**(points, update=0) → CvRect

> **Parameters**

>> **points** – Input 2D point set, stored in `std::vector` or `Mat`.

The function calculates and returns the minimal up-right bounding rectangle for the specified point set.

## contourArea

Calculates a contour area.

**C++:** double **contourArea**(InputArray **contour**, bool **oriented**=false )

**Python:** cv2.**contourArea**(contour[, oriented]) → retval

**C:** double **cvContourArea**(const CvArr* **contour**, CvSlice **slice**=CV_WHOLE_SEQ, int **oriented**=0 )

**Python:** cv.**ContourArea**(contour, slice=CV_WHOLE_SEQ) → float

> **Parameters**

>> **contour** – Input vector of 2D points (contour vertices), stored in `std::vector` or `Mat`.

>> **oriented** – Oriented area flag. If it is true, the function returns a signed area value, depending on the contour orientation (clockwise or counter-clockwise). Using this feature you can determine orientation of a contour by taking the sign of an area. By default, the parameter is `false`, which means that the absolute value is returned.

The function computes a contour area. Similarly to `moments()`, the area is computed using the Green formula. Thus, the returned area and the number of non-zero pixels, if you draw the contour using `drawContours()` or `fillPoly()`, can be different. Also, the function will most certainly give a wrong results for contours with self-intersections.

Example:

```
vector<Point> contour;
contour.push_back(Point2f(0, 0));
contour.push_back(Point2f(10, 0));
contour.push_back(Point2f(10, 10));
contour.push_back(Point2f(5, 4));

double area0 = contourArea(contour);
vector<Point> approx;
approxPolyDP(contour, approx, 5, true);
double area1 = contourArea(approx);

cout << "area0 =" << area0 << endl <<
        "area1 =" << area1 << endl <<
        "approx poly vertices" << approx.size() << endl;
```

## convexHull

Finds the convex hull of a point set.

**C++:** void **convexHull**(InputArray **points**, OutputArray **hull**, bool **clockwise**=false, bool **returnPoints**=true )

**Python:** cv2.**convexHull**(points[, hull[, clockwise[, returnPoints ] ] ]) → hull

**C:** CvSeq* **cvConvexHull2**(const CvArr* **input**, void* **hull_storage**=NULL, int **orientation**=CV_CLOCKWISE, int **return_points**=0 )

**Python:** cv.**ConvexHull2**(points, storage, orientation=CV_CLOCKWISE, return_points=0) → convexHull

>   **Parameters**

>>   **points** – Input 2D point set, stored in std::vector or Mat.

>>   **hull** – Output convex hull. It is either an integer vector of indices or vector of points. In the first case, the hull elements are 0-based indices of the convex hull points in the original array (since the set of convex hull points is a subset of the original point set). In the second case, hull elements are the convex hull points themselves.

>>   **hull_storage** – Output memory storage in the old API (cvConvexHull2 returns a sequence containing the convex hull points or their indices).

>>   **clockwise** – Orientation flag. If it is true, the output convex hull is oriented clockwise. Otherwise, it is oriented counter-clockwise. The usual screen coordinate system is assumed so that the origin is at the top-left corner, x axis is oriented to the right, and y axis is oriented downwards.

>>   **orientation** – Convex hull orientation parameter in the old API, CV_CLOCKWISE or CV_COUNTERCLOCKWISE.

>>   **returnPoints** – Operation flag. In case of a matrix, when the flag is true, the function returns convex hull points. Otherwise, it returns indices of the convex hull points. When the output array is std::vector, the flag is ignored, and the output depends on the type of the vector: std::vector<int> implies returnPoints=true, std::vector<Point> implies returnPoints=false.

The functions find the convex hull of a 2D point set using the Sklansky's algorithm [Sklansky82] that has *O(N logN)* complexity in the current implementation. See the OpenCV sample convexhull.cpp that demonstrates the usage of different function variants.

## convexityDefects

Finds the convexity defects of a contour.

**C++:** void **convexityDefects**(InputArray **contour**, InputArray **convexhull**, OutputArray **convexityDefects**)

**Python:** cv2.**convexityDefects**(contour, convexhull[, convexityDefects]) → convexityDefects

**C:** CvSeq* **cvConvexityDefects**(const CvArr* **contour**, const CvArr* **convexhull**, CvMemStorage* **storage**=NULL )

**Python:** cv.**ConvexityDefects**(contour, convexhull, storage) → convexityDefects

> **Parameters**
>
> > **contour** – Input contour.
> >
> > **convexhull** – Convex hull obtained using convexHull() that should contain indices of the contour points that make the hull.
> >
> > **convexityDefects** – The output vector of convexity defects. In C++ and the new Python/Java interface each convexity defect is represented as 4-element integer vector (a.k.a. cv::Vec4i): (start_index, end_index, farthest_pt_index, fixpt_depth), where indices are 0-based indices in the original contour of the convexity defect beginning, end and the farthest point, and fixpt_depth is fixed-point approximation (with 8 fractional bits) of the distance between the farthest contour point and the hull. That is, to get the floating-point value of the depth will be fixpt_depth/256.0. In C interface convexity defect is represented by CvConvexityDefect structure - see below.
> >
> > **storage** – Container for the output sequence of convexity defects. If it is NULL, the contour or hull (in that order) storage is used.
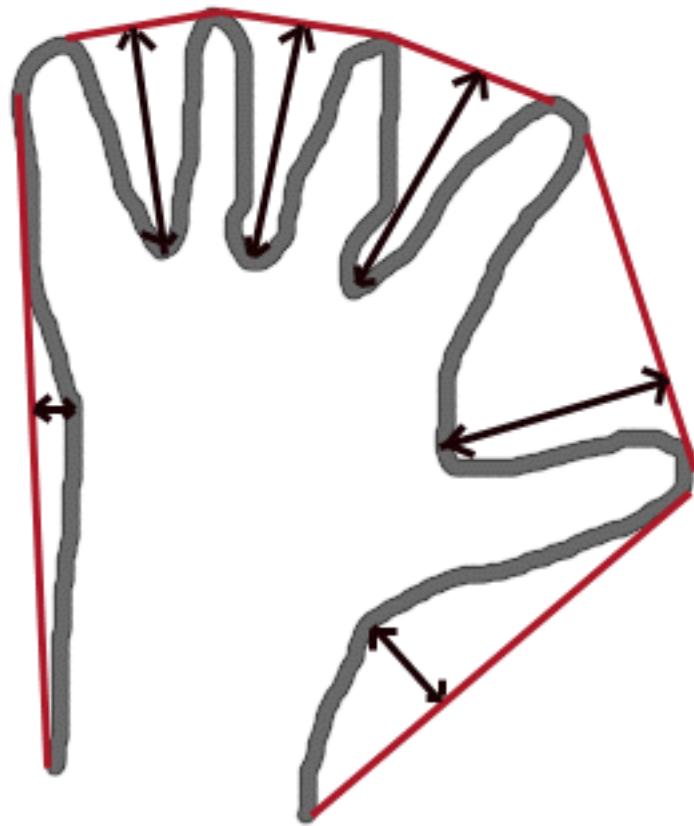
The function finds all convexity defects of the input contour and returns a sequence of the CvConvexityDefect structures, where CvConvexityDetect is defined as:

```
struct CvConvexityDefect
{
    CvPoint* start; // point of the contour where the defect begins
    CvPoint* end; // point of the contour where the defect ends
    CvPoint* depth_point; // the farthest from the convex hull point within the defect
    float depth; // distance between the farthest point and the convex hull
};
```

The figure below displays convexity defects of a hand contour:

## fitEllipse

Fits an ellipse around a set of 2D points.

**C++:** RotatedRect **fitEllipse**(InputArray **points**)

**Python:** cv2.**fitEllipse**(points) → retval

**C:** CvBox2D **cvFitEllipse2**(const CvArr* **points**)

**Python:** cv.**FitEllipse2**(points) → Box2D

> **Parameters**
>
> > **points** – Input 2D point set, stored in:
> >
> > – std::vector<> or Mat (C++ interface)
> >
> > – CvSeq* or CvMat* (C interface)
> >
> > – Nx2 numpy array (Python interface)

The function calculates the ellipse that fits (in a least-squares sense) a set of 2D points best of all. It returns the rotated rectangle in which the ellipse is inscribed. The algorithm [Fitzgibbon95] is used.

## fitLine

Fits a line to a 2D or 3D point set.

**C++:** void **fitLine**(InputArray **points**, OutputArray **line**, int **distType**, double **param**, double **reps**, double **aeps**)

**Python:** cv2.**fitLine**(points, distType, param, reps, aeps[, line ]) → line

**C:** void **cvFitLine**(const CvArr* **points**, int **dist_type**, double **param**, double **reps**, double **aeps**, float* **line**)

**Python:** cv.**FitLine**(points, dist_type, param, reps, aeps) → line

> **Parameters**

>> **points** – Input vector of 2D or 3D points, stored in std::vector<> or Mat.

>> **line** – Output line parameters. In case of 2D fitting, it should be a vector of 4 elements (like Vec4f) - (vx, vy, x0, y0), where (vx, vy) is a normalized vector collinear to the line and (x0, y0) is a point on the line. In case of 3D fitting, it should be a vector of 6 elements (like Vec6f) - (vx, vy, vz, x0, y0, z0), where (vx, vy, vz) is a normalized vector collinear to the line and (x0, y0, z0) is a point on the line.

>> **distType** – Distance used by the M-estimator (see the discussion below).

>> **param** – Numerical parameter ( C ) for some types of distances. If it is 0, an optimal value is chosen.

>> **reps** – Sufficient accuracy for the radius (distance between the coordinate origin and the line).

>> **aeps** – Sufficient accuracy for the angle. 0.01 would be a good default value for reps and aeps.

The function fitLine fits a line to a 2D or 3D point set by minimizing $\sum_i \rho(r_i)$ where $r_i$ is a distance between the $i^{th}$ point, the line and $\rho(r)$ is a distance function, one of the following:

- distType=CV_DIST_L2

$$\rho(r) = r^2/2 \quad \text{(the simplest and the fastest least-squares method)}$$

- distType=CV_DIST_L1

$$\rho(r) = r$$

- distType=CV_DIST_L12

$$\rho(r) = 2 \cdot (\sqrt{1 + \frac{r^2}{2}} - 1)$$

- distType=CV_DIST_FAIR

$$\rho(r) = C^2 \cdot \left( \frac{r}{C} - \log\left(1 + \frac{r}{C}\right) \right) \quad \text{where} \quad C = 1.3998$$

- distType=CV_DIST_WELSCH

$$\rho(r) = \frac{C^2}{2} \cdot \left( 1 - \exp\left( -\left(\frac{r}{C}\right)^2 \right) \right) \quad \text{where} \quad C = 2.9846$$

- distType=CV_DIST_HUBER

$$\rho(r) = \begin{cases} r^2/2 & \text{if } r < C \\ C \cdot (r - C/2) & \text{otherwise} \end{cases} \quad \text{where} \quad C = 1.345$$

The algorithm is based on the M-estimator ( http://en.wikipedia.org/wiki/M-estimator ) technique that iteratively fits the line using the weighted least-squares algorithm. After each iteration the weights $w_i$ are adjusted to be inversely proportional to $\rho(r_i)$ .

## isContourConvex

Tests a contour convexity.

**C++:** bool **isContourConvex**(InputArray **contour**)

**Python:** cv2.**isContourConvex**(contour) → retval

**C:** int **cvCheckContourConvexity**(const CvArr* **contour**)

**Python:** cv.**CheckContourConvexity**(contour) → int

> **Parameters**
>
> > **contour** – Input vector of 2D points, stored in:
> >
> > – `std::vector<>` or `Mat` (C++ interface)
> >
> > – `CvSeq*` or `CvMat*` (C interface)
> >
> > – Nx2 numpy array (Python interface)

The function tests whether the input contour is convex or not. The contour must be simple, that is, without self-intersections. Otherwise, the function output is undefined.

## minAreaRect

Finds a rotated rectangle of the minimum area enclosing the input 2D point set.

**C++:** RotatedRect **minAreaRect**(InputArray **points**)

**Python:** cv2.**minAreaRect**(points) → retval

**C:** CvBox2D **cvMinAreaRect2**(const CvArr* **points**, CvMemStorage* **storage**=NULL )

**Python:** cv.**MinAreaRect2**(points, storage=None) → Box2D

> **Parameters**
>
> > **points** – Input vector of 2D points, stored in:
> >
> > – `std::vector<>` or `Mat` (C++ interface)
> >
> > – `CvSeq*` or `CvMat*` (C interface)
> >
> > – Nx2 numpy array (Python interface)

The function calculates and returns the minimum-area bounding rectangle (possibly rotated) for a specified point set. See the OpenCV sample `minarea.cpp` .

## minEnclosingCircle

Finds a circle of the minimum area enclosing a 2D point set.

**C++:** void **minEnclosingCircle**(InputArray **points**, Point2f& **center**, float& **radius**)

**Python:** cv2.**minEnclosingCircle**(points) → center, radius

**C:** int **cvMinEnclosingCircle**(const CvArr* **points**, CvPoint2D32f* **center**, float* **radius**)

**Python:** cv.**MinEnclosingCircle**(points)-> (int, center, radius)

> **Parameters**
>
> > **points** – Input vector of 2D points, stored in:
> >
> > – std::vector<> or Mat (C++ interface)
> >
> > – CvSeq* or CvMat* (C interface)
> >
> > – Nx2 numpy array (Python interface)
> >
> > **center** – Output center of the circle.
> >
> > **radius** – Output radius of the circle.

The function finds the minimal enclosing circle of a 2D point set using an iterative algorithm. See the OpenCV sample minarea.cpp .

## matchShapes

Compares two shapes.

**C++:** double **matchShapes**(InputArray **contour1**, InputArray **contour2**, int **method**, double **parameter**)

**Python:** cv2.**matchShapes**(contour1, contour2, method, parameter) → retval

**C:** double **cvMatchShapes**(const void* **object1**, const void* **object2**, int **method**, double **parameter**=0 )

**Python:** cv.**MatchShapes**(object1, object2, method, parameter=0) → float

> **Parameters**
>
> > **object1** – First contour or grayscale image.
> >
> > **object2** – Second contour or grayscale image.
> >
> > **method** – Comparison method: CV_CONTOURS_MATCH_I1 , CV_CONTOURS_MATCH_I2 or CV_CONTOURS_MATCH_I3 (see the details below).
> >
> > **parameter** – Method-specific parameter (not supported now).

The function compares two shapes. All three implemented methods use the Hu invariants (see HuMoments() ) as follows ( $A$ denotes object1,:math:$B$ denotes object2 ):

- method=CV_CONTOURS_MATCH_I1

$$I_1(A, B) = \sum_{i=1\ldots7} \left| \frac{1}{m_i^A} - \frac{1}{m_i^B} \right|$$

- method=CV_CONTOURS_MATCH_I2

$$I_2(A, B) = \sum_{i=1...7} \left| m_i^A - m_i^B \right|$$

- method=CV_CONTOURS_MATCH_I3

$$I_3(A, B) = \max_{i=1...7} \frac{\left| m_i^A - m_i^B \right|}{\left| m_i^A \right|}$$

where

$$m_i^A = \text{sign}(h_i^A) \cdot \log h_i^A$$
$$m_i^B = \text{sign}(h_i^B) \cdot \log h_i^B$$

and $h_i^A, h_i^B$ are the Hu moments of A and B , respectively.

## pointPolygonTest

Performs a point-in-contour test.

**C++:** double **pointPolygonTest**(InputArray **contour**, Point2f **pt**, bool **measureDist**)

**Python:** cv2.**pointPolygonTest**(contour, pt, measureDist) → retval

**C:** double **cvPointPolygonTest**(const CvArr* **contour**, CvPoint2D32f **pt**, int **measure_dist**)

**Python:** cv.**PointPolygonTest**(contour, pt, measure_dist) → float
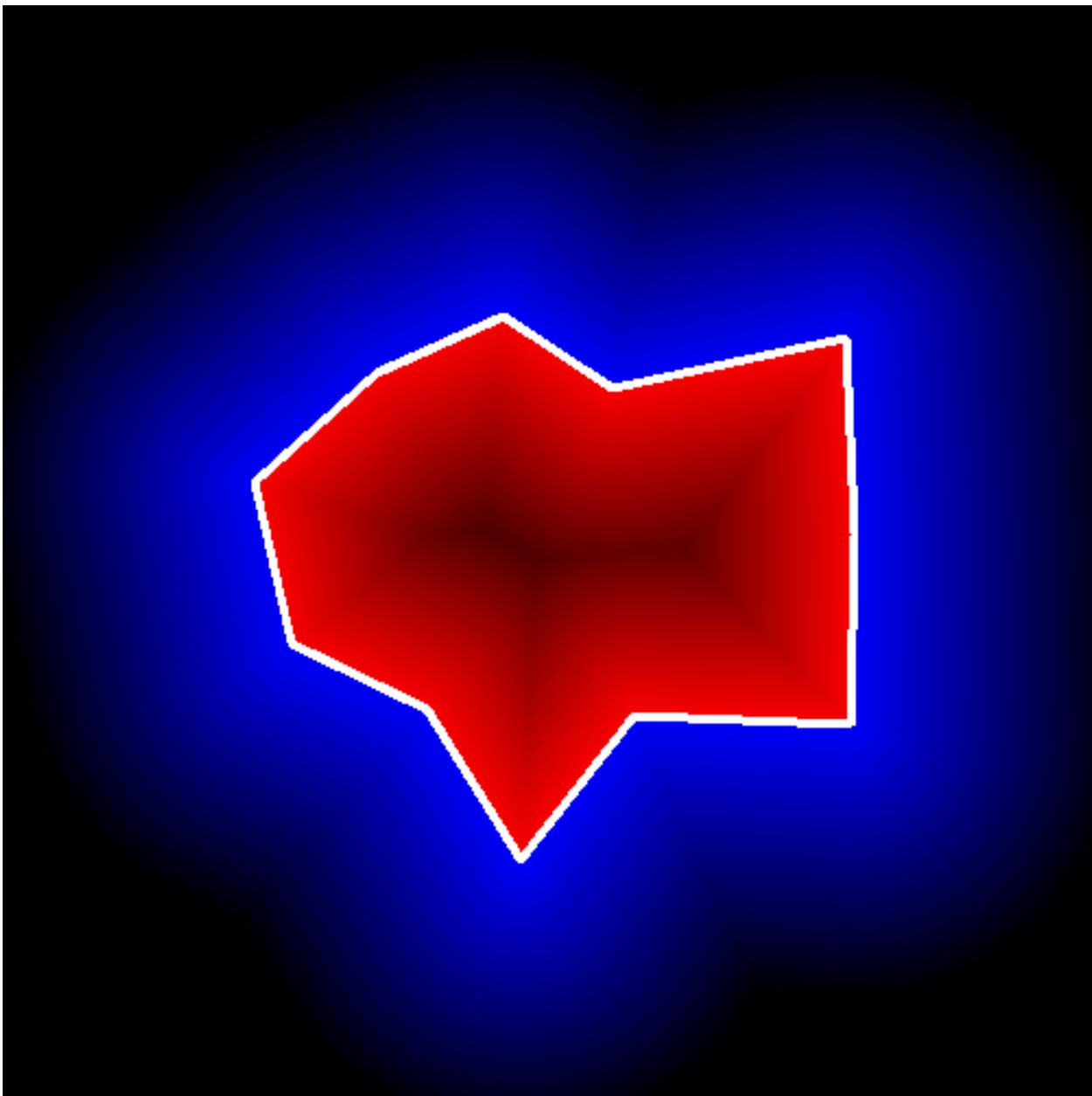
> **Parameters**
>
> > **contour** – Input contour.
> >
> > **pt** – Point tested against the contour.
> >
> > **measureDist** – If true, the function estimates the signed distance from the point to the nearest contour edge. Otherwise, the function only checks if the point is inside a contour or not.

The function determines whether the point is inside a contour, outside, or lies on an edge (or coincides with a vertex). It returns positive (inside), negative (outside), or zero (on an edge) value, correspondingly. When `measureDist=false` , the return value is +1, -1, and 0, respectively. Otherwise, the return value is a signed distance between the point and the nearest contour edge.

See below a sample output of the function where each image pixel is tested against the contour.

## 3.6 Motion Analysis and Object Tracking

### accumulate

Adds an image to the accumulator.

**C++:** void **accumulate**(InputArray **src**, InputOutputArray **dst**, InputArray **mask**=noArray() )

**Python:** cv2.**accumulate**(src, dst[, mask ]) → None

**C:** void **cvAcc**(const CvArr* **image**, CvArr* **sum**, const CvArr* **mask**=NULL )

**Python:** cv.**Acc**(image, sum, mask=None) → None

> **Parameters**

>> **src** – Input image as 1- or 3-channel, 8-bit or 32-bit floating point.

>> **dst** – Accumulator image with the same number of channels as input image, 32-bit or 64-bit floating-point.

>> **mask** – Optional operation mask.

The function adds `src` or some of its elements to `dst` :

$$\mathtt{dst}(x,y) \leftarrow \mathtt{dst}(x,y) + \mathtt{src}(x,y) \quad \text{if} \quad \mathtt{mask}(x,y) \neq 0$$

The function supports multi-channel images. Each channel is processed independently.

The functions `accumulate*` can be used, for example, to collect statistics of a scene background viewed by a still camera and for the further foreground-background segmentation.

**See Also:**

`accumulateSquare()`, `accumulateProduct()`, `accumulateWeighted()`

## accumulateSquare

Adds the square of a source image to the accumulator.

**C++:** void **accumulateSquare**(InputArray **src**, InputOutputArray **dst**, InputArray **mask**=noArray() )

**Python:** cv2.**accumulateSquare**(src, dst[, mask ]) → None

**C:** void **cvSquareAcc**(const CvArr* **image**, CvArr* **sqsum**, const CvArr* **mask**=NULL )

**Python:** cv.**SquareAcc**(image, sqsum, mask=None) → None

> **Parameters**

>> **src** – Input image as 1- or 3-channel, 8-bit or 32-bit floating point.

>> **dst** – Accumulator image with the same number of channels as input image, 32-bit or 64-bit floating-point.

>> **mask** – Optional operation mask.

The function adds the input image `src` or its selected region, raised to a power of 2, to the accumulator `dst` :

$$\mathtt{dst}(x,y) \leftarrow \mathtt{dst}(x,y) + \mathtt{src}(x,y)^2 \quad \text{if} \quad \mathtt{mask}(x,y) \neq 0$$

The function supports multi-channel images. Each channel is processed independently.

**See Also:**

`accumulateSquare()`, `accumulateProduct()`, `accumulateWeighted()`

## accumulateProduct

Adds the per-element product of two input images to the accumulator.

**C++:** void **accumulateProduct**(InputArray **src1**, InputArray **src2**, InputOutputArray **dst**, InputArray **mask**=noArray() )

**Python:** cv2.**accumulateProduct**(src1, src2, dst[, mask ]) → None

**C:** void **cvMultiplyAcc**(const CvArr* **image1**, const CvArr* **image2**, CvArr* **acc**, const CvArr* **mask**=NULL )

---

**Python:** cv.**MultiplyAcc**(image1, image2, acc, mask=None) → None

       **Parameters**

              **src1** – First input image, 1- or 3-channel, 8-bit or 32-bit floating point.

              **src2** – Second input image of the same type and the same size as src1 .

              **dst** – Accumulator with the same number of channels as input images, 32-bit or 64-bit floating-point.

              **mask** – Optional operation mask.

The function adds the product of two images or their selected regions to the accumulator dst :

$$dst(x, y) \leftarrow dst(x, y) + src1(x, y) \cdot src2(x, y) \quad \text{if} \quad mask(x, y) \neq 0$$

The function supports multi-channel images. Each channel is processed independently.

**See Also:**

accumulate(), accumulateSquare(), accumulateWeighted()

## accumulateWeighted

Updates a running average.

**C++:** void **accumulateWeighted**(InputArray **src**, InputOutputArray **dst**, double **alpha**, InputArray **mask**=noArray() )

**Python:** cv2.**accumulateWeighted**(src, dst, alpha[, mask]) → None

**C:** void **cvRunningAvg**(const CvArr* **image**, CvArr* **acc**, double **alpha**, const CvArr* **mask**=NULL )

**Python:** cv.**RunningAvg**(image, acc, alpha, mask=None) → None

       **Parameters**

              **src** – Input image as 1- or 3-channel, 8-bit or 32-bit floating point.

              **dst** – Accumulator image with the same number of channels as input image, 32-bit or 64-bit floating-point.

              **alpha** – Weight of the input image.

              **mask** – Optional operation mask.

The function calculates the weighted sum of the input image src and the accumulator dst so that dst becomes a running average of a frame sequence:

$$dst(x, y) \leftarrow (1 - alpha) \cdot dst(x, y) + alpha \cdot src(x, y) \quad \text{if} \quad mask(x, y) \neq 0$$

That is, alpha regulates the update speed (how fast the accumulator "forgets" about earlier images). The function supports multi-channel images. Each channel is processed independently.

**See Also:**

accumulate(), accumulateSquare(), accumulateProduct()

# phaseCorrelate

The function is used to detect translational shifts that occur between two images. The operation takes advantage of the Fourier shift theorem for detecting the translational shift in the frequency domain. It can be used for fast image registration as well as motion estimation. For more information please see http://en.wikipedia.org/wiki/Phase_correlation .

Calculates the cross-power spectrum of two supplied source arrays. The arrays are padded if needed with `getOptimalDFTSize()`.

**C++:** Point2d **phaseCorrelate**(InputArray **src1**, InputArray **src2**, InputArray **window**=noArray())

**C++:** Point2d **phaseCorrelateRes**(InputArray **src1**, InputArray **src2**, InputArray **window**, double* **response**=0)

>       **Parameters**

>>          **src1** – Source floating point array (CV_32FC1 or CV_64FC1)

>>          **src2** – Source floating point array (CV_32FC1 or CV_64FC1)

>>          **window** – Floating point array with windowing coefficients to reduce edge effects (optional).

>>          **response** – Signal power within the 5x5 centroid around the peak, between 0 and 1 (optional).

Return value: detected phase shift (sub-pixel) between the two arrays.

The function performs the following equations

- First it applies a Hanning window (see http://en.wikipedia.org/wiki/Hann_function) to each image to remove possible edge effects. This window is cached until the array size changes to speed up processing time.

- Next it computes the forward DFTs of each source array:

$$\mathbf{G}_a = \mathcal{F}\{src_1\}, \ \mathbf{G}_b = \mathcal{F}\{src_2\}$$

  where $\mathcal{F}$ is the forward DFT.

- It then computes the cross-power spectrum of each frequency domain array:

$$R = \frac{\mathbf{G}_a \mathbf{G}_b^*}{|\mathbf{G}_a \mathbf{G}_b^*|}$$

- Next the cross-correlation is converted back into the time domain via the inverse DFT:

$$r = \mathcal{F}^{-1}\{R\}$$

- Finally, it computes the peak location and computes a 5x5 weighted centroid around the peak to achieve sub-pixel accuracy.

$$(\Delta x, \Delta y) = \texttt{weightedCentroid}\{\arg \max_{(x,y)}\{r\}\}$$

- If non-zero, the response parameter is computed as the sum of the elements of r within the 5x5 centroid around the peak location. It is normalized to a maximum of 1 (meaning there is a single peak) and will be smaller when there are multiple peaks.

**See Also:**

`dft()`, `getOptimalDFTSize()`, `idft()`, `mulSpectrums() createHanningWindow()`

## createHanningWindow

This function computes a Hanning window coefficients in two dimensions. See http://en.wikipedia.org/wiki/Hann_function and http://en.wikipedia.org/wiki/Window_function for more information.

**C++:** void **createHanningWindow**(OutputArray **dst**, Size **winSize**, int **type**)

> **Parameters**
>
> > **dst** – Destination array to place Hann coefficients in
> >
> > **winSize** – The window size specifications
> >
> > **type** – Created array type

An example is shown below:

```
// create hanning window of size 100x100 and type CV_32F
Mat hann;
createHanningWindow(hann, Size(100, 100), CV_32F);
```

**See Also:**

`phaseCorrelate()`

# 3.7 Feature Detection

## Canny

Finds edges in an image using the [Canny86] algorithm.

**C++:** void **Canny**(InputArray **image**, OutputArray **edges**, double **threshold1**, double **threshold2**, int **apertureSize**=3, bool **L2gradient**=false )

**Python:** cv2.**Canny**(image, threshold1, threshold2$\big[$, edges$\big[$, apertureSize$\big[$, L2gradient $\big]\,\big]\,\big]$) $\rightarrow$ edges

**C:** void **cvCanny**(const CvArr* **image**, CvArr* **edges**, double **threshold1**, double **threshold2**, int **aperture_size**=3 )

**Python:** cv.**Canny**(image, edges, threshold1, threshold2, aperture_size=3) $\rightarrow$ None

> **Parameters**
>
> > **image** – single-channel 8-bit input image.
> >
> > **edges** – output edge map; it has the same size and type as `image` .
> >
> > **threshold1** – first threshold for the hysteresis procedure.
> >
> > **threshold2** – second threshold for the hysteresis procedure.
> >
> > **apertureSize** – aperture size for the `Sobel()` operator.

**L2gradient** – a flag, indicating whether a more accurate $L_2$ norm $= \sqrt{(dI/dx)^2 + (dI/dy)^2}$ should be used to calculate the image gradient magnitude ( L2gradient=true ), or whether the default $L_1$ norm $= |dI/dx| + |dI/dy|$ is enough ( L2gradient=false ).

The function finds edges in the input image image and marks them in the output map edges using the Canny algorithm. The smallest value between threshold1 and threshold2 is used for edge linking. The largest value is used to find initial segments of strong edges. See http://en.wikipedia.org/wiki/Canny_edge_detector

## cornerEigenValsAndVecs

Calculates eigenvalues and eigenvectors of image blocks for corner detection.

**C++:** void **cornerEigenValsAndVecs**(InputArray **src**, OutputArray **dst**, int **blockSize**, int **ksize**, int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**cornerEigenValsAndVecs**(src, blockSize, ksize[, dst[, borderType ]]) → dst

**C:** void **cvCornerEigenValsAndVecs**(const CvArr* **image**, CvArr* **eigenvv**, int **block_size**, int **aperture_size**=3 )

**Python:** cv.**CornerEigenValsAndVecs**(image, eigenvv, blockSize, aperture_size=3) → None

> **Parameters**
>
> > **src** – Input single-channel 8-bit or floating-point image.
> >
> > **dst** – Image to store the results. It has the same size as src and the type CV_32FC(6) .
> >
> > **blockSize** – Neighborhood size (see details below).
> >
> > **ksize** – Aperture parameter for the Sobel() operator.
> >
> > **borderType** – Pixel extrapolation method. See borderInterpolate() .

For every pixel p , the function cornerEigenValsAndVecs considers a blockSize $\times$ blockSize neighborhood $S(p)$ . It calculates the covariation matrix of derivatives over the neighborhood as:

$$M = \begin{bmatrix} \sum_{S(p)} (dI/dx)^2 & \sum_{S(p)} (dI/dx dI/dy)^2 \\ \sum_{S(p)} (dI/dx dI/dy)^2 & \sum_{S(p)} (dI/dy)^2 \end{bmatrix}$$

where the derivatives are computed using the Sobel() operator.

After that, it finds eigenvectors and eigenvalues of $M$ and stores them in the destination image as $(\lambda_1, \lambda_2, x_1, y_1, x_2, y_2)$ where

- $\lambda_1, \lambda_2$ are the non-sorted eigenvalues of $M$
- $x_1, y_1$ are the eigenvectors corresponding to $\lambda_1$
- $x_2, y_2$ are the eigenvectors corresponding to $\lambda_2$

The output of the function can be used for robust edge or corner detection.

**See Also:**

cornerMinEigenVal(), cornerHarris(), preCornerDetect()

## cornerHarris

Harris edge detector.

**C++:** void **cornerHarris**(InputArray **src**, OutputArray **dst**, int **blockSize**, int **ksize**, double **k**, int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**cornerHarris**(src, blockSize, ksize, k[, dst[, borderType ] ]) → dst

**C:** void **cvCornerHarris**(const CvArr* **image**, CvArr* **harris_responce**, int **block_size**, int **aperture_size**=3, double **k**=0.04 )

**Python:** cv.**CornerHarris**(image, harris_dst, blockSize, aperture_size=3, k=0.04) → None

> **Parameters**
>
> > **src** – Input single-channel 8-bit or floating-point image.
> >
> > **dst** – Image to store the Harris detector responses. It has the type CV_32FC1 and the same size as src .
> >
> > **blockSize** – Neighborhood size (see the details on cornerEigenValsAndVecs() ).
> >
> > **ksize** – Aperture parameter for the Sobel() operator.
> >
> > **k** – Harris detector free parameter. See the formula below.
> >
> > **borderType** – Pixel extrapolation method. See borderInterpolate() .

The function runs the Harris edge detector on the image. Similarly to cornerMinEigenVal() and cornerEigenValsAndVecs() , for each pixel $(x, y)$ it calculates a $2 \times 2$ gradient covariance matrix $M^{(x,y)}$ over a blockSize × blockSize neighborhood. Then, it computes the following characteristic:

$$\texttt{dst}(x, y) = \det M^{(x,y)} - k \cdot \left( \operatorname{tr} M^{(x,y)} \right)^2$$

Corners in the image can be found as the local maxima of this response map.

## cornerMinEigenVal

Calculates the minimal eigenvalue of gradient matrices for corner detection.

**C++:** void **cornerMinEigenVal**(InputArray **src**, OutputArray **dst**, int **blockSize**, int **ksize**=3, int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**cornerMinEigenVal**(src, blockSize[, dst[, ksize[, borderType ] ] ]) → dst

**C:** void **cvCornerMinEigenVal**(const CvArr* **image**, CvArr* **eigenval**, int **block_size**, int **aperture_size**=3 )

**Python:** cv.**CornerMinEigenVal**(image, eigenval, blockSize, aperture_size=3) → None

> **Parameters**
>
> > **src** – Input single-channel 8-bit or floating-point image.
> >
> > **dst** – Image to store the minimal eigenvalues. It has the type CV_32FC1 and the same size as src .
> >
> > **blockSize** – Neighborhood size (see the details on cornerEigenValsAndVecs() ).
> >
> > **ksize** – Aperture parameter for the Sobel() operator.
> >
> > **borderType** – Pixel extrapolation method. See borderInterpolate() .

The function is similar to cornerEigenValsAndVecs() but it calculates and stores only the minimal eigenvalue of the covariance matrix of derivatives, that is, $\min(\lambda_1, \lambda_2)$ in terms of the formulae in the cornerEigenValsAndVecs() description.

## cornerSubPix

Refines the corner locations.

**C++:** void **cornerSubPix**(InputArray **image**, InputOutputArray **corners**, Size **winSize**, Size **zeroZone**, TermCriteria **criteria**)

**Python:** cv2.**cornerSubPix**(image, corners, winSize, zeroZone, criteria) → None

**C:** void **cvFindCornerSubPix**(const CvArr* **image**, CvPoint2D32f* **corners**, int **count**, CvSize **win**, CvSize **zero_zone**, CvTermCriteria **criteria**)

**Python:** cv.**FindCornerSubPix**(image, corners, win, zero_zone, criteria) → corners
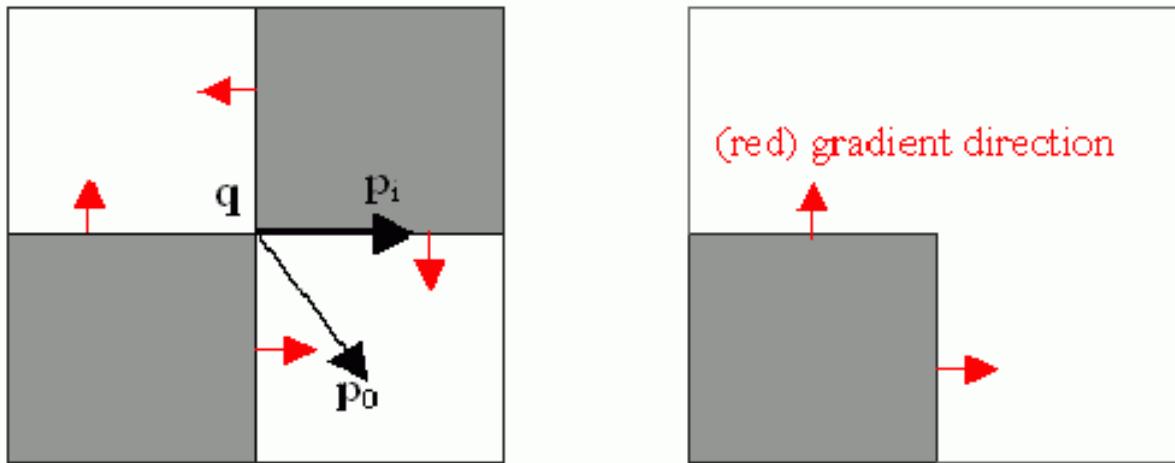
> **Parameters**
>
>> **image** – Input image.
>>
>> **corners** – Initial coordinates of the input corners and refined coordinates provided for output.
>>
>> **winSize** – Half of the side length of the search window. For example, if winSize=Size(5,5), then a $5*2+1 \times 5*2+1 = 11 \times 11$ search window is used.
>>
>> **zeroZone** – Half of the size of the dead region in the middle of the search zone over which the summation in the formula below is not done. It is used sometimes to avoid possible singularities of the autocorrelation matrix. The value of (-1,-1) indicates that there is no such a size.
>>
>> **criteria** – Criteria for termination of the iterative process of corner refinement. That is, the process of corner position refinement stops either after criteria.maxCount iterations or when the corner position moves by less than criteria.epsilon on some iteration.

The function iterates to find the sub-pixel accurate location of corners or radial saddle points, as shown on the figure below.



Sub-pixel accurate corner locator is based on the observation that every vector from the center $q$ to a point $p$ located within a neighborhood of $q$ is orthogonal to the image gradient at $p$ subject to image and measurement noise. Consider the expression:

$$\epsilon_i = DI_{p_i}^T \cdot (q - p_i)$$

where $DI_{p_i}$ is an image gradient at one of the points $p_i$ in a neighborhood of $q$. The value of $q$ is to be found so that $\epsilon_i$ is minimized. A system of equations may be set up with $\epsilon_i$ set to zero:

$$\sum_i (DI_{p_i} \cdot DI_{p_i}{}^T) - \sum_i (DI_{p_i} \cdot DI_{p_i}{}^T \cdot p_i)$$

where the gradients are summed within a neighborhood ("search window") of $q$. Calling the first gradient term G and the second gradient term b gives:

$$q = G^{-1} \cdot b$$

The algorithm sets the center of the neighborhood window at this new center $q$ and then iterates until the center stays within a set threshold.

## goodFeaturesToTrack

Determines strong corners on an image.

**C++:** void **goodFeaturesToTrack**(InputArray **image**, OutputArray **corners**, int **maxCorners**, double **qualityLevel**, double **minDistance**, InputArray **mask**=noArray(), int **blockSize**=3, bool **useHarrisDetector**=false, double **k**=0.04 )

**Python:** cv2.**goodFeaturesToTrack**(image, maxCorners, qualityLevel, minDistance[, corners[, mask[, blockSize[, useHarrisDetector[, k ]]]]]) → corners

**C:** void **cvGoodFeaturesToTrack**(const CvArr* **image**, CvArr* **eig_image**, CvArr* **temp_image**, CvPoint2D32f* **corners**, int* **corner_count**, double **quality_level**, double **min_distance**, const CvArr* **mask**=NULL, int **block_size**=3, int **use_harris**=0, double **k**=0.04 )

**Python:** cv.**GoodFeaturesToTrack**(image, eigImage, tempImage, cornerCount, qualityLevel, minDistance, mask=None, blockSize=3, useHarris=0, k=0.04) → cornerCount

> **Parameters**
>
> > **image** – Input 8-bit or floating-point 32-bit, single-channel image.
> >
> > **eig_image** – The parameter is ignored.
> >
> > **temp_image** – The parameter is ignored.
> >
> > **corners** – Output vector of detected corners.
> >
> > **maxCorners** – Maximum number of corners to return. If there are more corners than are found, the strongest of them is returned.
> >
> > **qualityLevel** – Parameter characterizing the minimal accepted quality of image corners. The parameter value is multiplied by the best corner quality measure, which is the minimal eigenvalue (see `cornerMinEigenVal()` ) or the Harris function response (see `cornerHarris()` ). The corners with the quality measure less than the product are rejected. For example, if the best corner has the quality measure = 1500, and the `qualityLevel=0.01` , then all the corners with the quality measure less than 15 are rejected.
> >
> > **minDistance** – Minimum possible Euclidean distance between the returned corners.
> >
> > **mask** – Optional region of interest. If the image is not empty (it needs to have the type `CV_8UC1` and the same size as `image` ), it specifies the region in which the corners are detected.
> >
> > **blockSize** – Size of an average block for computing a derivative covariation matrix over each pixel neighborhood. See `cornerEigenValsAndVecs()` .

**useHarrisDetector** – Parameter indicating whether to use a Harris detector (see `cornerHarris()`) or `cornerMinEigenVal()`.

**k** – Free parameter of the Harris detector.

The function finds the most prominent corners in the image or in the specified image region, as described in [Shi94]:

1. Function calculates the corner quality measure at every source image pixel using the `cornerMinEigenVal()` or `cornerHarris()`.

2. Function performs a non-maximum suppression (the local maximums in *3 x 3* neighborhood are retained).

3. The corners with the minimal eigenvalue less than `qualityLevel` $\cdot \max_{x,y} qualityMeasureMap(x,y)$ are rejected.

4. The remaining corners are sorted by the quality measure in the descending order.

5. Function throws away each corner for which there is a stronger corner at a distance less than `maxDistance`.

The function can be used to initialize a point-based tracker of an object.

---

**Note:** If the function is called with different values A and B of the parameter `qualityLevel`, and A > {B}, the vector of returned corners with `qualityLevel=A` will be the prefix of the output vector with `qualityLevel=B`.

---

**See Also:**

`cornerMinEigenVal()`, `cornerHarris()`, `calcOpticalFlowPyrLK()`, `estimateRigidTransform()`,

## HoughCircles

Finds circles in a grayscale image using the Hough transform.

**C++:** void **HoughCircles**(InputArray **image**, OutputArray **circles**, int **method**, double **dp**, double **minDist**, double **param1**=100, double **param2**=100, int **minRadius**=0, int **maxRadius**=0 )

**C:** CvSeq* **cvHoughCircles**(CvArr* **image**, void* **circle_storage**, int **method**, double **dp**, double **min_dist**, double **param1**=100, double **param2**=100, int **min_radius**=0, int **max_radius**=0 )

**Python:** cv2.**HoughCircles**(image, method, dp, minDist[, circles[, param1[, param2[, minRadius[, maxRadius ]]]]]) → circles

    **Parameters**

        **image** – 8-bit, single-channel, grayscale input image.

        **circles** – Output vector of found circles. Each vector is encoded as a 3-element floating-point vector $(x, y, radius)$.

        **circle_storage** – In C function this is a memory storage that will contain the output sequence of found circles.

        **method** – Detection method to use. Currently, the only implemented method is `CV_HOUGH_GRADIENT`, which is basically *21HT*, described in [Yuen90].

        **dp** – Inverse ratio of the accumulator resolution to the image resolution. For example, if `dp=1`, the accumulator has the same resolution as the input image. If `dp=2`, the accumulator has half as big width and height.

        **minDist** – Minimum distance between the centers of the detected circles. If the parameter is too small, multiple neighbor circles may be falsely detected in addition to a true one. If it is too large, some circles may be missed.

> > > **param1** – First method-specific parameter. In case of `CV_HOUGH_GRADIENT` , it is the higher
> > > threshold of the two passed to the `Canny()` edge detector (the lower one is twice smaller).
> > >
> > > **param2** – Second method-specific parameter. In case of `CV_HOUGH_GRADIENT` , it is the
> > > accumulator threshold for the circle centers at the detection stage. The smaller it is, the
> > > more false circles may be detected. Circles, corresponding to the larger accumulator values,
> > > will be returned first.
> > >
> > > **minRadius** – Minimum circle radius.
> > >
> > > **maxRadius** – Maximum circle radius.

The function finds circles in a grayscale image using a modification of the Hough transform.

Example:

```cpp
#include <cv.h>
#include <highgui.h>
#include <math.h>

using namespace cv;

int main(int argc, char** argv)
{
    Mat img, gray;
    if( argc != 2 && !(img=imread(argv[1], 1)).data)
        return -1;
    cvtColor(img, gray, CV_BGR2GRAY);
    // smooth it, otherwise a lot of false circles may be detected
    GaussianBlur( gray, gray, Size(9, 9), 2, 2 );
    vector<Vec3f> circles;
    HoughCircles(gray, circles, CV_HOUGH_GRADIENT,
                 2, gray->rows/4, 200, 100 );
    for( size_t i = 0; i < circles.size(); i++ )
    {
        Point center(cvRound(circles[i][0]), cvRound(circles[i][1]));
        int radius = cvRound(circles[i][2]);
        // draw the circle center
        circle( img, center, 3, Scalar(0,255,0), -1, 8, 0 );
        // draw the circle outline
        circle( img, center, radius, Scalar(0,0,255), 3, 8, 0 );
    }
    namedWindow( "circles", 1 );
    imshow( "circles", img );
    return 0;
}
```

**Note:** Usually the function detects the centers of circles well. However, it may fail to find correct radii. You can assist
to the function by specifying the radius range ( `minRadius` and `maxRadius` ) if you know it. Or, you may ignore the
returned radius, use only the center, and find the correct radius using an additional procedure.

**See Also:**

`fitEllipse()`, `minEnclosingCircle()`

## HoughLines

Finds lines in a binary image using the standard Hough transform.

**C++:** void **HoughLines**(InputArray **image**, OutputArray **lines**, double **rho**, double **theta**, int **threshold**, double **srn**=0, double **stn**=0 )

**Python:** cv2.**HoughLines**(image, rho, theta, threshold[, lines[, srn[, stn ] ] ]) → lines

**C:** CvSeq* **cvHoughLines2**(CvArr* **image**, void* **line_storage**, int **method**, double **rho**, double **theta**, int **threshold**, double **param1**=0, double **param2**=0 )

**Python:** cv.**HoughLines2**(image, storage, method, rho, theta, threshold, param1=0, param2=0) → lines

> **Parameters**

>> **image** – 8-bit, single-channel binary source image. The image may be modified by the function.

>> **lines** – Output vector of lines. Each line is represented by a two-element vector $(\rho, \theta)$ . $\rho$ is the distance from the coordinate origin $(0, 0)$ (top-left corner of the image). $\theta$ is the line rotation angle in radians ( $0 \sim$ vertical line, $\pi/2 \sim$ horizontal line ).

>> **rho** – Distance resolution of the accumulator in pixels.

>> **theta** – Angle resolution of the accumulator in radians.

>> **threshold** – Accumulator threshold parameter. Only those lines are returned that get enough votes ( $>$ `threshold` ).

>> **srn** – For the multi-scale Hough transform, it is a divisor for the distance resolution `rho` . The coarse accumulator distance resolution is `rho` and the accurate accumulator resolution is `rho/srn` . If both `srn=0` and `stn=0` , the classical Hough transform is used. Otherwise, both these parameters should be positive.

>> **stn** – For the multi-scale Hough transform, it is a divisor for the distance resolution `theta`.

>> **method** – One of the following Hough transform variants:

>> – **CV_HOUGH_STANDARD** classical or standard Hough transform. Every line is represented by two floating-point numbers $(\rho, \theta)$ , where $\rho$ is a distance between (0,0) point and the line, and $\theta$ is the angle between x-axis and the normal to the line. Thus, the matrix must be (the created sequence will be) of `CV_32FC2` type

>> – **CV_HOUGH_PROBABILISTIC** probabilistic Hough transform (more efficient in case if the picture contains a few long linear segments). It returns line segments rather than the whole line. Each segment is represented by starting and ending points, and the matrix must be (the created sequence will be) of the `CV_32SC4` type.

>> – **CV_HOUGH_MULTI_SCALE** multi-scale variant of the classical Hough transform. The lines are encoded the same way as `CV_HOUGH_STANDARD`.

>> **param1** – First method-dependent parameter:

>> – For the classical Hough transform, it is not used (0).

>> – For the probabilistic Hough transform, it is the minimum line length.

>> – For the multi-scale Hough transform, it is `srn`.

>> **param2** – Second method-dependent parameter:

>> – For the classical Hough transform, it is not used (0).

>> – For the probabilistic Hough transform, it is the maximum gap between line segments lying on the same line to treat them as a single line segment (that is, to join them).

>> – For the multi-scale Hough transform, it is `stn`.

The function implements the standard or standard multi-scale Hough transform algorithm for line detection. See
http://homepages.inf.ed.ac.uk/rbf/HIPR2/hough.htm for a good explanation of Hough transform. See also the example
in HoughLinesP() description.

## HoughLinesP

Finds line segments in a binary image using the probabilistic Hough transform.

**C++:** void **HoughLinesP**(InputArray **image**, OutputArray **lines**, double **rho**, double **theta**, int **threshold**, dou-
ble **minLineLength**=0, double **maxLineGap**=0 )

**Python:** cv2.**HoughLinesP**(image, rho, theta, threshold[, lines[, minLineLength[, maxLineGap ]]]) →
lines

Parameters

**image** – 8-bit, single-channel binary source image. The image may be modified by the
function.

**lines** – Output vector of lines. Each line is represented by a 4-element vector $(x_1, y_1, x_2, y_2)$
, where $(x_1, y_1)$ and $(x_2, y_2)$ are the ending points of each detected line segment.

**rho** – Distance resolution of the accumulator in pixels.

**theta** – Angle resolution of the accumulator in radians.

**threshold** – Accumulator threshold parameter. Only those lines are returned that get enough
votes ( > threshold ).

**minLineLength** – Minimum line length. Line segments shorter than that are rejected.

**maxLineGap** – Maximum allowed gap between points on the same line to link them.

The function implements the probabilistic Hough transform algorithm for line detection, described in [Matas00]. See
the line detection example below:

```
/* This is a standalone program. Pass an image name as the first parameter
of the program.  Switch between standard and probabilistic Hough transform
by changing "#if 1" to "#if 0" and back */
#include <cv.h>
#include <highgui.h>
#include <math.h>

using namespace cv;

int main(int argc, char** argv)
{
    Mat src, dst, color_dst;
    if( argc != 2 || !(src=imread(argv[1], 0)).data)
        return -1;

    Canny( src, dst, 50, 200, 3 );
    cvtColor( dst, color_dst, CV_GRAY2BGR );

#if 0
    vector<Vec2f> lines;
    HoughLines( dst, lines, 1, CV_PI/180, 100 );

    for( size_t i = 0; i < lines.size(); i++ )
    {
        float rho = lines[i][0];
```

```
        float theta = lines[i][1];
        double a = cos(theta), b = sin(theta);
        double x0 = a*rho, y0 = b*rho;
        Point pt1(cvRound(x0 + 1000*(-b)),
                  cvRound(y0 + 1000*(a)));
        Point pt2(cvRound(x0 - 1000*(-b)),
                  cvRound(y0 - 1000*(a)));
        line( color_dst, pt1, pt2, Scalar(0,0,255), 3, 8 );
    }
#else
    vector<Vec4i> lines;
    HoughLinesP( dst, lines, 1, CV_PI/180, 80, 30, 10 );
    for( size_t i = 0; i < lines.size(); i++ )
    {
        line( color_dst, Point(lines[i][0], lines[i][1]),
            Point(lines[i][2], lines[i][3]), Scalar(0,0,255), 3, 8 );
    }
#endif
    namedWindow( "Source", 1 );
    imshow( "Source", src );

    namedWindow( "Detected Lines", 1 );
    imshow( "Detected Lines", color_dst );

    waitKey(0);
    return 0;
}
```
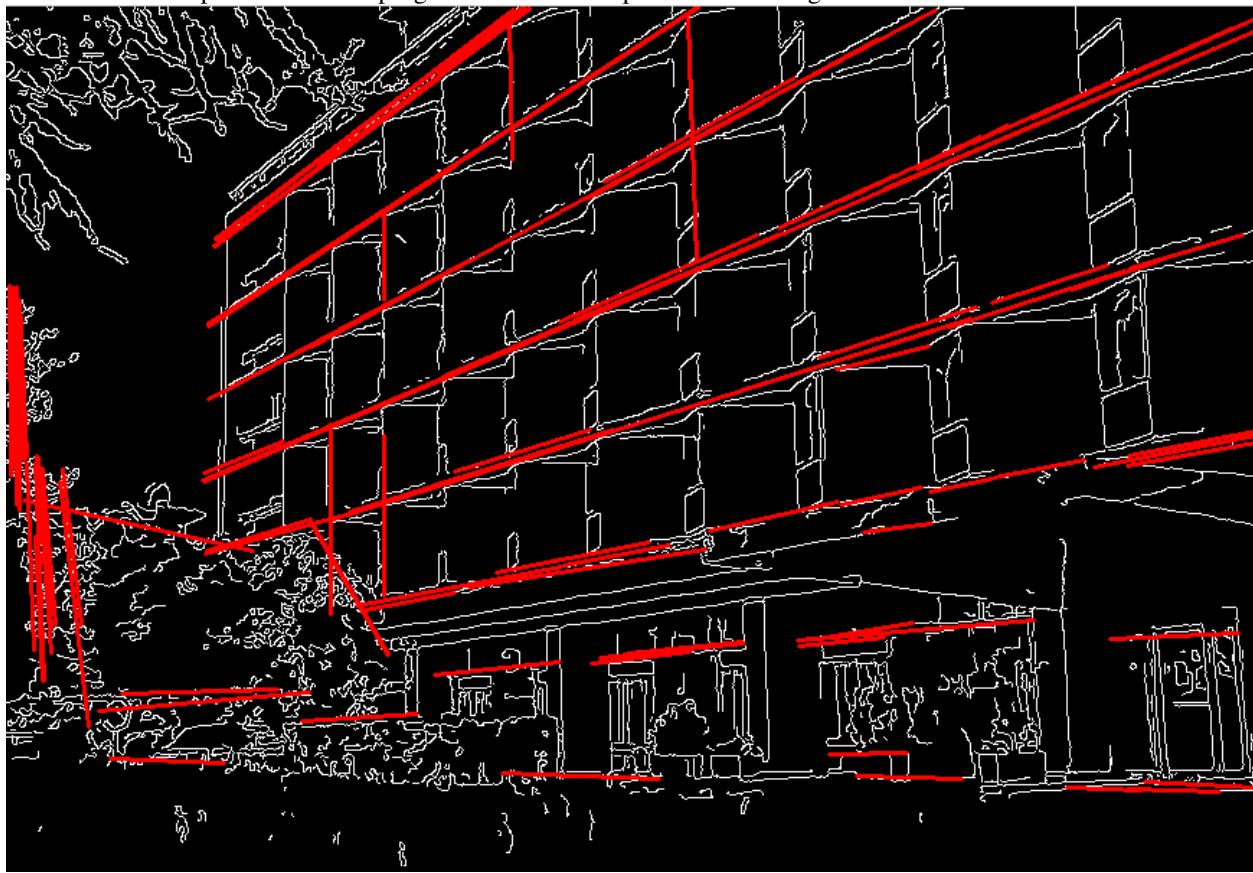
This is a sample picture the function parameters have been tuned for:

And this is the output of the above program in case of the probabilistic Hough transform:



## preCornerDetect

Calculates a feature map for corner detection.

**C++:** void **preCornerDetect**(InputArray **src**, OutputArray **dst**, int **ksize**, int **border-Type**=BORDER_DEFAULT )

**Python:** cv2.**preCornerDetect**(src, ksize[, dst[, borderType ]]) → dst

**C:** void **cvPreCornerDetect**(const CvArr* **image**, CvArr* **corners**, int **aperture_size**=3 )

**Python:** cv.**PreCornerDetect**(image, corners, apertureSize=3) → None

>  **Parameters**

>>  **src** – Source single-channel 8-bit of floating-point image.

>>  **dst** – Output image that has the type `CV_32F` and the same size as `src` .

>>  **ksize** – Aperture size of the `Sobel()` .

>>  **borderType** – Pixel extrapolation method. See `borderInterpolate()` .

The function calculates the complex spatial derivative-based function of the source image

$$\mathtt{dst} = (D_x\mathtt{src})^2 \cdot D_{yy}\mathtt{src} + (D_y\mathtt{src})^2 \cdot D_{xx}\mathtt{src} - 2D_x\mathtt{src} \cdot D_y\mathtt{src} \cdot D_{xy}\mathtt{src}$$

where $D_x$,:math:`D_y` are the first image derivatives, $D_{xx}$,:math:`D_{yy}` are the second image derivatives, and $D_{xy}$ is the mixed derivative.

The corners can be found as local maximums of the functions, as shown below:

```
Mat corners, dilated_corners;
preCornerDetect(image, corners, 3);
// dilation with 3x3 rectangular structuring element
dilate(corners, dilated_corners, Mat(), 1);
Mat corner_mask = corners == dilated_corners;
```

# 3.8 Object Detection

## matchTemplate

Compares a template against overlapped image regions.

**C++:** void **matchTemplate**(InputArray **image**, InputArray **templ**, OutputArray **result**, int **method**)

**Python:** cv2.**matchTemplate**(image, templ, method[, result]) → result

**C:** void **cvMatchTemplate**(const CvArr* **image**, const CvArr* **templ**, CvArr* **result**, int **method**)

**Python:** cv.**MatchTemplate**(image, templ, result, method) → None

> **Parameters**

>> **image** – Image where the search is running. It must be 8-bit or 32-bit floating-point.

>> **templ** – Searched template. It must be not greater than the source image and have the same data type.

>> **result** – Map of comparison results. It must be single-channel 32-bit floating-point. If image is $W \times H$ and templ is $w \times h$, then result is $(W - w + 1) \times (H - h + 1)$.

>> **method** – Parameter specifying the comparison method (see below).

The function slides through image , compares the overlapped patches of size $w \times h$ against templ using the specified method and stores the comparison results in result . Here are the formulae for the available comparison methods ( I denotes image, T template, R result ). The summation is done over template and/or the image patch: $x' = 0...w - 1, y' = 0...h - 1$ * method=CV_TM_SQDIFF

$$R(x, y) = \sum_{x', y'} (T(x', y') - I(x + x', y + y'))^2$$

- method=CV_TM_SQDIFF_NORMED

$$R(x, y) = \frac{\sum_{x', y'} (T(x', y') - I(x + x', y + y'))^2}{\sqrt{\sum_{x', y'} T(x', y')^2 \cdot \sum_{x', y'} I(x + x', y + y')^2}}$$

- method=CV_TM_CCORR

$$R(x, y) = \sum_{x', y'} (T(x', y') \cdot I(x + x', y + y'))$$

- method=CV_TM_CCORR_NORMED

$$R(x, y) = \frac{\sum_{x', y'} (T(x', y') \cdot I(x + x', y + y'))}{\sqrt{\sum_{x', y'} T(x', y')^2 \cdot \sum_{x', y'} I(x + x', y + y')^2}}$$

- method=CV_TM_CCOEFF

$$R(x, y) = \sum_{x', y'} (T'(x', y') \cdot I'(x + x', y + y'))$$

where

$$T'(x', y') = T(x', y') - 1/(w \cdot h) \cdot \sum_{x'', y''} T(x'', y'')$$
$$I'(x + x', y + y') = I(x + x', y + y') - 1/(w \cdot h) \cdot \sum_{x'', y''} I(x + x'', y + y'')$$

- method=CV_TM_CCOEFF_NORMED

$$R(x, y) = \frac{\sum_{x', y'} (T'(x', y') \cdot I'(x + x', y + y'))}{\sqrt{\sum_{x', y'} T'(x', y')^2 \cdot \sum_{x', y'} I'(x + x', y + y')^2}}$$

After the function finishes the comparison, the best matches can be found as global minimums (when `CV_TM_SQDIFF` was used) or maximums (when `CV_TM_CCORR` or `CV_TM_CCOEFF` was used) using the `minMaxLoc()` function. In case of a color image, template summation in the numerator and each sum in the denominator is done over all of the channels and separate mean values are used for each channel. That is, the function can take a color template and a color image. The result will still be a single-channel image, which is easier to analyze.

# HIGHGUI. HIGH-LEVEL GUI AND MEDIA I/O

While OpenCV was designed for use in full-scale applications and can be used within functionally rich UI frameworks (such as Qt*, WinForms*, or Cocoa*) or without any UI at all, sometimes there it is required to try functionality quickly and visualize the results. This is what the HighGUI module has been designed for.

It provides easy interface to:

- Create and manipulate windows that can display images and "remember" their content (no need to handle repaint events from OS).

- Add trackbars to the windows, handle simple mouse events as well as keyboard commands.

- Read and write images to/from disk or memory.

- Read video from camera or file and write video to a file.

## 4.1 User Interface

### createTrackbar

Creates a trackbar and attaches it to the specified window.

**C++:** int **createTrackbar**(const string& **trackbarname**, const string& **winname**, int* **value**, int **count**, TrackbarCallback **onChange**=0, void* **userdata**=0)

**C:** int **cvCreateTrackbar**(const char* **trackbar_name**, const char* **window_name**, int* **value**, int **count**, CvTrackbarCallback **on_change**=NULL )

**Python:** cv.**CreateTrackbar**(trackbarName, windowName, value, count, onChange) → None

> **Parameters**
>
>> **trackbarname** – Name of the created trackbar.
>>
>> **winname** – Name of the window that will be used as a parent of the created trackbar.
>>
>> **value** – Optional pointer to an integer variable whose value reflects the position of the slider. Upon creation, the slider position is defined by this variable.
>>
>> **count** – Maximal position of the slider. The minimal position is always 0.
>>
>> **onChange** – Pointer to the function to be called every time the slider changes position. This function should be prototyped as void Foo(int,void*); , where the first parameter is the

trackbar position and the second parameter is the user data (see the next parameter). If the callback is the NULL pointer, no callbacks are called, but only `value` is updated.

> **userdata** – User data that is passed as is to the callback. It can be used to handle trackbar events without using global variables.

The function `createTrackbar` creates a trackbar (a slider or range control) with the specified name and range, assigns a variable `value` to be a position synchronized with the trackbar and specifies the callback function `onChange` to be called on the trackbar position change. The created trackbar is displayed in the specified window `winname`.

---

**Note:** **[Qt Backend Only]** `winname` can be empty (or NULL) if the trackbar should be attached to the control panel.

---

Clicking the label of each trackbar enables editing the trackbar values manually.

## getTrackbarPos

Returns the trackbar position.

**C++:** int **getTrackbarPos**(const string& **trackbarname**, const string& **winname**)

**Python:** cv2.**getTrackbarPos**(trackbarname, winname) → retval

**C:** int **cvGetTrackbarPos**(const char* **trackbar_name**, const char* **window_name**)

**Python:** cv.**GetTrackbarPos**(trackbarName, windowName) → retval

> **Parameters**

> > **trackbarname** – Name of the trackbar.

> > **winname** – Name of the window that is the parent of the trackbar.

The function returns the current position of the specified trackbar.

---

**Note:** **[Qt Backend Only]** `winname` can be empty (or NULL) if the trackbar is attached to the control panel.

---

## imshow

Displays an image in the specified window.

**C++:** void **imshow**(const string& **winname**, InputArray **mat**)

**Python:** cv2.**imshow**(winname, mat) → None

**C:** void **cvShowImage**(const char* **name**, const CvArr* **image**)

**Python:** cv.**ShowImage**(name, image) → None

> **Parameters**

> > **winname** – Name of the window.

> > **image** – Image to be shown.

The function `imshow` displays an image in the specified window. If the window was created with the `CV_WINDOW_AUTOSIZE` flag, the image is shown with its original size. Otherwise, the image is scaled to fit the window. The function may scale the image, depending on its depth:

- If the image is 8-bit unsigned, it is displayed as is.

---

- If the image is 16-bit unsigned or 32-bit integer, the pixels are divided by 256. That is, the value range [0,255*256] is mapped to [0,255].

- If the image is 32-bit floating-point, the pixel values are multiplied by 255. That is, the value range [0,1] is mapped to [0,255].

## namedWindow

Creates a window.

**C++:** void **namedWindow**(const string& **winname**, int **flags**=WINDOW_AUTOSIZE )

**Python:** cv2.**namedWindow**(winname[, flags]) → None

**C:** int **cvNamedWindow**(const char* **name**, int **flags**=CV_WINDOW_AUTOSIZE )

**Python:** cv.**NamedWindow**(name, flags=CV_WINDOW_AUTOSIZE) → None

>    **Parameters**

>        **name** – Name of the window in the window caption that may be used as a window identifier.

>        **flags** – Flags of the window. Currently the only supported flag is CV_WINDOW_AUTOSIZE . If this is set, the window size is automatically adjusted to fit the displayed image (see imshow() ), and you cannot change the window size manually.

The function namedWindow creates a window that can be used as a placeholder for images and trackbars. Created windows are referred to by their names.

If a window with the same name already exists, the function does nothing.

You can call destroyWindow() or destroyAllWindows() to close the window and de-allocate any associated memory usage. For a simple program, you do not really have to call these functions because all the resources and windows of the application are closed automatically by the operating system upon exit.

---

**Note:** Qt backend supports additional flags:

- **CV_WINDOW_NORMAL or CV_WINDOW_AUTOSIZE:** CV_WINDOW_NORMAL enables you to resize the window, whereas CV_WINDOW_AUTOSIZE adjusts automatically the window size to fit the displayed image (see imshow() ), and you cannot change the window size manually.

- **CV_WINDOW_FREERATIO or CV_WINDOW_KEEPRATIO:** CV_WINDOW_FREERATIO adjusts the image with no respect to its ratio, whereas CV_WINDOW_KEEPRATIO keeps the image ratio.

- **CV_GUI_NORMAL or CV_GUI_EXPANDED:** CV_GUI_NORMAL is the old way to draw the window without statusbar and toolbar, whereas CV_GUI_EXPANDED is a new enhanced GUI.

By default, flags == CV_WINDOW_AUTOSIZE | CV_WINDOW_KEEPRATIO | CV_GUI_EXPANDED

---

## destroyWindow

Destroys a window.

**C++:** void **destroyWindow**(const string& **winname**)

**Python:** cv2.**destroyWindow**(winname) → None

**C:** void **cvDestroyWindow**(const char* **name**)

**Python:** cv.**DestroyWindow**(name) → None

---

> **Parameters**
>
> > **winname** – Name of the window to be destroyed.

The function `destroyWindow` destroys the window with the given name.

## destroyAllWindows

Destroys all of the HighGUI windows.

**C++:** void **destroyAllWindows**()

**Python:** cv2.**destroyAllWindows**() → None

**C:** void **cvDestroyAllWindows**()

**Python:** cv.**DestroyAllWindows**() → None

The function `destroyAllWindows` destroys all of the opened HighGUI windows.

## MoveWindow

Moves window to the specified position

**C++:** void **moveWindow**(const string& **winname**, int **x**, int **y**)

**Python:** cv2.**moveWindow**(winname, x, y) → None

**C:** void **cvMoveWindow**(const char* **name**, int **x**, int **y**)

**Python:** cv.**MoveWindow**(name, x, y) → None

> **Parameters**
>
> > **winname** – Window name
> >
> > **x** – The new x-coordinate of the window
> >
> > **y** – The new y-coordinate of the window

## ResizeWindow

Resizes window to the specified size

**C++:** void **resizeWindow**(const string& **winname**, int **width**, int **height**)

**Python:** cv2.**resizeWindow**(winname, width, height) → None

**C:** void **cvResizeWindow**(const char* **name**, int **width**, int **height**)

**Python:** cv.**ResizeWindow**(name, width, height) → None

> **Parameters**
>
> > **winname** – Window name
> >
> > **width** – The new window width
> >
> > **height** – The new window height

**Note:**

- The specified window size is for the image area. Toolbars are not counted.

> • Only windows created without CV_WINDOW_AUTOSIZE flag can be resized.

## SetMouseCallback

Sets mouse handler for the specified window

**C++:** void **setMouseCallback**(const string& **winname**, MouseCallback **onMouse**, void* **userdata**=0 )

**C:** void **cvSetMouseCallback**(const char* **window_name**, CvMouseCallback **on_mouse**, void* **param**=NULL )

**Python:** cv.**SetMouseCallback**(windowName, onMouse, param=None) → None

>    **Parameters**

>        **winname** – Window name

>        **onMouse** – Mouse callback. See OpenCV samples, such as http://code.opencv.org/projects/opencv/repository/revisions/master/entry/samples/cpp/ffilldemo.cpp, on how to specify and use the callback.

>        **userdata** – The optional parameter passed to the callback.

## setTrackbarPos

Sets the trackbar position.

**C++:** void **setTrackbarPos**(const string& **trackbarname**, const string& **winname**, int **pos**)

**Python:** cv2.**setTrackbarPos**(trackbarname, winname, pos) → None

**C:** void **cvSetTrackbarPos**(const char* **trackbar_name**, const char* **window_name**, int **pos**)

**Python:** cv.**SetTrackbarPos**(trackbarName, windowName, pos) → None

>    **Parameters**

>        **trackbarname** – Name of the trackbar.

>        **winname** – Name of the window that is the parent of trackbar.

>        **pos** – New position.

The function sets the position of the specified trackbar in the specified window.

**Note: [Qt Backend Only]** `winname` can be empty (or NULL) if the trackbar is attached to the control panel.

## waitKey

Waits for a pressed key.

**C++:** int **waitKey**(int **delay**=0)

**Python:** cv2.**waitKey**([delay]) → retval

**C:** int **cvWaitKey**(int **delay**=0 )

**Python:** cv.**WaitKey**(delay=0) → int

>    **Parameters**

> **delay** – Delay in milliseconds. 0 is the special value that means "forever".

The function `waitKey` waits for a key event infinitely (when `delay` ≤ 0 ) or for `delay` milliseconds, when it is positive. Since the OS has a minimum time between switching threads, the function will not wait exactly `delay` ms, it will wait at least `delay` ms, depending on what else is running on your computer at that time. It returns the code of the pressed key or -1 if no key was pressed before the specified time had elapsed.

---

**Note:** This function is the only method in HighGUI that can fetch and handle events, so it needs to be called periodically for normal event processing unless HighGUI is used within an environment that takes care of event processing.

---

---

**Note:** The function only works if there is at least one HighGUI window created and the window is active. If there are several HighGUI windows, any of them can be active.

---

## 4.2 Reading and Writing Images and Video

### imdecode

Reads an image from a buffer in memory.

**C++:** Mat **imdecode**(InputArray **buf**, int **flags**)

**C++:** Mat **imdecode**(InputArray **buf**, int **flags**, Mat* **dst**)

**C:** IplImage* **cvDecodeImage**(const CvMat* **buf**, int **iscolor**=CV_LOAD_IMAGE_COLOR)

**C:** CvMat* **cvDecodeImageM**(const CvMat* **buf**, int **iscolor**=CV_LOAD_IMAGE_COLOR)

**Python:** `cv2.imdecode`(buf, flags) → retval

> **Parameters**
>
>> **buf** – Input array or vector of bytes.
>>
>> **flags** – The same flags as in `imread()` .
>>
>> **dst** – The optional output placeholder for the decoded matrix. It can save the image reallocations when the function is called repeatedly for images of the same size.

The function reads an image from the specified buffer in the memory. If the buffer is too short or contains invalid data, the empty matrix/image is returned.

See `imread()` for the list of supported formats and flags description.

---

**Note:** In the case of color images, the decoded images will have the channels stored in `B G R` order.

---

### imencode

Encodes an image into a memory buffer.

**C++:** bool **imencode**(const string& **ext**, InputArray **img**, vector<uchar>& **buf**, const vector<int>& **params**=vector<int>())

**C:** CvMat* **cvEncodeImage**(const char* **ext**, const CvArr* **image**, const int* **params**=0 )

---

**Python:** cv2.**imencode**(ext, img[, params ]) → retval, buf

> **Parameters**

>> **ext** – File extension that defines the output format.

>> **img** – Image to be written.

>> **buf** – Output buffer resized to fit the compressed image.

>> **params** – Format-specific parameters. See imwrite() .

The function compresses the image and stores it in the memory buffer that is resized to fit the result. See imwrite() for the list of supported formats and flags description.

---

**Note:** cvEncodeImage returns single-row matrix of type CV_8UC1 that contains encoded image as array of bytes.

---

## imread

Loads an image from a file.

**C++:** Mat **imread**(const string& **filename**, int **flags**=1 )

**Python:** cv2.**imread**(filename[, flags ]) → retval

**C:** IplImage* **cvLoadImage**(const char* **filename**, int **iscolor**=CV_LOAD_IMAGE_COLOR )

**C:** CvMat* **cvLoadImageM**(const char* **filename**, int **iscolor**=CV_LOAD_IMAGE_COLOR )

**Python:** cv.**LoadImage**(filename, iscolor=CV_LOAD_IMAGE_COLOR) → None

**Python:** cv.**LoadImageM**(filename, iscolor=CV_LOAD_IMAGE_COLOR) → None

> **Parameters**

>> **filename** – Name of file to be loaded.

>> **flags** – Flags specifying the color type of a loaded image:

>> – 1 -

>> – CV_LOAD_IMAGE_ANYDEPTH -

>> CV_LOAD_IMAGE_COLOR CV_LOAD_IMAGE_GRAYSCALE

>> – **>0 Return a 3-channel color image.**

>>> ---

>>> **Note:** In the current implementation the alpha channel, if any, is stripped from the output image. Use negative value if you need the alpha channel.

>>> ---

>> – **=0** Return a grayscale image.

>> – **<0** Return the loaded image as is (with alpha channel).

The function imread loads an image from the specified file and returns it. If the image cannot be read (because of missing file, improper permissions, unsupported or invalid format), the function returns an empty matrix ( Mat::data==NULL ). Currently, the following file formats are supported:

- Windows bitmaps - *.bmp, *.dib (always supported)

- JPEG files - *.jpeg, *.jpg, *.jpe (see the *Notes* section)

- JPEG 2000 files - *.jp2 (see the *Notes* section)

- Portable Network Graphics - *.png (see the *Notes* section)

- Portable image format - *.pbm, *.pgm, *.ppm (always supported)

- Sun rasters - *.sr, *.ras (always supported)

- TIFF files - *.tiff, *.tif (see the *Notes* section)

---

**Note:**

- The function determines the type of an image by the content, not by the file extension.

- On Microsoft Windows* OS and MacOSX*, the codecs shipped with an OpenCV image (libjpeg, libpng, libtiff, and libjasper) are used by default. So, OpenCV can always read JPEGs, PNGs, and TIFFs. On MacOSX, there is also an option to use native MacOSX image readers. But beware that currently these native image loaders give images with different pixel values because of the color management embedded into MacOSX.

- On Linux*, BSD flavors and other Unix-like open-source operating systems, OpenCV looks for codecs supplied with an OS image. Install the relevant packages (do not forget the development files, for example, "libjpeg-dev", in Debian* and Ubuntu*) to get the codec support or turn on the OPENCV_BUILD_3RDPARTY_LIBS flag in CMake.

---

**Note:** In the case of color images, the decoded images will have the channels stored in B G R order.

---

## imwrite

Saves an image to a specified file.

**C++:** bool **imwrite**(const string& **filename**, InputArray **img**, const vector<int>& **params**=vector<int>() )

**Python:** cv2.**imwrite**(filename, img[, params ]) → retval

**C:** int **cvSaveImage**(const char* **filename**, const CvArr* **image**, const int* **params**=0 )

**Python:** cv.**SaveImage**(filename, image) → None

> **Parameters**
>
> > **filename** – Name of the file.
> >
> > **image** – Image to be saved.
> >
> > **params** – Format-specific save parameters encoded as pairs paramId_1, paramValue_1, paramId_2, paramValue_2, ... . The following parameters are currently supported:
> >
> > - For JPEG, it can be a quality ( CV_IMWRITE_JPEG_QUALITY ) from 0 to 100 (the higher is the better). Default value is 95.
> >
> > - For PNG, it can be the compression level ( CV_IMWRITE_PNG_COMPRESSION ) from 0 to 9. A higher value means a smaller size and longer compression time. Default value is 3.
> >
> > - For PPM, PGM, or PBM, it can be a binary format flag ( CV_IMWRITE_PXM_BINARY ), 0 or 1. Default value is 1.

The function imwrite saves the image to the specified file. The image format is chosen based on the filename extension (see imread() for the list of extensions). Only 8-bit (or 16-bit unsigned (CV_16U) in case of PNG, JPEG 2000, and TIFF) single-channel or 3-channel (with 'BGR' channel order) images can be saved using this function. If the format, depth or channel order is different, use Mat::convertTo() , and cvtColor() to convert it before saving. Or, use the universal XML I/O functions to save the image to XML or YAML format.

---

It is possible to store PNG images with an alpha channel using this function. To do this, create 8-bit (or 16-bit) 4-channel image BGRA, where the alpha channel goes last. Fully transparent pixels should have alpha set to 0, fully opaque pixels should have alpha set to 255/65535. The sample below shows how to create such a BGRA image and store to PNG file. It also demonstrates how to set custom compression parameters

```
#include <vector>
#include <stdio.h>
#include <opencv2/opencv.hpp>

using namespace cv;
using namespace std;

void createAlphaMat(Mat &mat)
{
    for (int i = 0; i < mat.rows; ++i) {
        for (int j = 0; j < mat.cols; ++j) {
            Vec4b& rgba = mat.at<Vec4b>(i, j);
            rgba[0] = UCHAR_MAX;
            rgba[1] = saturate_cast<uchar>((float (mat.cols - j)) / ((float)mat.cols) * UCHAR_MAX);
            rgba[2] = saturate_cast<uchar>((float (mat.rows - i)) / ((float)mat.rows) * UCHAR_MAX);
            rgba[3] = saturate_cast<uchar>(0.5 * (rgba[1] + rgba[2]));
        }
    }
}

int main(int argv, char **argc)
{
    // Create mat with alpha channel
    Mat mat(480, 640, CV_8UC4);
    createAlphaMat(mat);

    vector<int> compression_params;
    compression_params.push_back(CV_IMWRITE_PNG_COMPRESSION);
    compression_params.push_back(9);

    try {
        imwrite("alpha.png", mat, compression_params);
    }
    catch (runtime_error& ex) {
        fprintf(stderr, "Exception converting image to PNG format: %s\n", ex.what());
        return 1;
    }

    fprintf(stdout, "Saved PNG file with alpha data.\n");
    return 0;
}
```

## VideoCapture

**class** `VideoCapture`

Class for video capturing from video files or cameras. The class provides C++ API for capturing video from cameras or for reading video files. Here is how the class can be used:

```
#include "opencv2/opencv.hpp"

using namespace cv;
```

---

```cpp
int main(int, char**)
{
    VideoCapture cap(0); // open the default camera
    if(!cap.isOpened())  // check if we succeeded
        return -1;

    Mat edges;
    namedWindow("edges",1);
    for(;;)
    {
        Mat frame;
        cap >> frame; // get a new frame from camera
        cvtColor(frame, edges, CV_BGR2GRAY);
        GaussianBlur(edges, edges, Size(7,7), 1.5, 1.5);
        Canny(edges, edges, 0, 30, 3);
        imshow("edges", edges);
        if(waitKey(30) >= 0) break;
    }
    // the camera will be deinitialized automatically in VideoCapture destructor
    return 0;
}
```

**Note:** In C API the black-box structure `CvCapture` is used instead of `VideoCapture`.

## VideoCapture::VideoCapture

VideoCapture constructors.

**C++:** `VideoCapture::`**`VideoCapture`**`()`

**C++:** `VideoCapture::`**`VideoCapture`**`(const string& `**filename**`)`

**C++:** `VideoCapture::`**`VideoCapture`**`(int `**device**`)`

**Python:** `cv2.`**`VideoCapture`**`()` → <VideoCapture object>

**Python:** `cv2.`**`VideoCapture`**`(filename)` → <VideoCapture object>

**Python:** `cv2.`**`VideoCapture`**`(device)` → <VideoCapture object>

**C:** `CvCapture* `**`cvCaptureFromCAM`**`(int `**device**`)`

**Python:** `cv.`**`CaptureFromCAM`**`(index)` → CvCapture

**C:** `CvCapture* `**`cvCaptureFromFile`**`(const char* `**filename**`)`

**Python:** `cv.`**`CaptureFromFile`**`(filename)` → CvCapture

> **Parameters**
>
> > **filename** – name of the opened video file
> >
> > **device** – id of the opened video capturing device (i.e. a camera index). If there is a single camera connected, just pass 0.

**Note:** In C API, when you finished working with video, release `CvCapture` structure with `cvReleaseCapture()`, or use `Ptr<CvCapture>` that calls `cvReleaseCapture()` automatically in the destructor.

## VideoCapture::open

Open video file or a capturing device for video capturing

**C++:** bool VideoCapture::**open**(const string& **filename**)

**C++:** bool VideoCapture::**open**(int **device**)

**Python:** cv2.VideoCapture.**open**(filename) → retval

**Python:** cv2.VideoCapture.**open**(device) → retval

>>> **Parameters**

>>>> **filename** – name of the opened video file

>>>> **device** – id of the opened video capturing device (i.e. a camera index).

The methods first call VideoCapture::release() to close the already opened file or camera.

## VideoCapture::isOpened

Returns true if video capturing has been initialized already.

**C++:** bool VideoCapture::**isOpened**()

**Python:** cv2.VideoCapture.**isOpened**() → retval

If the previous call to VideoCapture constructor or VideoCapture::open succeeded, the method returns true.

## VideoCapture::release

Closes video file or capturing device.

**C++:** void VideoCapture::**release**()

**Python:** cv2.VideoCapture.**release**() → None

**C:** void **cvReleaseCapture**(CvCapture** **capture**)

The methods are automatically called by subsequent VideoCapture::open() and by VideoCapture destructor.

The C function also deallocates memory and clears *capture pointer.

## VideoCapture::grab

Grabs the next frame from video file or capturing device.

**C++:** bool VideoCapture::**grab**()

**Python:** cv2.VideoCapture.**grab**() → retval

**C:** int **cvGrabFrame**(CvCapture* **capture**)

**Python:** cv.**GrabFrame**(capture) → int

The methods/functions grab the next frame from video file or camera and return true (non-zero) in the case of success.

The primary use of the function is in multi-camera environments, especially when the cameras do not have hardware synchronization. That is, you call VideoCapture::grab() for each camera and after that call the slower method VideoCapture::retrieve() to decode and get frame from each camera. This way the overhead on demosaicing or motion jpeg decompression etc. is eliminated and the retrieved frames from different cameras will be closer in time.

---

Also, when a connected camera is multi-head (for example, a stereo camera or a Kinect device), the correct way of retrieving data from it is to call *VideoCapture::grab* first and then call `VideoCapture::retrieve()` one or more times with different values of the channel parameter. See http://code.opencv.org/projects/opencv/repository/revisions/master/entry/samples/cpp/kinect_maps.cpp

## VideoCapture::retrieve

Decodes and returns the grabbed video frame.

**C++:** bool VideoCapture::**retrieve**(Mat& **image**, int **channel**=0)

**Python:** cv2.VideoCapture.**retrieve**([image[, channel]]) → retval, image

**C:** IplImage* **cvRetrieveFrame**(CvCapture* **capture**, int **streamIdx**=0 )

**Python:** cv.**RetrieveFrame**(capture) → image

The methods/functions decode and return the just grabbed frame. If no frames has been grabbed (camera has been disconnected, or there are no more frames in video file), the methods return false and the functions return NULL pointer.

---

**Note:** OpenCV 1.x functions cvRetrieveFrame and cv.RetrieveFrame return image stored inside the video capturing structure. It is not allowed to modify or release the image! You can copy the frame using `cvCloneImage()` and then do whatever you want with the copy.

---

## VideoCapture::read

Grabs, decodes and returns the next video frame.

**C++:** VideoCapture& VideoCapture::**operator>>**(Mat& **image**)

**C++:** bool VideoCapture::**read**(Mat& **image**)

**Python:** cv2.VideoCapture.**read**([image]) → retval, image

**C:** IplImage* **cvQueryFrame**(CvCapture* **capture**)

**Python:** cv.**QueryFrame**(capture) → image

The methods/functions combine `VideoCapture::grab()` and `VideoCapture::retrieve()` in one call. This is the most convenient method for reading video files or capturing data from decode and return the just grabbed frame. If no frames has been grabbed (camera has been disconnected, or there are no more frames in video file), the methods return false and the functions return NULL pointer.

---

**Note:** OpenCV 1.x functions cvRetrieveFrame and cv.RetrieveFrame return image stored inside the video capturing structure. It is not allowed to modify or release the image! You can copy the frame using `cvCloneImage()` and then do whatever you want with the copy.

---

## VideoCapture::get

Returns the specified `VideoCapture` property

**C++:** double VideoCapture::**get**(int **propId**)

**Python:** cv2.VideoCapture.**get**(propId) → retval

**C:** double **cvGetCaptureProperty**(CvCapture* **capture**, int **property_id**)

**Python:** cv.**GetCaptureProperty**(capture, property_id) → float

> **Parameters**
>
>> **propId** – Property identifier. It can be one of the following:
>>
>> – **CV_CAP_PROP_POS_MSEC** Current position of the video file in milliseconds or video capture timestamp.
>>
>> – **CV_CAP_PROP_POS_FRAMES** 0-based index of the frame to be decoded/captured next.
>>
>> – **CV_CAP_PROP_POS_AVI_RATIO** Relative position of the video file: 0 - start of the film, 1 - end of the film.
>>
>> – **CV_CAP_PROP_FRAME_WIDTH** Width of the frames in the video stream.
>>
>> – **CV_CAP_PROP_FRAME_HEIGHT** Height of the frames in the video stream.
>>
>> – **CV_CAP_PROP_FPS** Frame rate.
>>
>> – **CV_CAP_PROP_FOURCC** 4-character code of codec.
>>
>> – **CV_CAP_PROP_FRAME_COUNT** Number of frames in the video file.
>>
>> – **CV_CAP_PROP_FORMAT** Format of the Mat objects returned by `retrieve()`.
>>
>> – **CV_CAP_PROP_MODE** Backend-specific value indicating the current capture mode.
>>
>> – **CV_CAP_PROP_BRIGHTNESS** Brightness of the image (only for cameras).
>>
>> – **CV_CAP_PROP_CONTRAST** Contrast of the image (only for cameras).
>>
>> – **CV_CAP_PROP_SATURATION** Saturation of the image (only for cameras).
>>
>> – **CV_CAP_PROP_HUE** Hue of the image (only for cameras).
>>
>> – **CV_CAP_PROP_GAIN** Gain of the image (only for cameras).
>>
>> – **CV_CAP_PROP_EXPOSURE** Exposure (only for cameras).
>>
>> – **CV_CAP_PROP_CONVERT_RGB** Boolean flags indicating whether images should be converted to RGB.
>>
>> – **CV_CAP_PROP_WHITE_BALANCE** Currently not supported
>>
>> – **CV_CAP_PROP_RECTIFICATION** Rectification flag for stereo cameras (note: only supported by DC1394 v 2.x backend currently)

**Note**: When querying a property that is not supported by the backend used by the `VideoCapture` class, value 0 is returned.

## VideoCapture::set

Sets a property in the `VideoCapture`.

**C++:** bool VideoCapture::**set**(int **propId**, double **value**)

**Python:** cv2.VideoCapture.**set**(propId, value) → retval

**C:** int **cvSetCaptureProperty**(CvCapture* **capture**, int **property_id**, double **value**)

**Python:** cv.**SetCaptureProperty**(capture, property_id, value) → retval

> **Parameters**

> **propId** – Property identifier. It can be one of the following:
>
> – **CV_CAP_PROP_POS_MSEC** Current position of the video file in milliseconds.
>
> – **CV_CAP_PROP_POS_FRAMES** 0-based index of the frame to be decoded/captured next.
>
> – **CV_CAP_PROP_POS_AVI_RATIO** Relative position of the video file: 0 - start of the film, 1 - end of the film.
>
> – **CV_CAP_PROP_FRAME_WIDTH** Width of the frames in the video stream.
>
> – **CV_CAP_PROP_FRAME_HEIGHT** Height of the frames in the video stream.
>
> – **CV_CAP_PROP_FPS** Frame rate.
>
> – **CV_CAP_PROP_FOURCC** 4-character code of codec.
>
> – **CV_CAP_PROP_FRAME_COUNT** Number of frames in the video file.
>
> – **CV_CAP_PROP_FORMAT** Format of the Mat objects returned by `retrieve()`.
>
> – **CV_CAP_PROP_MODE** Backend-specific value indicating the current capture mode.
>
> – **CV_CAP_PROP_BRIGHTNESS** Brightness of the image (only for cameras).
>
> – **CV_CAP_PROP_CONTRAST** Contrast of the image (only for cameras).
>
> – **CV_CAP_PROP_SATURATION** Saturation of the image (only for cameras).
>
> – **CV_CAP_PROP_HUE** Hue of the image (only for cameras).
>
> – **CV_CAP_PROP_GAIN** Gain of the image (only for cameras).
>
> – **CV_CAP_PROP_EXPOSURE** Exposure (only for cameras).
>
> – **CV_CAP_PROP_CONVERT_RGB** Boolean flags indicating whether images should be converted to RGB.
>
> – **CV_CAP_PROP_WHITE_BALANCE** Currently unsupported
>
> – **CV_CAP_PROP_RECTIFICATION** Rectification flag for stereo cameras (note: only supported by DC1394 v 2.x backend currently)
>
> **value** – Value of the property.

## VideoWriter

**class VideoWriter**

Video writer class.

## VideoWriter::VideoWriter

VideoWriter constructors

**C++:** VideoWriter::**VideoWriter**()

**C++:** VideoWriter::**VideoWriter**(const string& **filename**, int **fourcc**, double **fps**, Size **frameSize**, bool **isColor**=true)

**Python:** cv2.**VideoWriter**([filename, fourcc, fps, frameSize[, isColor]]) → <VideoWriter object>

**C:** CvVideoWriter* **cvCreateVideoWriter**(const char* **filename**, int **fourcc**, double **fps**, CvSize **frame_size**, int **is_color**=1 )

**Python:** cv.**CreateVideoWriter**(filename, fourcc, fps, frame_size, is_color=true) → CvVideoWriter

**Python:** cv2.VideoWriter.**isOpened**() → retval

**Python:** cv2.VideoWriter.**open**(filename, fourcc, fps, frameSize[, isColor]) → retval

**Python:** cv2.VideoWriter.**write**(image) → None

> **Parameters**
>
> > **filename** – Name of the output video file.
> >
> > **fourcc** – 4-character code of codec used to compress the frames. For example, CV_FOURCC('P','I','M','1') is a MPEG-1 codec, CV_FOURCC('M','J','P','G') is a motion-jpeg codec etc.
> >
> > **fps** – Framerate of the created video stream.
> >
> > **frameSize** – Size of the video frames.
> >
> > **isColor** – If it is not zero, the encoder will expect and encode color frames, otherwise it will work with grayscale frames (the flag is currently supported on Windows only).

The constructors/functions initialize video writers. On Linux FFMPEG is used to write videos; on Windows FFMPEG or VFW is used; on MacOSX QTKit is used.

## ReleaseVideoWriter

Releases the AVI writer.

**C:** void **cvReleaseVideoWriter**(CvVideoWriter** **writer**)

The function should be called after you finished using CvVideoWriter opened with `CreateVideoWriter()`.

## VideoWriter::open

Initializes or reinitializes video writer.

**C++:** bool VideoWriter::**open**(const string& **filename**, int **fourcc**, double **fps**, Size **frameSize**, bool **isColor**=true)

**Python:** cv2.VideoWriter.**open**(filename, fourcc, fps, frameSize[, isColor]) → retval

The method opens video writer. Parameters are the same as in the constructor `VideoWriter::VideoWriter()`.

## VideoWriter::isOpened

Returns true if video writer has been successfully initialized.

**C++:** bool VideoWriter::**isOpened**()

**Python:** cv2.VideoWriter.**isOpened**() → retval

## VideoWriter::write

Writes the next video frame

**C++:** VideoWriter& VideoWriter::**operator<<**(const Mat& **image**)

**C++:** void VideoWriter::**write**(const Mat& **image**)

**Python:** `cv2.VideoWriter.write`(image) → None

**C:** int **cvWriteFrame**(CvVideoWriter* **writer**, const IplImage* **image**)

**Python:** `cv.WriteFrame`(writer, image) → int
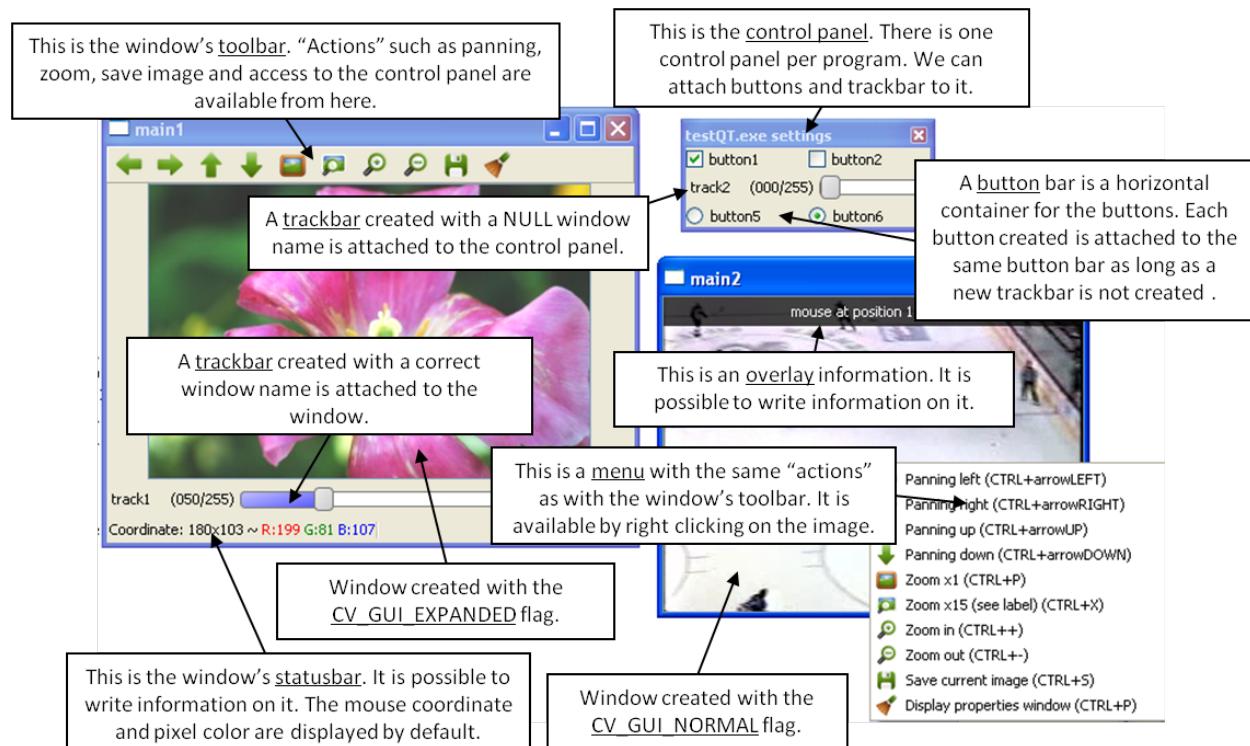
        **Parameters**

                **writer** – Video writer structure (OpenCV 1.x API)

                **image** – The written frame

The functions/methods write the specified image to video file. It must have the same size as has been specified when opening the video writer.

## 4.3 Qt New Functions



This figure explains new functionality implemented with Qt* GUI. The new GUI provides a statusbar, a toolbar, and a control panel. The control panel can have trackbars and buttonbars attached to it. If you cannot see the control panel, press Ctrl+P or right-click any Qt window and select **Display properties window**.

- To attach a trackbar, the window name parameter must be NULL.

- To attach a buttonbar, a button must be created. If the last bar attached to the control panel is a buttonbar, the new button is added to the right of the last button. If the last bar attached to the control panel is a trackbar, or the control panel is empty, a new buttonbar is created. Then, a new button is attached to it.

See below the example used to generate the figure:

```
int main(int argc, char *argv[])
    int value = 50;
    int value2 = 0;
```

```
cvNamedWindow("main1",CV_WINDOW_NORMAL);
cvNamedWindow("main2",CV_WINDOW_AUTOSIZE | CV_GUI_NORMAL);

cvCreateTrackbar( "track1", "main1", &value, 255,  NULL);//OK tested
char* nameb1 = "button1";
char* nameb2 = "button2";
cvCreateButton(nameb1,callbackButton,nameb1,CV_CHECKBOX,1);

cvCreateButton(nameb2,callbackButton,nameb2,CV_CHECKBOX,0);
cvCreateTrackbar( "track2", NULL, &value2, 255, NULL);
cvCreateButton("button5",callbackButton1,NULL,CV_RADIOBOX,0);
cvCreateButton("button6",callbackButton2,NULL,CV_RADIOBOX,1);

cvSetMouseCallback( "main2",on_mouse,NULL );

IplImage* img1 = cvLoadImage("files/flower.jpg");
IplImage* img2 = cvCreateImage(cvGetSize(img1),8,3);
CvCapture* video = cvCaptureFromFile("files/hockey.avi");
IplImage* img3 = cvCreateImage(cvGetSize(cvQueryFrame(video)),8,3);

while(cvWaitKey(33) != 27)
{
    cvAddS(img1,cvScalarAll(value),img2);
    cvAddS(cvQueryFrame(video),cvScalarAll(value2),img3);
    cvShowImage("main1",img2);
    cvShowImage("main2",img3);
}

cvDestroyAllWindows();
cvReleaseImage(&img1);
cvReleaseImage(&img2);
cvReleaseImage(&img3);
cvReleaseCapture(&video);
return 0;
}
```

## setWindowProperty

Changes parameters of a window dynamically.

**C++:** void **setWindowProperty**(const string& **winname**, int **prop_id**, double **prop_value**)

**Python:** cv2.**setWindowProperty**(winname, prop_id, prop_value) → None

**C:** void **cvSetWindowProperty**(const char* **name**, int **prop_id**, double **prop_value**)

> **Parameters**
>
> > **name** – Name of the window.
> >
> > **prop_id** – Window property to edit. The following operation flags are available:
> >
> > - **CV_WND_PROP_FULLSCREEN** Change if the window is fullscreen ( CV_WINDOW_NORMAL or CV_WINDOW_FULLSCREEN ).
> >
> > - **CV_WND_PROP_AUTOSIZE** Change if the window is resizable (CV_WINDOW_NORMAL or CV_WINDOW_AUTOSIZE ).
> >
> > - **CV_WND_PROP_ASPECTRATIO** Change if the aspect ratio of the image is preserved ( CV_WINDOW_FREERATIO or CV_WINDOW_KEEPRATIO ).

> > **prop_value** – New value of the window property. The following operation flags are available:

> > – **CV_WINDOW_NORMAL** Change the window to normal size or make the window resizable.

> > – **CV_WINDOW_AUTOSIZE** Constrain the size by the displayed image. The window is not resizable.

> > – **CV_WINDOW_FULLSCREEN** Change the window to fullscreen.

> > – **CV_WINDOW_FREERATIO** Make the window resizable without any ratio constraints.

> > – **CV_WINDOW_KEEPRATIO** Make the window resizable, but preserve the proportions of the displayed image.

The function `setWindowProperty` enables changing properties of a window.


## getWindowProperty

Provides parameters of a window.

**C++:** double **getWindowProperty**(const string& **winname**, int **prop_id**)

**Python:** cv2.**getWindowProperty**(winname, prop_id) → retval

**C:** double **cvGetWindowProperty**(const char* **name**, int **prop_id**)

> **Parameters**

> > **name** – Name of the window.

> > **prop_id** – Window property to retrieve. The following operation flags are available:

> > – **CV_WND_PROP_FULLSCREEN** Change if the window is fullscreen ( `CV_WINDOW_NORMAL` or `CV_WINDOW_FULLSCREEN` ).

> > – **CV_WND_PROP_AUTOSIZE** Change if the window is resizable (`CV_WINDOW_NORMAL` or `CV_WINDOW_AUTOSIZE` ).

> > – **CV_WND_PROP_ASPECTRATIO** Change if the aspect ratio of the image is preserved (`CV_WINDOW_FREERATIO` or `CV_WINDOW_KEEPRATIO` ).

See `setWindowProperty()` to know the meaning of the returned values.

The function `getWindowProperty` returns properties of a window.


## fontQt

Creates the font to draw a text on an image.

**C++:** CvFont **fontQt**(const string& **nameFont**, int **pointSize**=-1, Scalar **color**=Scalar::all(0), int **weight**=CV_FONT_NORMAL, int **style**=CV_STYLE_NORMAL, int **spacing**=0)

**C:** CvFont **cvFontQt**(const char* **nameFont**, int **pointSize**=-1, CvScalar **color**=cvScalarAll(0), int **weight**=CV_FONT_NORMAL, int **style**=CV_STYLE_NORMAL, int **spacing**=0)

> **Parameters**

> > **nameFont** – Name of the font. The name should match the name of a system font (such as *Times*). If the font is not found, a default one is used.

**pointSize** – Size of the font. If not specified, equal zero or negative, the point size of the font is set to a system-dependent default value. Generally, this is 12 points.

**color** – Color of the font in BGRA where A = 255 is fully transparent. Use the macro CV _ RGB for simplicity.

**weight** – Font weight. The following operation flags are available:

– **CV_FONT_LIGHT** Weight of 25

– **CV_FONT_NORMAL** Weight of 50

– **CV_FONT_DEMIBOLD** Weight of 63

– **CV_FONT_BOLD** Weight of 75

– **CV_FONT_BLACK** Weight of 87

You can also specify a positive integer for better control.

**style** – Font style. The following operation flags are available:

– **CV_STYLE_NORMAL** Normal font

– **CV_STYLE_ITALIC** Italic font

– **CV_STYLE_OBLIQUE** Oblique font

**spacing** – Spacing between characters. It can be negative or positive.

The function `fontQt` creates a CvFont object. This CvFont is not compatible with `putText` .

A basic usage of this function is the following:

```
CvFont font = fontQt(''Times'');
addText( img1, ''Hello World !'', Point(50,50), font);
```

## addText

Creates the font to draw a text on an image.

**C++:** void **addText**(const Mat& **img**, const string& **text**, Point **org**, CvFont **font**)

**C:** void **cvAddText**(const CvArr* **img**, const char* **text**, CvPoint **org**, CvFont* **arg2**)

> **Parameters**

> > **img** – 8-bit 3-channel image where the text should be drawn.

> > **text** – Text to write on an image.

> > **org** – Point(x,y) where the text should start on an image.

> > **font** – Font to use to draw a text.

The function `addText` draws *text* on an image *img* using a specific font *font* (see example `fontQt()` )

## displayOverlay

Displays a text on a window image as an overlay for a specified duration.

**C++:** void **displayOverlay**(const string& **winname**, const string& **text**, int **delayms**=0 )

**C:** void **cvDisplayOverlay**(const char* **name**, const char* **text**, int **delayms**=0)

> **Parameters**

> > **name** – Name of the window.

> > **text** – Overlay text to write on a window image.

> > **delayms** – The period (in milliseconds), during which the overlay text is displayed. If this function is called before the previous overlay text timed out, the timer is restarted and the text is updated. If this value is zero, the text never disappears.

The function `displayOverlay` displays useful information/tips on top of the window for a certain amount of time *delayms*. The function does not modify the image, displayed in the window, that is, after the specified delay the original content of the window is restored.

## displayStatusBar

Displays a text on the window statusbar during the specified period of time.

**C++:** void **displayStatusBar**(const string& **winname**, const string& **text**, int **delayms**=0 )

**C:** void **cvDisplayStatusBar**(const char* **name**, const char* **text**, int **delayms**=0)

> **Parameters**

> > **name** – Name of the window.

> > **text** – Text to write on the window statusbar.

> > **delayms** – Duration (in milliseconds) to display the text. If this function is called before the previous text timed out, the timer is restarted and the text is updated. If this value is zero, the text never disappears.

The function `displayOverlay` displays useful information/tips on top of the window for a certain amount of time *delayms* . This information is displayed on the window statusbar (the window must be created with the `CV_GUI_EXPANDED` flags).

## setOpenGlDrawCallback

Sets a callback function to be called to draw on top of displayed image.

**C++:** void **setOpenGlDrawCallback**(const string& **winname**, OpenGlDrawCallback **onOpenGlDraw**, void* **userdata**=0 )

**C:** void **cvSetOpenGlDrawCallback**(const char* **window_name**, CvOpenGlDrawCallback **callback**, void* **userdata**=NULL )

> **Parameters**

> > **window_name** – Name of the window.

> > **onOpenGlDraw** – Pointer to the function to be called every frame. This function should be prototyped as `void Foo(void*)` .

> > **userdata** – Pointer passed to the callback function. *(Optional)*

The function `setOpenGlDrawCallback` can be used to draw 3D data on the window. See the example of callback function below:

```
void on_opengl(void* param)
{
    glLoadIdentity();
```

```
    glTranslated(0.0, 0.0, -1.0);

    glRotatef( 55, 1, 0, 0 );
    glRotatef( 45, 0, 1, 0 );
    glRotatef( 0, 0, 0, 1 );

    static const int coords[6][4][3] = {
        { { +1, -1, -1 }, { -1, -1, -1 }, { -1, +1, -1 }, { +1, +1, -1 } },
        { { +1, +1, -1 }, { -1, +1, -1 }, { -1, +1, +1 }, { +1, +1, +1 } },
        { { +1, -1, +1 }, { +1, -1, -1 }, { +1, +1, -1 }, { +1, +1, +1 } },
        { { -1, -1, -1 }, { -1, -1, +1 }, { -1, +1, +1 }, { -1, +1, -1 } },
        { { +1, -1, +1 }, { -1, -1, +1 }, { -1, -1, -1 }, { +1, -1, -1 } },
        { { -1, -1, +1 }, { +1, -1, +1 }, { +1, +1, +1 }, { -1, +1, +1 } }
    };

    for (int i = 0; i < 6; ++i) {
            glColor3ub( i*20, 100+i*10, i*42 );
            glBegin(GL_QUADS);
            for (int j = 0; j < 4; ++j) {
                    glVertex3d(0.2 * coords[i][j][0], 0.2 * coords[i][j][1], 0.2 * coords[i][j][2]);
            }
            glEnd();
    }
}
```

## saveWindowParameters

Saves parameters of the specified window.

**C++:** void **saveWindowParameters**(const string& **windowName**)

**C:** void **cvSaveWindowParameters**(const char* **name**)

> **Parameters**
>
> > **name** – Name of the window.

The function `saveWindowParameters` saves size, location, flags, trackbars value, zoom and panning location of the window `window_name` .

## loadWindowParameters

Loads parameters of the specified window.

**C++:** void **loadWindowParameters**(const string& **windowName**)

**C:** void **cvLoadWindowParameters**(const char* **name**)

> **Parameters**
>
> > **name** – Name of the window.

The function `loadWindowParameters` loads size, location, flags, trackbars value, zoom and panning location of the window `window_name` .

## createButton

Attaches a button to the control panel.

**C++:** int **createButton**(const string& **bar_name**, ButtonCallback **on_change**, void* **userdata**=NULL, int **type**=CV_PUSH_BUTTON, bool **initial_button_state**=0 )

**C:** int **cvCreateButton**(const char* **button_name**=NULL, CvButtonCallback **on_change**=NULL, void* **userdata**=NULL, int **button_type**=CV_PUSH_BUTTON, int **initial_button_state**=0 )

>    Parameters

>>        **button_name** – Name of the button.

>>        **on_change** – Pointer to the function to be called every time the button changes its state. This function should be prototyped as void Foo(int state,*void); . *state* is the current state of the button. It could be -1 for a push button, 0 or 1 for a check/radio box button.

>>        **userdata** – Pointer passed to the callback function.

>>        **button_type** – Optional type of the button.

>>        – **CV_PUSH_BUTTON** Push button

>>        – **CV_CHECKBOX** Checkbox button

>>        – **CV_RADIOBOX** Radiobox button. The radiobox on the same buttonbar (same line) are exclusive, that is only one can be selected at a time.

>>        **initial_button_state** – Default state of the button. Use for checkbox and radiobox. Its value could be 0 or 1. *(Optional)*

The function createButton attaches a button to the control panel. Each button is added to a buttonbar to the right of the last button. A new buttonbar is created if nothing was attached to the control panel before, or if the last element attached to the control panel was a trackbar.

See below various examples of the createButton function call:

```
createButton(NULL,callbackButton);//create a push button "button 0", that will call callbackButton.
createButton("button2",callbackButton,NULL,CV_CHECKBOX,0);
createButton("button3",callbackButton,&value);
createButton("button5",callbackButton1,NULL,CV_RADIOBOX);
createButton("button6",callbackButton2,NULL,CV_PUSH_BUTTON,1);
```

# VIDEO. VIDEO ANALYSIS

## 5.1 Motion Analysis and Object Tracking

### calcOpticalFlowPyrLK

Calculates an optical flow for a sparse feature set using the iterative Lucas-Kanade method with pyramids.

**C++:** void **calcOpticalFlowPyrLK**(InputArray **prevImg**, InputArray **nextImg**, InputArray **prevPts**, InputOutputArray **nextPts**, OutputArray **status**, OutputArray **err**, Size **winSize**=Size(21,21), int **maxLevel**=3, TermCriteria **criteria**=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 0.01), int **flags**=0, double **minEigThreshold**=1e-4 )

**Python:** cv2.**calcOpticalFlowPyrLK**(prevImg, nextImg, prevPts[, nextPts[, status[, err[, winSize[, maxLevel[, criteria[, flags[, minEigThreshold ]]]]]]]]) → nextPts, status, err

**C:** void **cvCalcOpticalFlowPyrLK**(const CvArr* **prev**, const CvArr* **curr**, CvArr* **prev_pyr**, CvArr* **curr_pyr**, const CvPoint2D32f* **prev_features**, CvPoint2D32f* **curr_features**, int **count**, CvSize **win_size**, int **level**, char* **status**, float* **track_error**, CvTermCriteria **criteria**, int **flags**)

**Python:** cv.**CalcOpticalFlowPyrLK**(prev, curr, prevPyr, currPyr, prevFeatures, winSize, level, criteria, flags, guesses=None) -> (currFeatures, status, track_error)

> **Parameters**
>
> > **prevImg** – first 8-bit input image or pyramid constructed by buildOpticalFlowPyramid().
> >
> > **nextImg** – second input image or pyramid of the same size and the same type as prevImg.
> >
> > **prevPts** – vector of 2D points for which the flow needs to be found; point coordinates must be single-precision floating-point numbers.
> >
> > **nextPts** – output vector of 2D points (with single-precision floating-point coordinates) containing the calculated new positions of input features in the second image; when OPTFLOW_USE_INITIAL_FLOW flag is passed, the vector must have the same size as in the input.
> >
> > **status** – output status vector; each element of the vector is set to 1 if the flow for the corresponding features has been found, otherwise, it is set to 0.
> >
> > **err** – output vector of errors; each element of the vector is set to an error for the corresponding feature, type of the error measure can be set in flags parameter; if the flow wasn't found then the error is not defined (use the status parameter to find such cases).

> **winSize** – size of the search window at each pyramid level.
>
> **maxLevel** – 0-based maximal pyramid level number; if set to 0, pyramids are not used (single level), if set to 1, two levels are used, and so on; if pyramids are passed to input then algorithm will use as many levels as pyramids have but no more than `maxLevel`.
>
> **criteria** – parameter, specifying the termination criteria of the iterative search algorithm (after the specified maximum number of iterations `criteria.maxCount` or when the search window moves by less than `criteria.epsilon`.
>
> **flags** – operation flags:
>
> – **OPTFLOW_USE_INITIAL_FLOW** uses initial estimations, stored in `nextPts`; if the flag is not set, then `prevPts` is copied to `nextPts` and is considered the initial estimate.
>
> – **OPTFLOW_LK_GET_MIN_EIGENVALS** use minimum eigen values as an error measure (see `minEigThreshold` description); if the flag is not set, then L1 distance between patches around the original and a moved point, divided by number of pixels in a window, is used as a error measure.
>
> **minEigThreshold** – the algorithm calculates the minimum eigen value of a 2x2 normal matrix of optical flow equations (this matrix is called a spatial gradient matrix in [Bouguet00]), divided by number of pixels in a window; if this value is less than `minEigThreshold`, then a corresponding feature is filtered out and its flow is not processed, so it allows to remove bad points and get a performance boost.

The function implements a sparse iterative version of the Lucas-Kanade optical flow in pyramids. See [Bouguet00]. The function is parallelized with the TBB library.

## buildOpticalFlowPyramid

Constructs the image pyramid which can be passed to `calcOpticalFlowPyrLK()`.

**C++:** int **buildOpticalFlowPyramid**(InputArray **img**, OutputArrayOfArrays **pyramid**, Size **winSize**, int **maxLevel**, bool **withDerivatives**=true, int **pyrBorder**=BORDER_REFLECT_101, int **derivBorder**=BORDER_CONSTANT, bool **tryReuseInputImage**=true)

**Python:** cv2.**buildOpticalFlowPyramid**(img, winSize, maxLevel[, pyramid[, withDerivatives[, pyrBorder[, derivBorder[, tryReuseInputImage ]]]]]) → retval, pyramid

> **Parameters**
>
> > **img** – 8-bit input image.
> >
> > **pyramid** – output pyramid.
> >
> > **winSize** – window size of optical flow algorithm. Must be not less than `winSize` argument of `calcOpticalFlowPyrLK()`. It is needed to calculate required padding for pyramid levels.
> >
> > **maxLevel** – 0-based maximal pyramid level number.
> >
> > **withDerivatives** – set to precompute gradients for the every pyramid level. If pyramid is constructed without the gradients then `calcOpticalFlowPyrLK()` will calculate them internally.
> >
> > **pyrBorder** – the border mode for pyramid layers.
> >
> > **derivBorder** – the border mode for gradients.

**tryReuseInputImage** – put ROI of input image into the pyramid if possible. You can pass `false` to force data copying.

**Returns** number of levels in constructed pyramid. Can be less than `maxLevel`.

## calcOpticalFlowFarneback

Computes a dense optical flow using the Gunnar Farneback's algorithm.

**C++:** void **calcOpticalFlowFarneback**(InputArray **prev**, InputArray **next**, InputOutputArray **flow**, double **pyr_scale**, int **levels**, int **winsize**, int **iterations**, int **poly_n**, double **poly_sigma**, int **flags**)

**C:** void **cvCalcOpticalFlowFarneback**(const CvArr* **prev**, const CvArr* **next**, CvArr* **flow**, double **pyr_scale**, int **levels**, int **winsize**, int **iterations**, int **poly_n**, double **poly_sigma**, int **flags**)

**Python:** cv2.**calcOpticalFlowFarneback**(prev, next, pyr_scale, levels, winsize, iterations, poly_n, poly_sigma, flags[, flow ]) → flow

> **Parameters**

> > **prev** – first 8-bit single-channel input image.

> > **next** – second input image of the same size and the same type as `prev`.

> > **flow** – computed flow image that has the same size as `prev` and type `CV_32FC2`.

> > **pyr_scale** – parameter, specifying the image scale (<1) to build pyramids for each image; `pyr_scale=0.5` means a classical pyramid, where each next layer is twice smaller than the previous one.

> > **levels** – number of pyramid layers including the initial image; `levels=1` means that no extra layers are created and only the original images are used.

> > **winsize** – averaging window size; larger values increase the algorithm robustness to image noise and give more chances for fast motion detection, but yield more blurred motion field.

> > **iterations** – number of iterations the algorithm does at each pyramid level.

> > **poly_n** – size of the pixel neighborhood used to find polynomial expansion in each pixel; larger values mean that the image will be approximated with smoother surfaces, yielding more robust algorithm and more blurred motion field, typically `poly_n`=5 or 7.

> > **poly_sigma** – standard deviation of the Gaussian that is used to smooth derivatives used as a basis for the polynomial expansion; for `poly_n=5`, you can set `poly_sigma=1.1`, for `poly_n=7`, a good value would be `poly_sigma=1.5`.

> > **flags** – operation flags that can be a combination of the following:

> > – **OPTFLOW_USE_INITIAL_FLOW** uses the input `flow` as an initial flow approximation.

> > – **OPTFLOW_FARNEBACK_GAUSSIAN** uses the Gaussian `winsize` × `winsize` filter instead of a box filter of the same size for optical flow estimation; usually, this option gives z more accurate flow than with a box filter, at the cost of lower speed; normally, `winsize` for a Gaussian window should be set to a larger value to achieve the same level of robustness.

The function finds an optical flow for each `prev` pixel using the [Farneback2003] algorithm so that

$$\mathtt{prev}(y,x) \sim \mathtt{next}(y + \mathtt{flow}(y,x)[1], x + \mathtt{flow}(y,x)[0])$$

# estimateRigidTransform

Computes an optimal affine transformation between two 2D point sets.

**C++:** Mat **estimateRigidTransform**(InputArray **src**, InputArray **dst**, bool **fullAffine**)

**Python:** cv2.**estimateRigidTransform**(src, dst, fullAffine) → retval

> **Parameters**
>
>> **src** – First input 2D point set stored in std::vector or Mat, or an image stored in Mat.
>>
>> **dst** – Second input 2D point set of the same size and the same type as A, or another image.
>>
>> **fullAffine** – If true, the function finds an optimal affine transformation with no additional restrictions (6 degrees of freedom). Otherwise, the class of transformations to choose from is limited to combinations of translation, rotation, and uniform scaling (5 degrees of freedom).

The function finds an optimal affine transform *[A|b]* (a 2 x 3 floating-point matrix) that approximates best the affine transformation between:

- Two point sets

- Two raster images. In this case, the function first finds some features in the src image and finds the corresponding features in dst image. After that, the problem is reduced to the first case.

In case of point sets, the problem is formulated as follows: you need to find a 2x2 matrix *A* and 2x1 vector *b* so that:

$$[A^*|b^*] = \arg\min_{[A|b]} \sum_i \|dst[i] - A src[i]^T - b\|^2$$

> where src[i] and dst[i] are the i-th points in src and dst, respectively
>
> [A|b] can be either arbitrary (when fullAffine=true) or have a form of
>
>> $$\begin{bmatrix} a_{11} & a_{12} & b_1 \\ -a_{12} & a_{11} & b_2 \end{bmatrix}$$
>
> when fullAffine=false.

**See Also:**

getAffineTransform(), getPerspectiveTransform(), findHomography()

# updateMotionHistory

Updates the motion history image by a moving silhouette.

**C++:** void **updateMotionHistory**(InputArray **silhouette**, InputOutputArray **mhi**, double **timestamp**, double **duration**)

**Python:** cv2.**updateMotionHistory**(silhouette, mhi, timestamp, duration) → None

**C:** void **cvUpdateMotionHistory**(const CvArr* **silhouette**, CvArr* **mhi**, double **timestamp**, double **duration**)

**Python:** cv.**UpdateMotionHistory**(silhouette, mhi, timestamp, duration) → None

> **Parameters**
>
>> **silhouette** – Silhouette mask that has non-zero pixels where the motion occurs.
>>
>> **mhi** – Motion history image that is updated by the function (single-channel, 32-bit floating-point).

> **timestamp** – Current time in milliseconds or other units.

> **duration** – Maximal duration of the motion track in the same units as `timestamp` .

The function updates the motion history image as follows:

$$\text{mhi}(x, y) = \begin{cases} \text{timestamp} & \text{if } \text{silhouette}(x, y) \neq 0 \\ 0 & \text{if } \text{silhouette}(x, y) = 0 \text{ and } \text{mhi} < (\text{timestamp} - \text{duration}) \\ \text{mhi}(x, y) & \text{otherwise} \end{cases}$$

That is, MHI pixels where the motion occurs are set to the current `timestamp` , while the pixels where the motion happened last time a long time ago are cleared.

The function, together with `calcMotionGradient()` and `calcGlobalOrientation()` , implements a motion templates technique described in [Davis97] and [Bradski00]. See also the OpenCV sample `motempl.c` that demonstrates the use of all the motion template functions.

## calcMotionGradient

Calculates a gradient orientation of a motion history image.

**C++:** void **calcMotionGradient**(InputArray **mhi**, OutputArray **mask**, OutputArray **orientation**, double **delta1**, double **delta2**, int **apertureSize**=3 )

**Python:** cv2.**calcMotionGradient**(mhi, delta1, delta2[, mask[, orientation[, apertureSize ] ] ]) → mask, orientation

**C:** void **cvCalcMotionGradient**(const CvArr* **mhi**, CvArr* **mask**, CvArr* **orientation**, double **delta1**, double **delta2**, int **aperture_size**=3 )

**Python:** cv.**CalcMotionGradient**(mhi, mask, orientation, delta1, delta2, apertureSize=3) → None

> **Parameters**

> > **mhi** – Motion history single-channel floating-point image.

> > **mask** – Output mask image that has the type `CV_8UC1` and the same size as `mhi` . Its non-zero elements mark pixels where the motion gradient data is correct.

> > **orientation** – Output motion gradient orientation image that has the same type and the same size as `mhi` . Each pixel of the image is a motion orientation, from 0 to 360 degrees.

> > **delta1** – Minimal (or maximal) allowed difference between `mhi` values within a pixel neighborhood.

> > **delta2** – Maximal (or minimal) allowed difference between `mhi` values within a pixel neighborhood. That is, the function finds the minimum ( $m(x, y)$ ) and maximum ( $M(x, y)$ ) `mhi` values over $3 \times 3$ neighborhood of each pixel and marks the motion orientation at $(x, y)$ as valid only if

> > $$\min(\text{delta1}, \text{delta2}) \leq M(x, y) - m(x, y) \leq \max(\text{delta1}, \text{delta2}).$$

> > **apertureSize** – Aperture size of the `Sobel()` operator.

The function calculates a gradient orientation at each pixel $(x, y)$ as:

$$\text{orientation}(x, y) = \arctan \frac{\text{dmhi}/\text{dy}}{\text{dmhi}/\text{dx}}$$

In fact, `fastAtan2()` and `phase()` are used so that the computed angle is measured in degrees and covers the full range 0..360. Also, the `mask` is filled to indicate pixels where the computed angle is valid.

## calcGlobalOrientation

Calculates a global motion orientation in a selected region.

**C++:** double **calcGlobalOrientation**(InputArray **orientation**, InputArray **mask**, InputArray **mhi**, double **timestamp**, double **duration**)

**Python:** cv2.**calcGlobalOrientation**(orientation, mask, mhi, timestamp, duration) → retval

**C:** double **cvCalcGlobalOrientation**(const CvArr* **orientation**, const CvArr* **mask**, const CvArr* **mhi**, double **timestamp**, double **duration**)

**Python:** cv.**CalcGlobalOrientation**(orientation, mask, mhi, timestamp, duration) → float

> **Parameters**
>
> > **orientation** – Motion gradient orientation image calculated by the function `calcMotionGradient()`.
> >
> > **mask** – Mask image. It may be a conjunction of a valid gradient mask, also calculated by `calcMotionGradient()`, and the mask of a region whose direction needs to be calculated.
> >
> > **mhi** – Motion history image calculated by `updateMotionHistory()`.
> >
> > **timestamp** – Timestamp passed to `updateMotionHistory()`.
> >
> > **duration** – Maximum duration of a motion track in milliseconds, passed to `updateMotionHistory()`.

The function calculates an average motion direction in the selected region and returns the angle between 0 degrees and 360 degrees. The average direction is computed from the weighted orientation histogram, where a recent motion has a larger weight and the motion occurred in the past has a smaller weight, as recorded in `mhi`.

## segmentMotion

Splits a motion history image into a few parts corresponding to separate independent motions (for example, left hand, right hand).

**C++:** void **segmentMotion**(InputArray **mhi**, OutputArray **segmask**, vector<Rect>& **boundingRects**, double **timestamp**, double **segThresh**)

**Python:** cv2.**segmentMotion**(mhi, timestamp, segThresh[, segmask]) → segmask, boundingRects

**C:** CvSeq* **cvSegmentMotion**(const CvArr* **mhi**, CvArr* **seg_mask**, CvMemStorage* **storage**, double **timestamp**, double **seg_thresh**)

**Python:** cv.**SegmentMotion**(mhi, seg_mask, storage, timestamp, seg_thresh) → boundingRects

> **Parameters**
>
> > **mhi** – Motion history image.
> >
> > **segmask** – Image where the found mask should be stored, single-channel, 32-bit floating-point.
> >
> > **boundingRects** – Vector containing ROIs of motion connected components.
> >
> > **timestamp** – Current time in milliseconds or other units.
> >
> > **segThresh** – Segmentation threshold that is recommended to be equal to the interval between motion history "steps" or greater.

The function finds all of the motion segments and marks them in `segmask` with individual values (1,2,...). It also computes a vector with ROIs of motion connected components. After that the motion direction for every component can be calculated with `calcGlobalOrientation()` using the extracted mask of the particular component.

## CamShift

Finds an object center, size, and orientation.

**C++:** RotatedRect **CamShift**(InputArray **probImage**, Rect& **window**, TermCriteria **criteria**)

**Python:** cv2.**CamShift**(probImage, window, criteria) → retval, window

**C:** int **cvCamShift**(const CvArr* **prob_image**, CvRect **window**, CvTermCriteria **criteria**, CvConnected-Comp* **comp**, CvBox2D* **box**=NULL )

**Python:** cv.**CamShift**(prob_image, window, criteria) -> (int, comp, box)

> **Parameters**
>
>> **probImage** – Back projection of the object histogram. See `calcBackProject()` .
>>
>> **window** – Initial search window.
>>
>> **criteria** – Stop criteria for the underlying `meanShift()` .
>
> **Returns** (in old interfaces) Number of iterations CAMSHIFT took to converge

The function implements the CAMSHIFT object tracking algorithm [Bradski98]. First, it finds an object center using `meanShift()` and then adjusts the window size and finds the optimal rotation. The function returns the rotated rectangle structure that includes the object position, size, and orientation. The next position of the search window can be obtained with `RotatedRect::boundingRect()` .

See the OpenCV sample `camshiftdemo.c` that tracks colored objects.

## meanShift

Finds an object on a back projection image.

**C++:** int **meanShift**(InputArray **probImage**, Rect& **window**, TermCriteria **criteria**)

**Python:** cv2.**meanShift**(probImage, window, criteria) → retval, window

**C:** int **cvMeanShift**(const CvArr* **prob_image**, CvRect **window**, CvTermCriteria **criteria**, CvConnected-Comp* **comp**)

**Python:** cv.**MeanShift**(prob_image, window, criteria) → comp

> **Parameters**
>
>> **probImage** – Back projection of the object histogram. See `calcBackProject()` for details.
>>
>> **window** – Initial search window.
>>
>> **criteria** – Stop criteria for the iterative search algorithm.
>
> **Returns** Number of iterations CAMSHIFT took to converge.

The function implements the iterative object search algorithm. It takes the input back projection of an object and the initial position. The mass center in `window` of the back projection image is computed and the search window center shifts to the mass center. The procedure is repeated until the specified number of iterations `criteria.maxCount` is done or until the window center shifts by less than `criteria.epsilon` . The algorithm is used inside `CamShift()` and, unlike `CamShift()` , the search window size or orientation do not change during the search. You can simply pass the output of `calcBackProject()` to this function. But better results can be obtained if you pre-filter the back projection and remove the noise. For example, you can do this by retrieving connected components with `findContours()` , throwing away contours with small area ( `contourArea()` ), and rendering the remaining contours with `drawContours()` .

# KalmanFilter

class **KalmanFilter**

Kalman filter class.

The class implements a standard Kalman filter http://en.wikipedia.org/wiki/Kalman_filter, [Welch95]. However, you can modify `transitionMatrix`, `controlMatrix`, and `measurementMatrix` to get an extended Kalman filter functionality. See the OpenCV sample `kalman.cpp` .

## KalmanFilter::KalmanFilter

The constructors.

**C++:** KalmanFilter::**KalmanFilter**()

**C++:** KalmanFilter::**KalmanFilter**(int **dynamParams**, int **measureParams**, int **controlParams**=0, int **type**=CV_32F)

**Python:** cv2.**KalmanFilter**([dynamParams, measureParams[, controlParams[, type ] ] ]) → <KalmanFilter object>

**C:** CvKalman* **cvCreateKalman**(int **dynam_params**, int **measure_params**, int **control_params**=0 )

**Python:** cv.**CreateKalman**(dynam_params, measure_params, control_params=0) → CvKalman

The full constructor.

> **Parameters**
>
> > **dynamParams** – Dimensionality of the state.
> >
> > **measureParams** – Dimensionality of the measurement.
> >
> > **controlParams** – Dimensionality of the control vector.
> >
> > **type** – Type of the created matrices that should be `CV_32F` or `CV_64F`.

**Note:** In C API when CvKalman* kalmanFilter structure is not needed anymore, it should be released with cvReleaseKalman(&kalmanFilter)

## KalmanFilter::init

Re-initializes Kalman filter. The previous content is destroyed.

**C++:** void KalmanFilter::**init**(int **dynamParams**, int **measureParams**, int **controlParams**=0, int **type**=CV_32F)

> **Parameters**
>
> > **dynamParams** – Dimensionalityensionality of the state.
> >
> > **measureParams** – Dimensionality of the measurement.
> >
> > **controlParams** – Dimensionality of the control vector.
> >
> > **type** – Type of the created matrices that should be `CV_32F` or `CV_64F`.

## KalmanFilter::predict

Computes a predicted state.

**C++:** const Mat& KalmanFilter::**predict**(const Mat& **control**=Mat())

**Python:** cv2.KalmanFilter.**predict**([control]) → retval

**C:** const CvMat* **cvKalmanPredict**(CvKalman* **kalman**, const CvMat* **control**=NULL)

**Python:** cv.**KalmanPredict**(kalman, control=None) → mat

> **Parameters**
>
> > **control** – The optional input control

## KalmanFilter::correct

Updates the predicted state from the measurement.

**C++:** const Mat& KalmanFilter::**correct**(const Mat& **measurement**)

**Python:** cv2.KalmanFilter.**correct**(measurement) → retval

**C:** const CvMat* **cvKalmanCorrect**(CvKalman* **kalman**, const CvMat* **measurement**)

**Python:** cv.**KalmanCorrect**(kalman, measurement) → mat

> **Parameters**
>
> > **measurement** – The measured system parameters

## BackgroundSubtractor

class **BackgroundSubtractor** : **public** **Algorithm**

Base class for background/foreground segmentation.

```cpp
class BackgroundSubtractor : public Algorithm
{
public:
    virtual ~BackgroundSubtractor();
    virtual void operator()(InputArray image, OutputArray fgmask, double learningRate=0);
    virtual void getBackgroundImage(OutputArray backgroundImage) const;
};
```

The class is only used to define the common interface for the whole family of background/foreground segmentation algorithms.

## BackgroundSubtractor::operator()

Computes a foreground mask.

**C++:** void BackgroundSubtractor::**operator()**(InputArray **image**, OutputArray **fgmask**, double **learningRate**=0)

**Python:** cv2.BackgroundSubtractor.**apply**(image[, fgmask[, learningRate]]) → fgmask

> **Parameters**
>
> > **image** – Next video frame.

**fgmask** – The output foreground mask as an 8-bit binary image.

## BackgroundSubtractor::getBackgroundImage

Computes a background image.

**C++:** void BackgroundSubtractor::**getBackgroundImage**(OutputArray **backgroundImage**) const

> **Parameters**

> > **backgroundImage** – The output background image.

**Note:** Sometimes the background image can be very blurry, as it contain the average background statistics.

## BackgroundSubtractorMOG

class **BackgroundSubtractorMOG** : **public BackgroundSubtractor**

Gaussian Mixture-based Background/Foreground Segmentation Algorithm.

The class implements the algorithm described in P. KadewTraKuPong and R. Bowden, *An improved adaptive background mixture model for real-time tracking with shadow detection*, Proc. 2nd European Workshop on Advanced Video-Based Surveillance Systems, 2001: http://personal.ee.surrey.ac.uk/Personal/R.Bowden/publications/avbs01/avbs01.pdf

## BackgroundSubtractorMOG::BackgroundSubtractorMOG

The constructors.

**C++:** BackgroundSubtractorMOG::**BackgroundSubtractorMOG**()

**C++:** BackgroundSubtractorMOG::**BackgroundSubtractorMOG**(int **history**, int **nmixtures**, double **backgroundRatio**, double **noiseSigma**=0)

**Python:** cv2.**BackgroundSubtractorMOG**([history, nmixtures, backgroundRatio[, noiseSigma]]) → <BackgroundSubtractorMOG object>

> **Parameters**

> > **history** – Length of the history.

> > **nmixtures** – Number of Gaussian mixtures.

> > **backgroundRatio** – Background ratio.

> > **noiseSigma** – Noise strength.

Default constructor sets all parameters to default values.

## BackgroundSubtractorMOG::operator()

Updates the background model and returns the foreground mask

**C++:** void BackgroundSubtractorMOG::**operator()**(InputArray **image**, OutputArray **fgmask**, double **learningRate**=0)

Parameters are the same as in BackgroundSubtractor::operator()

## BackgroundSubtractorMOG2

Gaussian Mixture-based Background/Foreground Segmentation Algorithm.

**class BackgroundSubtractorMOG2** : **public BackgroundSubtractor**

Here are important members of the class that control the algorithm, which you can set after constructing the class instance:

int **nmixtures**

Maximum allowed number of mixture components. Actual number is determined dynamically per pixel.

float **backgroundRatio**

Threshold defining whether the component is significant enough to be included into the background model ( corresponds to TB=1-cf from the paper??which paper??). cf=0.1 => TB=0.9 is default. For alpha=0.001, it means that the mode should exist for approximately 105 frames before it is considered foreground.

float **varThresholdGen**

Threshold for the squared Mahalanobis distance that helps decide when a sample is close to the existing components (corresponds to Tg). If it is not close to any component, a new component is generated. 3 sigma => Tg=3*3=9 is default. A smaller Tg value generates more components. A higher Tg value may result in a small number of components but they can grow too large.

float **fVarInit**

Initial variance for the newly generated components. It affects the speed of adaptation. The parameter value is based on your estimate of the typical standard deviation from the images. OpenCV uses 15 as a reasonable value.

float **fVarMin**

Parameter used to further control the variance.

float **fVarMax**

Parameter used to further control the variance.

float **fCT**

Complexity reduction parameter. This parameter defines the number of samples needed to accept to prove the component exists. CT=0.05 is a default value for all the samples. By setting CT=0 you get an algorithm very similar to the standard Stauffer&Grimson algorithm.

uchar **nShadowDetection**

The value for marking shadow pixels in the output foreground mask. Default value is 127.

float **fTau**

Shadow threshold. The shadow is detected if the pixel is a darker version of the background. Tau is a threshold defining how much darker the shadow can be. Tau= 0.5 means that if a pixel is more than twice darker then it is not shadow. See Prati,Mikic,Trivedi,Cucchiarra, *Detecting Moving Shadows...*, IEEE PAMI,2003.

The class implements the Gaussian mixture model background subtraction described in:

- Z.Zivkovic, *Improved adaptive Gausian mixture model for background subtraction*, International Conference Pattern Recognition, UK, August, 2004, http://www.zoranz.net/Publications/zivkovic2004ICPR.pdf. The code is very fast and performs also shadow detection. Number of Gausssian components is adapted per pixel.

- Z.Zivkovic, F. van der Heijden, *Efficient Adaptive Density Estimapion per Image Pixel for the Task of Background Subtraction*, Pattern Recognition Letters, vol. 27, no. 7, pages 773-780, 2006. The algorithm similar to the standard Stauffer&Grimson algorithm with additional selection of the number of the Gaussian components based on: Z.Zivkovic, F.van der Heijden, Recursive unsupervised learning of finite mixture models, IEEE Trans. on Pattern Analysis and Machine Intelligence, vol.26, no.5, pages 651-656, 2004.

## BackgroundSubtractorMOG2::BackgroundSubtractorMOG2

The constructors.

**C++:** BackgroundSubtractorMOG2::**BackgroundSubtractorMOG2**()

**C++:** BackgroundSubtractorMOG2::**BackgroundSubtractorMOG2**(int **history**, float **varThreshold**, bool **bShadowDetection**=true )

> **Parameters**
>
>> **history** – Length of the history.
>>
>> **varThreshold** – Threshold on the squared Mahalanobis distance to decide whether it is well described by the background model (see Cthr??). This parameter does not affect the background update. A typical value could be 4 sigma, that is, varThreshold=4∗4=16; (see Tb??).
>>
>> **bShadowDetection** – Parameter defining whether shadow detection should be enabled (true or false).

## BackgroundSubtractorMOG2::operator()

Updates the background model and computes the foreground mask

**C++:** void BackgroundSubtractorMOG2::**operator()**(InputArray **image**, OutputArray **fgmask**, double **learningRate**=-1)

> See BackgroundSubtractor::operator().

## BackgroundSubtractorMOG2::getBackgroundImage

Returns background image

**C++:** void BackgroundSubtractorMOG2::**getBackgroundImage**(OutputArray **backgroundImage**)

See BackgroundSubtractor::getBackgroundImage().

## calcOpticalFlowSF

Calculate an optical flow using "SimpleFlow" algorithm.

**C++:** void **calcOpticalFlowSF**(Mat& **from**, Mat& **to**, Mat& **flow**, int **layers**, int **averaging_block_size**, int **max_flow**)

**C++:** void **calcOpticalFlowSF**(Mat& **from**, Mat& **to**, Mat& **flow**, int **layers**, int **averaging_block_size**, int **max_flow**, double **sigma_dist**, double **sigma_color**, int **postprocess_window**, double **sigma_dist_fix**, double **sigma_color_fix**, double **occ_thr**, int **upscale_averaging_radius**, double **upscale_sigma_dist**, double **upscale_sigma_color**, double **speed_up_thr**)

> **Parameters**
>
>> **prev** – First 8-bit 3-channel image.
>>
>> **next** – Second 8-bit 3-channel image
>>
>> **flowX** – X-coordinate of estimated flow
>>
>> **flowY** – Y-coordinate of estimated flow
>>
>> **layers** – Number of layers

> **averaging_block_size** – Size of block through which we sum up when calculate cost function for pixel
>
> **max_flow** – maximal flow that we search at each level
>
> **sigma_dist** – vector smooth spatial sigma parameter
>
> **sigma_color** – vector smooth color sigma parameter
>
> **postprocess_window** – window size for postprocess cross bilateral filter
>
> **sigma_dist_fix** – spatial sigma for postprocess cross bilateralf filter
>
> **sigma_color_fix** – color sigma for postprocess cross bilateral filter
>
> **occ_thr** – threshold for detecting occlusions
>
> **upscale_averaging_radiud** – window size for bilateral upscale operation
>
> **upscale_sigma_dist** – spatial sigma for bilateral upscale operation
>
> **upscale_sigma_color** – color sigma for bilateral upscale operation
>
> **speed_up_thr** – threshold to detect point with irregular flow - where flow should be recalculated after upscale

See [Tao2012]. And site of project - http://graphics.berkeley.edu/papers/Tao-SAN-2012-05/.

# CALIB3D. CAMERA CALIBRATION AND 3D RECONSTRUCTION

## 6.1 Camera Calibration and 3D Reconstruction

The functions in this section use a so-called pinhole camera model. In this model, a scene view is formed by projecting 3D points into the image plane using a perspective transformation.

$$s\, m' = A[R|t]M'$$

or

$$s \begin{bmatrix} u \\ v \\ 1 \end{bmatrix} = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} r_{11} & r_{12} & r_{13} & t_1 \\ r_{21} & r_{22} & r_{23} & t_2 \\ r_{31} & r_{32} & r_{33} & t_3 \end{bmatrix} \begin{bmatrix} X \\ Y \\ Z \\ 1 \end{bmatrix}$$

where:

- $(X, Y, Z)$ are the coordinates of a 3D point in the world coordinate space
- $(u, v)$ are the coordinates of the projection point in pixels
- $A$ is a camera matrix, or a matrix of intrinsic parameters
- $(cx, cy)$ is a principal point that is usually at the image center
- $fx, fy$ are the focal lengths expressed in pixel units.

Thus, if an image from the camera is scaled by a factor, all of these parameters should be scaled (multiplied/divided, respectively) by the same factor. The matrix of intrinsic parameters does not depend on the scene viewed. So, once estimated, it can be re-used as long as the focal length is fixed (in case of zoom lens). The joint rotation-translation matrix $[R|t]$ is called a matrix of extrinsic parameters. It is used to describe the camera motion around a static scene, or vice versa, rigid motion of an object in front of a still camera. That is, $[R|t]$ translates coordinates of a point $(X, Y, Z)$ to a coordinate system, fixed with respect to the camera. The transformation above is equivalent to the following (when $z \neq 0$):

$$\begin{bmatrix} x \\ y \\ z \end{bmatrix} = R \begin{bmatrix} X \\ Y \\ Z \end{bmatrix} + t$$

$$x' = x/z$$
$$y' = y/z$$
$$u = f_x * x' + c_x$$
$$v = f_y * y' + c_y$$

Real lenses usually have some distortion, mostly radial distortion and slight tangential distortion. So, the above model is extended as:

$$
\begin{bmatrix} x \\ y \\ z \end{bmatrix} = R \begin{bmatrix} X \\ Y \\ Z \end{bmatrix} + t
$$
$$
x' = x/z
$$
$$
y' = y/z
$$
$$
x'' = x' \frac{1 + k_1 r^2 + k_2 r^4 + k_3 r^6}{1 + k_4 r^2 + k_5 r^4 + k_6 r^6} + 2p_1 x' y' + p_2 (r^2 + 2x'^2)
$$
$$
y'' = y' \frac{1 + k_1 r^2 + k_2 r^4 + k_3 r^6}{1 + k_4 r^2 + k_5 r^4 + k_6 r^6} + p_1 (r^2 + 2y'^2) + 2p_2 x' y'
$$
$$
\text{where} \quad r^2 = x'^2 + y'^2
$$
$$
u = f_x * x'' + c_x
$$
$$
v = f_y * y'' + c_y
$$

$k_1$, $k_2$, $k_3$, $k_4$, $k_5$, and $k_6$ are radial distortion coefficients. $p_1$ and $p_2$ are tangential distortion coefficients. Higher-order coefficients are not considered in OpenCV. In the functions below the coefficients are passed or returned as

$$
(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])
$$

vector. That is, if the vector contains four elements, it means that $k_3 = 0$. The distortion coefficients do not depend on the scene viewed. Thus, they also belong to the intrinsic camera parameters. And they remain the same regardless of the captured image resolution. If, for example, a camera has been calibrated on images of 320 x 240 resolution, absolutely the same distortion coefficients can be used for 640 x 480 images from the same camera while $f_x$, $f_y$, $c_x$, and $c_y$ need to be scaled appropriately.

The functions below use the above model to do the following:

- Project 3D points to the image plane given intrinsic and extrinsic parameters.

- Compute extrinsic parameters given intrinsic parameters, a few 3D points, and their projections.

- Estimate intrinsic and extrinsic camera parameters from several views of a known calibration pattern (every view is described by several 3D-2D point correspondences).

- Estimate the relative position and orientation of the stereo camera "heads" and compute the *rectification* transformation that makes the camera optical axes parallel.

## calibrateCamera

Finds the camera intrinsic and extrinsic parameters from several views of a calibration pattern.

**C++:** double **calibrateCamera**(InputArrayOfArrays **objectPoints**, InputArrayOfArrays **imagePoints**, Size **imageSize**, InputOutputArray **cameraMatrix**, InputOutputArray **distCoeffs**, OutputArrayOfArrays **rvecs**, OutputArrayOfArrays **tvecs**, int **flags**=0, TermCriteria **criteria**=TermCriteria( TermCriteria::COUNT+TermCriteria::EPS, 30, DBL_EPSILON) )

**Python:** cv2.**calibrateCamera**(objectPoints, imagePoints, imageSize[, cameraMatrix[, distCoeffs[, rvecs[, tvecs[, flags[, criteria ]]]]]]) → retval, cameraMatrix, distCoeffs, rvecs, tvecs

**C:** double **cvCalibrateCamera2**(const CvMat* **object_points**, const CvMat* **image_points**, const CvMat* **point_counts**, CvSize **image_size**, CvMat* **camera_matrix**, CvMat* **distortion_coeffs**, CvMat* **rotation_vectors**=NULL, CvMat* **translation_vectors**=NULL, int **flags**=0, CvTermCriteria **term_crit**=cvTermCriteria( CV_TERMCRIT_ITER+CV_TERMCRIT_EPS,30,DBL_EPSILON) )

**Python:** cv.**CalibrateCamera2**(objectPoints, imagePoints, pointCounts, imageSize, cameraMatrix, distCo-
effs, rvecs, tvecs, flags=0) → None

**Parameters**

**objectPoints** – In the new interface it is a vector of vectors of calibration pattern points in
the calibration pattern coordinate space. The outer vector contains as many elements as the
number of the pattern views. If the same calibration pattern is shown in each view and it
is fully visible, all the vectors will be the same. Although, it is possible to use partially
occluded patterns, or even different patterns in different views. Then, the vectors will be
different. The points are 3D, but since they are in a pattern coordinate system, then, if the rig
is planar, it may make sense to put the model to a XY coordinate plane so that Z-coordinate
of each input object point is 0.

In the old interface all the vectors of object points from different views are concatenated
together.

**imagePoints** – In the new interface it is a vector of vectors of the projections
of calibration pattern points. `imagePoints.size()` and `objectPoints.size()` and
`imagePoints[i].size()` must be equal to `objectPoints[i].size()` for each i.

In the old interface all the vectors of object points from different views are concatenated
together.

**point_counts** – In the old interface this is a vector of integers, containing as many elements,
as the number of views of the calibration pattern. Each element is the number of points in
each view. Usually, all the elements are the same and equal to the number of feature points
on the calibration pattern.

**imageSize** – Size of the image used only to initialize the intrinsic camera matrix.

**cameraMatrix** – Output 3x3 floating-point camera matrix $A = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ . If
`CV_CALIB_USE_INTRINSIC_GUESS` and/or `CV_CALIB_FIX_ASPECT_RATIO` are specified,
some or all of `fx, fy, cx, cy` must be initialized before calling the function.

**distCoeffs** – Output vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4,
5, or 8 elements.

**rvecs** – Output vector of rotation vectors (see `Rodrigues()` ) estimated for each pattern
view. That is, each k-th rotation vector together with the corresponding k-th translation
vector (see the next output parameter description) brings the calibration pattern from the
model coordinate space (in which object points are specified) to the world coordinate space,
that is, a real position of the calibration pattern in the k-th pattern view (k=0.. $M$ -1).

**tvecs** – Output vector of translation vectors estimated for each pattern view.

**flags** – Different flags that may be zero or a combination of the following values:

– **CV_CALIB_USE_INTRINSIC_GUESS** `cameraMatrix` contains valid initial values of
`fx, fy, cx, cy` that are optimized further. Otherwise, (`cx, cy`) is initially set to the
image center ( `imageSize` is used), and focal distances are computed in a least-squares
fashion. Note, that if intrinsic parameters are known, there is no need to use this function
just to estimate extrinsic parameters. Use `solvePnP()` instead.

– **CV_CALIB_FIX_PRINCIPAL_POINT** The principal point is not changed during the
global optimization. It stays at the center or at a different location specified when
`CV_CALIB_USE_INTRINSIC_GUESS` is set too.

– **CV_CALIB_FIX_ASPECT_RATIO** The functions considers only `fy` as a free pa-
rameter. The ratio `fx/fy` stays the same as in the input `cameraMatrix` . When

> CV_CALIB_USE_INTRINSIC_GUESS is not set, the actual input values of fx and fy are
> ignored, only their ratio is computed and used further.

> – **CV_CALIB_ZERO_TANGENT_DIST** Tangential distortion coefficients $(p_1, p_2)$ are
> set to zeros and stay zero.

> – **CV_CALIB_FIX_K1,...,CV_CALIB_FIX_K6** The corresponding radial distortion co-
> efficient is not changed during the optimization. If CV_CALIB_USE_INTRINSIC_GUESS is
> set, the coefficient from the supplied distCoeffs matrix is used. Otherwise, it is set to
> 0.

> – **CV_CALIB_RATIONAL_MODEL** Coefficients k4, k5, and k6 are enabled. To pro-
> vide the backward compatibility, this extra flag should be explicitly specified to make the
> calibration function use the rational model and return 8 coefficients. If the flag is not set,
> the function computes and returns only 5 distortion coefficients.

> **criteria** – Termination criteria for the iterative optimization algorithm.

> **term_crit** – same as criteria.

The function estimates the intrinsic camera parameters and extrinsic parameters for each of the views. The algorithm
is based on [Zhang2000] and [BouguetMCT]. The coordinates of 3D object points and their corresponding 2D pro-
jections in each view must be specified. That may be achieved by using an object with a known geometry and easily
detectable feature points. Such an object is called a calibration rig or calibration pattern, and OpenCV has built-in
support for a chessboard as a calibration rig (see findChessboardCorners() ). Currently, initialization of intrinsic
parameters (when CV_CALIB_USE_INTRINSIC_GUESS is not set) is only implemented for planar calibration patterns
(where Z-coordinates of the object points must be all zeros). 3D calibration rigs can also be used as long as initial
cameraMatrix is provided.

The algorithm performs the following steps:

1. Compute the initial intrinsic parameters (the option only available for planar calibration patterns) or read
   them from the input parameters. The distortion coefficients are all set to zeros initially unless some of
   CV_CALIB_FIX_K? are specified.

2. Estimate the initial camera pose as if the intrinsic parameters have been already known. This is done using
   solvePnP() .

3. Run the global Levenberg-Marquardt optimization algorithm to minimize the reprojection error, that is, the
   total sum of squared distances between the observed feature points imagePoints and the projected (using the
   current estimates for camera parameters and the poses) object points objectPoints. See projectPoints()
   for details.

The function returns the final re-projection error.

---

**Note:** If you use a non-square (=non-NxN) grid and findChessboardCorners() for calibration,
and calibrateCamera returns bad values (zero distortion coefficients, an image center very far from
(w/2-0.5,h/2-0.5), and/or large differences between $f_x$ and $f_y$ (ratios of 10:1 or more)), then you
have probably used patternSize=cvSize(rows,cols) instead of using patternSize=cvSize(cols,rows) in
findChessboardCorners() .

---

**See Also:**

findChessboardCorners(), solvePnP(), initCameraMatrix2D(), stereoCalibrate(), undistort()

## calibrationMatrixValues

Computes useful camera characteristics from the camera matrix.

**C++:** void **calibrationMatrixValues**(InputArray **cameraMatrix**, Size **imageSize**, double **apertureWidth**, double **apertureHeight**, double& **fovx**, double& **fovy**, double& **focalLength**, Point2d& **principalPoint**, double& **aspectRatio**)

**Python:** cv2.**calibrationMatrixValues**(cameraMatrix, imageSize, apertureWidth, apertureHeight) → fovx, fovy, focalLength, principalPoint, aspectRatio

> Parameters
>
> > **cameraMatrix** – Input camera matrix that can be estimated by `calibrateCamera()` or `stereoCalibrate()` .
> >
> > **imageSize** – Input image size in pixels.
> >
> > **apertureWidth** – Physical width of the sensor.
> >
> > **apertureHeight** – Physical height of the sensor.
> >
> > **fovx** – Output field of view in degrees along the horizontal sensor axis.
> >
> > **fovy** – Output field of view in degrees along the vertical sensor axis.
> >
> > **focalLength** – Focal length of the lens in mm.
> >
> > **principalPoint** – Principal point in pixels.
> >
> > **aspectRatio** – $f_y/f_x$

The function computes various useful camera characteristics from the previously estimated camera matrix.

## composeRT

Combines two rotation-and-shift transformations.

**C++:** void **composeRT**(InputArray **rvec1**, InputArray **tvec1**, InputArray **rvec2**, InputArray **tvec2**, OutputArray **rvec3**, OutputArray **tvec3**, OutputArray **dr3dr1**=noArray(), OutputArray **dr3dt1**=noArray(), OutputArray **dr3dr2**=noArray(), OutputArray **dr3dt2**=noArray(), OutputArray **dt3dr1**=noArray(), OutputArray **dt3dt1**=noArray(), OutputArray **dt3dr2**=noArray(), OutputArray **dt3dt2**=noArray() )

**Python:** cv2.**composeRT**(rvec1, tvec1, rvec2, tvec2[, rvec3[, tvec3[, dr3dr1[, dr3dt1[, dr3dr2[, dr3dt2[, dt3dr1[, dt3dt1[, dt3dr2[, dt3dt2 ]]]]]]]]]]) → rvec3, tvec3, dr3dr1, dr3dt1, dr3dr2, dr3dt2, dt3dr1, dt3dt1, dt3dr2, dt3dt2

> Parameters
>
> > **rvec1** – First rotation vector.
> >
> > **tvec1** – First translation vector.
> >
> > **rvec2** – Second rotation vector.
> >
> > **tvec2** – Second translation vector.
> >
> > **rvec3** – Output rotation vector of the superposition.
> >
> > **tvec3** – Output translation vector of the superposition.
> >
> > **d\*d\*** – Optional output derivatives of `rvec3` or `tvec3` with regard to `rvec1`, `rvec2`, `tvec1` and `tvec2`, respectively.

The functions compute:

$$\begin{aligned} \texttt{rvec3} &= \mathrm{rodrigues}^{-1}\left(\mathrm{rodrigues}(\texttt{rvec2}) \cdot \mathrm{rodrigues}(\texttt{rvec1})\right) \\ \texttt{tvec3} &= \mathrm{rodrigues}(\texttt{rvec2}) \cdot \texttt{tvec1} + \texttt{tvec2} \end{aligned},$$

where rodrigues denotes a rotation vector to a rotation matrix transformation, and rodrigues$^{-1}$ denotes the inverse transformation. See `Rodrigues()` for details.

Also, the functions can compute the derivatives of the output vectors with regards to the input vectors (see `matMulDeriv()` ). The functions are used inside `stereoCalibrate()` but can also be used in your own code where Levenberg-Marquardt or another gradient-based solver is used to optimize a function that contains a matrix multiplication.

## computeCorrespondEpilines

For points in an image of a stereo pair, computes the corresponding epilines in the other image.

**C++:** void **computeCorrespondEpilines**(InputArray **points**, int **whichImage**, InputArray **F**, OutputArray **lines**)

**C:** void **cvComputeCorrespondEpilines**(const CvMat\* **points**, int **which_image**, const CvMat\* **fundamental_matrix**, CvMat\* **correspondent_lines**)

**Python:** cv.**ComputeCorrespondEpilines**(points, whichImage, F, lines) → None

> **Parameters**
>
> > **points** – Input points. $N \times 1$ or $1 \times N$ matrix of type `CV_32FC2` or `vector<Point2f>` .
> >
> > **whichImage** – Index of the image (1 or 2) that contains the `points` .
> >
> > **F** – Fundamental matrix that can be estimated using `findFundamentalMat()` or `stereoRectify()` .
> >
> > **lines** – Output vector of the epipolar lines corresponding to the points in the other image. Each line $ax + by + c = 0$ is encoded by 3 numbers $(a, b, c)$ .

For every point in one of the two images of a stereo pair, the function finds the equation of the corresponding epipolar line in the other image.

From the fundamental matrix definition (see `findFundamentalMat()` ), line $l_i^{(2)}$ in the second image for the point $p_i^{(1)}$ in the first image (when `whichImage=1` ) is computed as:

$$l_i^{(2)} = F p_i^{(1)}$$

And vice versa, when `whichImage=2`, $l_i^{(1)}$ is computed from $p_i^{(2)}$ as:

$$l_i^{(1)} = F^T p_i^{(2)}$$

Line coefficients are defined up to a scale. They are normalized so that $a_i^2 + b_i^2 = 1$ .

## convertPointsToHomogeneous

Converts points from Euclidean to homogeneous space.

**C++:** void **convertPointsToHomogeneous**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**convertPointsToHomogeneous**(src[, dst]) → dst

> **Parameters**

>**src** – Input vector of `N`-dimensional points.

>**dst** – Output vector of `N+1`-dimensional points.

The function converts points from Euclidean to homogeneous space by appending 1's to the tuple of point coordinates. That is, each point (`x1, x2, ..., xn`) is converted to (`x1, x2, ..., xn, 1`).

## convertPointsFromHomogeneous

Converts points from homogeneous to Euclidean space.

**C++:** void **convertPointsFromHomogeneous**(InputArray **src**, OutputArray **dst**)

**Python:** cv2.**convertPointsFromHomogeneous**(src[, dst]) → dst

>**Parameters**

>>**src** – Input vector of `N`-dimensional points.

>>**dst** – Output vector of `N-1`-dimensional points.

The function converts points homogeneous to Euclidean space using perspective projection. That is, each point (`x1, x2, ...  x(n-1), xn`) is converted to (`x1/xn, x2/xn, ..., x(n-1)/xn`). When xn=0, the output point coordinates will be (`0,0,0,...`).

## convertPointsHomogeneous

Converts points to/from homogeneous coordinates.

**C++:** void **convertPointsHomogeneous**(InputArray **src**, OutputArray **dst**)

**C:** void **cvConvertPointsHomogeneous**(const CvMat* **src**, CvMat* **dst**)

**Python:** cv.**ConvertPointsHomogeneous**(src, dst) → None

>**Parameters**

>>**src** – Input array or vector of 2D, 3D, or 4D points.

>>**dst** – Output vector of 2D, 3D, or 4D points.

The function converts 2D or 3D points from/to homogeneous coordinates by calling either `convertPointsToHomogeneous()` or `convertPointsFromHomogeneous()`.

**Note:** The function is obsolete. Use one of the previous two functions instead.

## correctMatches

Refines coordinates of corresponding points.

**C++:** void **correctMatches**(InputArray **F**, InputArray **points1**, InputArray **points2**, OutputArray **newPoints1**, OutputArray **newPoints2**)

**Python:** cv2.**correctMatches**(F, points1, points2[, newPoints1[, newPoints2]]) → newPoints1, newPoints2

**C:** void **cvCorrectMatches**(CvMat* **F**, CvMat* **points1**, CvMat* **points2**, CvMat* **new_points1**, CvMat* **new_points2**)

>**Parameters**

> **F** – 3x3 fundamental matrix.
>
> **points1** – 1xN array containing the first set of points.
>
> **points2** – 1xN array containing the second set of points.
>
> **newPoints1** – The optimized points1.
>
> **newPoints2** – The optimized points2.

The function implements the Optimal Triangulation Method (see Multiple View Geometry for details). For each given point correspondence points1[i] <-> points2[i], and a fundamental matrix F, it computes the corrected correspondences newPoints1[i] <-> newPoints2[i] that minimize the geometric error $d(\mathrm{points1[i]}, \mathrm{newPoints1[i]})^2 + d(\mathrm{points2[i]}, \mathrm{newPoints2[i]})^2$ (where $d(a, b)$ is the geometric distance between points $a$ and $b$ ) subject to the epipolar constraint $\mathrm{newPoints2}^\mathsf{T} * F * \mathrm{newPoints1} = 0$ .

## decomposeProjectionMatrix

Decomposes a projection matrix into a rotation matrix and a camera matrix.

**C++:** void **decomposeProjectionMatrix**(InputArray **projMatrix**, OutputArray **cameraMatrix**, OutputArray **rotMatrix**, OutputArray **transVect**, OutputArray **rotMatrixX**=noArray(), OutputArray **rotMatrixY**=noArray(), OutputArray **rotMatrixZ**=noArray(), OutputArray **eulerAngles**=noArray() )

**Python:** cv2.**decomposeProjectionMatrix**(projMatrix[, cameraMatrix[, rotMatrix[, transVect[, rotMatrixX[, rotMatrixY[, rotMatrixZ[, eulerAngles ]]]]]]]) → cameraMatrix, rotMatrix, transVect, rotMatrixX, rotMatrixY, rotMatrixZ, eulerAngles

**C:** void **cvDecomposeProjectionMatrix**(const CvMat* **projMatr**, CvMat* **calibMatr**, CvMat* **rotMatr**, CvMat* **posVect**, CvMat* **rotMatrX**=NULL, CvMat* **rotMatrY**=NULL, CvMat* **rotMatrZ**=NULL, CvPoint3D64f* **eulerAngles**=NULL )

**Python:** cv.**DecomposeProjectionMatrix**(projMatr, cameraMatrix, rotMatrix, transVect, rotMatrX=None, rotMatrY=None, rotMatrZ=None) → eulerAngles

> **Parameters**
>
> > **projMatrix** – 3x4 input projection matrix P.
> >
> > **cameraMatrix** – Output 3x3 camera matrix K.
> >
> > **rotMatrix** – Output 3x3 external rotation matrix R.
> >
> > **transVect** – Output 4x1 translation vector T.
> >
> > **rotMatrX** – Optional 3x3 rotation matrix around x-axis.
> >
> > **rotMatrY** – Optional 3x3 rotation matrix around y-axis.
> >
> > **rotMatrZ** – Optional 3x3 rotation matrix around z-axis.
> >
> > **eulerAngles** – Optional three-element vector containing three Euler angles of rotation in degrees.

The function computes a decomposition of a projection matrix into a calibration and a rotation matrix and the position of a camera.

It optionally returns three rotation matrices, one for each axis, and three Euler angles that could be used in OpenGL. Note, there is always more than one sequence of rotations about the three principle axes that results in the same

orientation of an object, eg. see [Slabaugh]. Returned tree rotation matrices and corresponding three Euler angules are only one of the possible solutions.

The function is based on `RQDecomp3x3()` .

## drawChessboardCorners

Renders the detected chessboard corners.

**C++:** void **drawChessboardCorners**(InputOutputArray **image**, Size **patternSize**, InputArray **corners**, bool **patternWasFound**)

**Python:** cv2.**drawChessboardCorners**(image, patternSize, corners, patternWasFound) → None

**C:** void **cvDrawChessboardCorners**(CvArr* **image**, CvSize **pattern_size**, CvPoint2D32f* **corners**, int **count**, int **pattern_was_found**)

**Python:** cv.**DrawChessboardCorners**(image, patternSize, corners, patternWasFound) → None

> **Parameters**
>
>> **image** – Destination image. It must be an 8-bit color image.
>>
>> **patternSize** – Number of inner corners per a chessboard row and column (`patternSize = cv::Size(points_per_row,points_per_column)`).
>>
>> **corners** – Array of detected corners, the output of `findChessboardCorners`.
>>
>> **patternWasFound** – Parameter indicating whether the complete board was found or not. The return value of `findChessboardCorners()` should be passed here.

The function draws individual chessboard corners detected either as red circles if the board was not found, or as colored corners connected with lines if the board was found.

## findChessboardCorners

Finds the positions of internal corners of the chessboard.

**C++:** bool **findChessboardCorners**(InputArray **image**, Size **patternSize**, OutputArray **corners**, int **flags**=CALIB_CB_ADAPTIVE_THRESH+CALIB_CB_NORMALIZE_IMAGE )

**Python:** cv2.**findChessboardCorners**(image, patternSize[, corners[, flags ]]) → retval, corners

**C:** int **cvFindChessboardCorners**(const void* **image**, CvSize **pattern_size**, CvPoint2D32f* **corners**, int* **corner_count**=NULL, int **flags**=CV_CALIB_CB_ADAPTIVE_THRESH+CV_CALIB_CB_NORMALIZE_IMAGE )

**Python:** cv.**FindChessboardCorners**(image, patternSize, flags=CV_CALIB_CB_ADAPTIVE_THRESH) → corners

> **Parameters**
>
>> **image** – Source chessboard view. It must be an 8-bit grayscale or color image.
>>
>> **patternSize** – Number of inner corners per a chessboard row and column ( `patternSize = cvSize(points_per_row,points_per_colum) = cvSize(columns,rows)` ).
>>
>> **corners** – Output array of detected corners.
>>
>> **flags** – Various operation flags that can be zero or a combination of the following values:

- **CV_CALIB_CB_ADAPTIVE_THRESH** Use adaptive thresholding to convert the image to black and white, rather than a fixed threshold level (computed from the average image brightness).

- **CV_CALIB_CB_NORMALIZE_IMAGE** Normalize the image gamma with `equalizeHist()` before applying fixed or adaptive thresholding.

- **CV_CALIB_CB_FILTER_QUADS** Use additional criteria (like contour area, perimeter, square-like shape) to filter out false quads extracted at the contour retrieval stage.

- **CALIB_CB_FAST_CHECK** Run a fast check on the image that looks for chessboard corners, and shortcut the call if none is found. This can drastically speed up the call in the degenerate condition when no chessboard is observed.

The function attempts to determine whether the input image is a view of the chessboard pattern and locate the internal chessboard corners. The function returns a non-zero value if all of the corners are found and they are placed in a certain order (row by row, left to right in every row). Otherwise, if the function fails to find all the corners or reorder them, it returns 0. For example, a regular chessboard has 8 x 8 squares and 7 x 7 internal corners, that is, points where the black squares touch each other. The detected coordinates are approximate, and to determine their positions more accurately, the function calls `cornerSubPix()`. You also may use the function `cornerSubPix()` with different parameters if returned coordinates are not accurate enough.

Sample usage of detecting and drawing chessboard corners:

```
Size patternsize(8,6); //interior number of corners
Mat gray = ....; //source image
vector<Point2f> corners; //this will be filled by the detected corners

//CALIB_CB_FAST_CHECK saves a lot of time on images
//that do not contain any chessboard corners
bool patternfound = findChessboardCorners(gray, patternsize, corners,
        CALIB_CB_ADAPTIVE_THRESH + CALIB_CB_NORMALIZE_IMAGE
        + CALIB_CB_FAST_CHECK);

if(patternfound)
  cornerSubPix(gray, corners, Size(11, 11), Size(-1, -1),
    TermCriteria(CV_TERMCRIT_EPS + CV_TERMCRIT_ITER, 30, 0.1));

drawChessboardCorners(img, patternsize, Mat(corners), patternfound);
```

**Note:** The function requires white space (like a square-thick border, the wider the better) around the board to make the detection more robust in various environments. Otherwise, if there is no border and the background is dark, the outer black squares cannot be segmented properly and so the square grouping and ordering algorithm fails.

## findCirclesGrid

Finds centers in the grid of circles.

**C++:** bool **findCirclesGrid**(InputArray **image**, Size **patternSize**, OutputArray **centers**, int **flags**=CALIB_CB_SYMMETRIC_GRID, const Ptr<FeatureDetector>& **blobDetector**=new SimpleBlobDetector() )

**Python:** cv2.**findCirclesGridDefault**(image, patternSize[, centers[, flags]]) → retval, centers

        **Parameters**

                **image** – grid view of input circles; it must be an 8-bit grayscale or color image.

**patternSize** – number of circles per row and column ( patternSize = Size(points_per_row, points_per_colum) ).

**centers** – output array of detected centers.

**flags** – various operation flags that can be one of the following values:

  – **CALIB_CB_SYMMETRIC_GRID** uses symmetric pattern of circles.

  – **CALIB_CB_ASYMMETRIC_GRID** uses asymmetric pattern of circles.

  – **CALIB_CB_CLUSTERING** uses a special algorithm for grid detection. It is more robust to perspective distortions but much more sensitive to background clutter.

**blobDetector** – feature detector that finds blobs like dark circles on light background.

The function attempts to determine whether the input image contains a grid of circles. If it is, the function locates centers of the circles. The function returns a non-zero value if all of the centers have been found and they have been placed in a certain order (row by row, left to right in every row). Otherwise, if the function fails to find all the corners or reorder them, it returns 0.

Sample usage of detecting and drawing the centers of circles:

```
Size patternsize(7,7); //number of centers
Mat gray = ....; //source image
vector<Point2f> centers; //this will be filled by the detected centers

bool patternfound = findCirclesGrid(gray, patternsize, centers);

drawChessboardCorners(img, patternsize, Mat(centers), patternfound);
```

**Note:** The function requires white space (like a square-thick border, the wider the better) around the board to make the detection more robust in various environments.

## solvePnP

Finds an object pose from 3D-2D point correspondences.

**C++:** bool **solvePnP**(InputArray **objectPoints**, InputArray **imagePoints**, InputArray **cameraMatrix**, InputArray **distCoeffs**, OutputArray **rvec**, OutputArray **tvec**, bool **useExtrinsicGuess**=false, int **flags**=ITERATIVE )

**Python:** cv2.**solvePnP**(objectPoints, imagePoints, cameraMatrix, distCoeffs[, rvec[, tvec[, useExtrinsicGuess[, flags ] ] ] ]) → retval, rvec, tvec

**C:** void **cvFindExtrinsicCameraParams2**(const CvMat* **object_points**, const CvMat* **image_points**, const CvMat* **camera_matrix**, const CvMat* **distortion_coeffs**, CvMat* **rotation_vector**, CvMat* **translation_vector**, int **use_extrinsic_guess**=0 )

**Python:** cv.**FindExtrinsicCameraParams2**(objectPoints, imagePoints, cameraMatrix, distCoeffs, rvec, tvec, useExtrinsicGuess=0) → None

   Parameters

      **objectPoints** – Array of object points in the object coordinate space, 3xN/Nx3 1-channel or 1xN/Nx1 3-channel, where N is the number of points. vector<Point3f> can be also passed here.

      **imagePoints** – Array of corresponding image points, 2xN/Nx2 1-channel or 1xN/Nx1 2-channel, where N is the number of points. vector<Point2f> can be also passed here.

$$\textbf{cameraMatrix} - \text{Input camera matrix } A = \begin{bmatrix} fx & 0 & cx \\ 0 & fy & cy \\ 0 & 0 & 1 \end{bmatrix}.$$

**distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.

**rvec** – Output rotation vector (see `Rodrigues()` ) that, together with `tvec` , brings points from the model coordinate system to the camera coordinate system.

**tvec** – Output translation vector.

**useExtrinsicGuess** – If true (1), the function uses the provided `rvec` and `tvec` values as initial approximations of the rotation and translation vectors, respectively, and further optimizes them.

**flags** – Method for solving a PnP problem:

– **CV_ITERATIVE** Iterative method is based on Levenberg-Marquardt optimization. In this case the function finds such a pose that minimizes reprojection error, that is the sum of squared distances between the observed projections `imagePoints` and the projected (using `projectPoints()` ) `objectPoints` .

– **CV_P3P** Method is based on the paper of X.S. Gao, X.-R. Hou, J. Tang, H.-F. Chang "Complete Solution Classification for the Perspective-Three-Point Problem". In this case the function requires exactly four object and image points.

– **CV_EPNP** Method has been introduced by F.Moreno-Noguer, V.Lepetit and P.Fua in the paper "EPnP: Efficient Perspective-n-Point Camera Pose Estimation".

The function estimates the object pose given a set of object points, their corresponding image projections, as well as the camera matrix and the distortion coefficients.

## solvePnPRansac

Finds an object pose from 3D-2D point correspondences using the RANSAC scheme.

**C++:** void **solvePnPRansac**(InputArray **objectPoints**, InputArray **imagePoints**, InputArray **cameraMatrix**, InputArray **distCoeffs**, OutputArray **rvec**, OutputArray **tvec**, bool **useExtrinsicGuess**=false, int **iterationsCount**=100, float **reprojectionError**=8.0, int **minInliersCount**=100, OutputArray **inliers**=noArray(), int **flags**=ITERATIVE )

**Python:** cv2.**solvePnPRansac**(objectPoints, imagePoints, cameraMatrix, distCoeffs[, rvec[, tvec[, useExtrinsicGuess[, iterationsCount[, reprojectionError[, minInliersCount[, inliers[, flags ]]]]]]]]) → rvec, tvec, inliers

**Parameters**

**objectPoints** – Array of object points in the object coordinate space, 3xN/Nx3 1-channel or 1xN/Nx1 3-channel, where N is the number of points. `vector<Point3f>` can be also passed here.

**imagePoints** – Array of corresponding image points, 2xN/Nx2 1-channel or 1xN/Nx1 2-channel, where N is the number of points. `vector<Point2f>` can be also passed here.

$$\textbf{cameraMatrix} - \text{Input camera matrix } A = \begin{bmatrix} fx & 0 & cx \\ 0 & fy & cy \\ 0 & 0 & 1 \end{bmatrix}.$$

> **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
>
> **rvec** – Output rotation vector (see `Rodrigues()` ) that, together with `tvec` , brings points from the model coordinate system to the camera coordinate system.
>
> **tvec** – Output translation vector.
>
> **useExtrinsicGuess** – If true (1), the function uses the provided `rvec` and `tvec` values as initial approximations of the rotation and translation vectors, respectively, and further optimizes them.
>
> **iterationsCount** – Number of iterations.
>
> **reprojectionError** – Inlier threshold value used by the RANSAC procedure. The parameter value is the maximum allowed distance between the observed and computed point projections to consider it an inlier.
>
> **minInliersCount** – Number of inliers. If the algorithm at some stage finds more inliers than `minInliersCount` , it finishes.
>
> **inliers** – Output vector that contains indices of inliers in `objectPoints` and `imagePoints` .
>
> **flags** – Method for solving a PnP problem (see `solvePnP()` ).

The function estimates an object pose given a set of object points, their corresponding image projections, as well as the camera matrix and the distortion coefficients. This function finds such a pose that minimizes reprojection error, that is, the sum of squared distances between the observed projections `imagePoints` and the projected (using `projectPoints()` ) `objectPoints`. The use of RANSAC makes the function resistant to outliers. The function is parallelized with the TBB library.

## findFundamentalMat

Calculates a fundamental matrix from the corresponding points in two images.

**C++:** Mat **findFundamentalMat**(InputArray **points1**, InputArray **points2**, int **method**=FM_RANSAC, double **param1**=3., double **param2**=0.99, OutputArray **mask**=noArray() )

**Python:** cv2.**findFundamentalMat**(points1, points2[, method[, param1[, param2[, mask ]]]]) → retval, mask

**C:** int **cvFindFundamentalMat**(const CvMat* **points1**, const CvMat* **points2**, CvMat* **fundamental_matrix**, int **method**=CV_FM_RANSAC, double **param1**=3., double **param2**=0.99, CvMat* **status**=NULL )

**Python:** cv.**FindFundamentalMat**(points1, points2, fundamentalMatrix, method=CV_FM_RANSAC, param1=1., param2=0.99, status=None) → retval

> **Parameters**
>
> > **points1** – Array of `N` points from the first image. The point coordinates should be floating-point (single or double precision).
> >
> > **points2** – Array of the second image points of the same size and format as `points1` .
> >
> > **method** – Method for computing a fundamental matrix.
> >
> > – **CV_FM_7POINT** for a 7-point algorithm. $N = 7$
> >
> > – **CV_FM_8POINT** for an 8-point algorithm. $N \geq 8$
> >
> > – **CV_FM_RANSAC** for the RANSAC algorithm. $N \geq 8$

---

- **CV_FM_LMEDS** for the LMedS algorithm. $N \geq 8$

**param1** – Parameter used for RANSAC. It is the maximum distance from a point to an epipolar line in pixels, beyond which the point is considered an outlier and is not used for computing the final fundamental matrix. It can be set to something like 1-3, depending on the accuracy of the point localization, image resolution, and the image noise.

**param2** – Parameter used for the RANSAC or LMedS methods only. It specifies a desirable level of confidence (probability) that the estimated matrix is correct.

**status** – Output array of N elements, every element of which is set to 0 for outliers and to 1 for the other points. The array is computed only in the RANSAC and LMedS methods. For other methods, it is set to all 1's.

The epipolar geometry is described by the following equation:

$$[p_2; 1]^T F [p_1; 1] = 0$$

where $F$ is a fundamental matrix, $p_1$ and $p_2$ are corresponding points in the first and the second images, respectively.

The function calculates the fundamental matrix using one of four methods listed above and returns the found fundamental matrix. Normally just one matrix is found. But in case of the 7-point algorithm, the function may return up to 3 solutions ( $9 \times 3$ matrix that stores all 3 matrices sequentially).

The calculated fundamental matrix may be passed further to `computeCorrespondEpilines()` that finds the epipolar lines corresponding to the specified points. It can also be passed to `stereoRectifyUncalibrated()` to compute the rectification transformation.

```
// Example. Estimation of fundamental matrix using the RANSAC algorithm
int point_count = 100;
vector<Point2f> points1(point_count);
vector<Point2f> points2(point_count);

// initialize the points here ... */
for( int i = 0; i < point_count; i++ )
{
    points1[i] = ...;
    points2[i] = ...;
}

Mat fundamental_matrix =
 findFundamentalMat(points1, points2, FM_RANSAC, 3, 0.99);
```

## findHomography

Finds a perspective transformation between two planes.

**C++:** Mat **findHomography**(InputArray **srcPoints**, InputArray **dstPoints**, int **method**=0, double **ransacReprojThreshold**=3, OutputArray **mask**=noArray() )

**Python:** cv2.**findHomography**(srcPoints, dstPoints[, method[, ransacReprojThreshold[, mask]]]) → retval, mask

**C:** int **cvFindHomography**(const CvMat* **src_points**, const CvMat* **dst_points**, CvMat* **homography**, int **method**=0, double **ransacReprojThreshold**=3, CvMat* **mask**=0 )

**Python:** cv.**FindHomography**(srcPoints, dstPoints, H, method=0, ransacReprojThreshold=3.0, status=None) → None

Parameters

**srcPoints** – Coordinates of the points in the original plane, a matrix of the type `CV_32FC2` or `vector<Point2f>` .

**dstPoints** – Coordinates of the points in the target plane, a matrix of the type `CV_32FC2` or a `vector<Point2f>` .

**method** – Method used to computed a homography matrix. The following methods are possible:

- **0** - a regular method using all the points

- **CV_RANSAC** - RANSAC-based robust method

- **CV_LMEDS** - Least-Median robust method

**ransacReprojThreshold** – Maximum allowed reprojection error to treat a point pair as an inlier (used in the RANSAC method only). That is, if

$$\|\text{dstPoints}_i - \text{convertPointsHomogeneous}(H * \text{srcPoints}_i)\| > \text{ransacReprojThreshold}$$

then the point i is considered an outlier. If `srcPoints` and `dstPoints` are measured in pixels, it usually makes sense to set this parameter somewhere in the range of 1 to 10.

**mask** – Optional output mask set by a robust method ( `CV_RANSAC` or `CV_LMEDS` ). Note that the input mask values are ignored.

The functions find and return the perspective transformation H between the source and the destination planes:

$$s_i \begin{bmatrix} x_i' \\ y_i' \\ 1 \end{bmatrix} \sim H \begin{bmatrix} x_i \\ y_i \\ 1 \end{bmatrix}$$

so that the back-projection error

$$\sum_i \left( x_i' - \frac{h_{11}x_i + h_{12}y_i + h_{13}}{h_{31}x_i + h_{32}y_i + h_{33}} \right)^2 + \left( y_i' - \frac{h_{21}x_i + h_{22}y_i + h_{23}}{h_{31}x_i + h_{32}y_i + h_{33}} \right)^2$$

is minimized. If the parameter `method` is set to the default value 0, the function uses all the point pairs to compute an initial homography estimate with a simple least-squares scheme.

However, if not all of the point pairs ( $\text{srcPoints}_i$,:math:*dstPoints_i* ) fit the rigid perspective transformation (that is, there are some outliers), this initial estimate will be poor. In this case, you can use one of the two robust methods. Both methods, `RANSAC` and `LMeDS` , try many different random subsets of the corresponding point pairs (of four pairs each), estimate the homography matrix using this subset and a simple least-square algorithm, and then compute the quality/goodness of the computed homography (which is the number of inliers for RANSAC or the median reprojection error for LMeDs). The best subset is then used to produce the initial estimate of the homography matrix and the mask of inliers/outliers.

Regardless of the method, robust or not, the computed homography matrix is refined further (using inliers only in case of a robust method) with the Levenberg-Marquardt method to reduce the re-projection error even more.

The method `RANSAC` can handle practically any ratio of outliers but it needs a threshold to distinguish inliers from outliers. The method `LMeDS` does not need any threshold but it works correctly only when there are more than 50% of inliers. Finally, if there are no outliers and the noise is rather small, use the default method (`method=0`).

The function is used to find initial intrinsic and extrinsic matrices. Homography matrix is determined up to a scale. Thus, it is normalized so that $h_{33} = 1$ .

**See Also:**

`getAffineTransform()`, `getPerspectiveTransform()`, `estimateRigidTransform()`, `warpPerspective()`, `perspectiveTransform()`

## estimateAffine3D

Computes an optimal affine transformation between two 3D point sets.

**C++:** int **estimateAffine3D**(InputArray **src**, InputArray **dst**, OutputArray **out**, OutputArray **inliers**, double **ransacThreshold**=3, double **confidence**=0.99)

**Python:** cv2.**estimateAffine3D**(src, dst[, out[, inliers[, ransacThreshold[, confidence ]]]]) → retval, out, inliers

> **Parameters**

>> **src** – First input 3D point set.

>> **dst** – Second input 3D point set.

>> **out** – Output 3D affine transformation matrix $3 \times 4$ .

>> **inliers** – Output vector indicating which points are inliers.

>> **ransacThreshold** – Maximum reprojection error in the RANSAC algorithm to consider a point as an inlier.

>> **confidence** – Confidence level, between 0 and 1, for the estimated transformation. Anything between 0.95 and 0.99 is usually good enough. Values too close to 1 can slow down the estimation significantly. Values lower than 0.8-0.9 can result in an incorrectly estimated transformation.

The function estimates an optimal 3D affine transformation between two 3D point sets using the RANSAC algorithm.

## filterSpeckles

Filters off small noise blobs (speckles) in the disparity map

**C++:** void **filterSpeckles**(InputOutputArray **img**, double **newVal**, int **maxSpeckleSize**, double **maxDiff**, InputOutputArray **buf**=noArray() )

**Python:** cv2.**filterSpeckles**(img, newVal, maxSpeckleSize, maxDiff[, buf ]) → None

> **Parameters**

>> **img** – The input 16-bit signed disparity image

>> **newVal** – The disparity value used to paint-off the speckles

>> **maxSpeckleSize** – The maximum speckle size to consider it a speckle. Larger blobs are not affected by the algorithm

>> **maxDiff** – Maximum difference between neighbor disparity pixels to put them into the same blob. Note that since StereoBM, StereoSGBM and may be other algorithms return a fixed-point disparity map, where disparity values are multiplied by 16, this scale factor should be taken into account when specifying this parameter value.

>> **buf** – The optional temporary buffer to avoid memory allocation within the function.

## getOptimalNewCameraMatrix

Returns the new camera matrix based on the free scaling parameter.

**C++:** Mat **getOptimalNewCameraMatrix**(InputArray **cameraMatrix**, InputArray **distCoeffs**, Size **imageSize**, double **alpha**, Size **newImgSize**=Size(), Rect* **validPixROI**=0, bool **centerPrincipalPoint**=false )

**Python:** cv2.**getOptimalNewCameraMatrix**(cameraMatrix, distCoeffs, imageSize, alpha$[$, newImgSize$[$, centerPrincipalPoint $]]$) → retval, validPixROI

**C:** void **cvGetOptimalNewCameraMatrix**(const CvMat* **camera_matrix**, const CvMat* **dist_coeffs**, CvSize **image_size**, double **alpha**, CvMat* **new_camera_matrix**, CvSize **new_imag_size**=cvSize(0,0), CvRect* **valid_pixel_ROI**=0, int **center_principal_point**=0 )

**Python:** cv.**GetOptimalNewCameraMatrix**(cameraMatrix, distCoeffs, imageSize, alpha, newCameraMatrix, newImageSize=(0, 0), validPixROI=0, centerPrincipalPoint=0) → None

> **Parameters**
>
> > **cameraMatrix** – Input camera matrix.
> >
> > **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
> >
> > **imageSize** – Original image size.
> >
> > **alpha** – Free scaling parameter between 0 (when all the pixels in the undistorted image are valid) and 1 (when all the source image pixels are retained in the undistorted image). See stereoRectify() for details.
> >
> > **new_camera_matrix** – Output new camera matrix.
> >
> > **new_imag_size** – Image size after rectification. By default,it is set to imageSize .
> >
> > **validPixROI** – Optional output rectangle that outlines all-good-pixels region in the undistorted image. See roi1, roi2 description in stereoRectify() .
> >
> > **centerPrincipalPoint** – Optional flag that indicates whether in the new camera matrix the principal point should be at the image center or not. By default, the principal point is chosen to best fit a subset of the source image (determined by alpha) to the corrected image.

The function computes and returns the optimal new camera matrix based on the free scaling parameter. By varying this parameter, you may retrieve only sensible pixels alpha=0 , keep all the original image pixels if there is valuable information in the corners alpha=1 , or get something in between. When alpha>0 , the undistortion result is likely to have some black pixels corresponding to "virtual" pixels outside of the captured distorted image. The original camera matrix, distortion coefficients, the computed new camera matrix, and newImageSize should be passed to initUndistortRectifyMap() to produce the maps for remap() .

## initCameraMatrix2D

Finds an initial camera matrix from 3D-2D point correspondences.

**C++:** Mat **initCameraMatrix2D**(InputArrayOfArrays **objectPoints**, InputArrayOfArrays **imagePoints**, Size **imageSize**, double **aspectRatio**=1.)

**Python:** cv2.**initCameraMatrix2D**(objectPoints, imagePoints, imageSize$[$, aspectRatio $]$) → retval

**C:** void **cvInitIntrinsicParams2D**(const CvMat* **object_points**, const CvMat* **image_points**, const CvMat* **npoints**, CvSize **image_size**, CvMat* **camera_matrix**, double **aspect_ratio**=1. )

**Python:** cv.**InitIntrinsicParams2D**(objectPoints, imagePoints, npoints, imageSize, cameraMatrix, aspectRatio=1.) → None

> **Parameters**

**objectPoints** – Vector of vectors of the calibration pattern points in the calibration pattern coordinate space. In the old interface all the per-view vectors are concatenated. See `calibrateCamera()` for details.

**imagePoints** – Vector of vectors of the projections of the calibration pattern points. In the old interface all the per-view vectors are concatenated.

**npoints** – The integer vector of point counters for each view.

**imageSize** – Image size in pixels used to initialize the principal point.

**aspectRatio** – If it is zero or negative, both $f_x$ and $f_y$ are estimated independently. Otherwise, $f_x = f_y *$ `aspectRatio` .

The function estimates and returns an initial camera matrix for the camera calibration process. Currently, the function only supports planar calibration patterns, which are patterns where each object point has z-coordinate =0.

## matMulDeriv

Computes partial derivatives of the matrix product for each multiplied matrix.

**C++:** void **matMulDeriv**(InputArray **A**, InputArray **B**, OutputArray **dABdA**, OutputArray **dABdB**)

**Python:** cv2.**matMulDeriv**(A, B[, dABdA[, dABdB]]) → dABdA, dABdB

> **Parameters**
>
>> **A** – First multiplied matrix.
>>
>> **B** – Second multiplied matrix.
>>
>> **dABdA** – First output derivative matrix d(A*B)/dA of size `A.rows*B.cols` $\times$ $A.rows * A.cols$ .
>>
>> **dABdB** – Second output derivative matrix d(A*B)/dB of size `A.rows*B.cols` $\times$ $B.rows * B.cols$ .

The function computes partial derivatives of the elements of the matrix product $A * B$ with regard to the elements of each of the two input matrices. The function is used to compute the Jacobian matrices in `stereoCalibrate()` but can also be used in any other similar optimization function.

## projectPoints

Projects 3D points to an image plane.

**C++:** void **projectPoints**(InputArray **objectPoints**, InputArray **rvec**, InputArray **tvec**, InputArray **cameraMatrix**, InputArray **distCoeffs**, OutputArray **imagePoints**, OutputArray **jacobian**=noArray(), double **aspectRatio**=0 )

**Python:** cv2.**projectPoints**(objectPoints, rvec, tvec, cameraMatrix, distCoeffs[, imagePoints[, jacobian[, aspectRatio]]]) → imagePoints, jacobian

**C:** void **cvProjectPoints2**(const CvMat* **object_points**, const CvMat* **rotation_vector**, const CvMat* **translation_vector**, const CvMat* **camera_matrix**, const CvMat* **distortion_coeffs**, CvMat* **image_points**, CvMat* **dpdrot**=NULL, CvMat* **dpdt**=NULL, CvMat* **dpdf**=NULL, CvMat* **dpdc**=NULL, CvMat* **dpddist**=NULL, double **aspect_ratio**=0 )

**Python:** cv.**ProjectPoints2**(objectPoints, rvec, tvec, cameraMatrix, distCoeffs, imagePoints, dpdrot=None, dpdt=None, dpdf=None, dpdc=None, dpddist=None) → None

> **Parameters**

> **objectPoints** – Array of object points, 3xN/Nx3 1-channel or 1xN/Nx1 3-channel (or `vector<Point3f>`), where N is the number of points in the view.
>
> **rvec** – Rotation vector. See `Rodrigues()` for details.
>
> **tvec** – Translation vector.
>
> **cameraMatrix** – Camera matrix $A = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$.
>
> **distCoeffs** – Input vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
>
> **imagePoints** – Output array of image points, 2xN/Nx2 1-channel or 1xN/Nx1 2-channel, or `vector<Point2f>`.
>
> **jacobian** – Optional output 2Nx(10+<numDistCoeffs>) jacobian matrix of derivatives of image points with respect to components of the rotation vector, translation vector, focal lengths, coordinates of the principal point and the distortion coefficients. In the old interface different components of the jacobian are returned via different output parameters.
>
> **aspectRatio** – Optional "fixed aspect ratio" parameter. If the parameter is not 0, the function assumes that the aspect ratio (*fx/fy*) is fixed and correspondingly adjusts the jacobian matrix.

The function computes projections of 3D points to the image plane given intrinsic and extrinsic camera parameters. Optionally, the function computes Jacobians - matrices of partial derivatives of image points coordinates (as functions of all the input parameters) with respect to the particular parameters, intrinsic and/or extrinsic. The Jacobians are used during the global optimization in `calibrateCamera()`, `solvePnP()`, and `stereoCalibrate()` . The function itself can also be used to compute a re-projection error given the current intrinsic and extrinsic parameters.

---

**Note:** By setting rvec=tvec=(0,0,0) or by setting `cameraMatrix` to a 3x3 identity matrix, or by passing zero distortion coefficients, you can get various useful partial cases of the function. This means that you can compute the distorted coordinates for a sparse set of points or apply a perspective transformation (and also compute the derivatives) in the ideal zero-distortion setup.

---

## reprojectImageTo3D

Reprojects a disparity image to 3D space.

**C++:** void **reprojectImageTo3D**(InputArray **disparity**, OutputArray **_3dImage**, InputArray **Q**, bool **handleMissingValues**=false, int **ddepth**=-1 )

**Python:** cv2.**reprojectImageTo3D**(disparity, Q[, _3dImage[, handleMissingValues[, ddepth ]]]) → _3dImage

**C:** void **cvReprojectImageTo3D**(const CvArr* **disparityImage**, CvArr* **_3dImage**, const CvMat* **Q**, int **handleMissingValues**=0 )

**Python:** cv.**ReprojectImageTo3D**(disparity, _3dImage, Q, handleMissingValues=0) → None

> **Parameters**
>
> > **disparity** – Input single-channel 8-bit unsigned, 16-bit signed, 32-bit signed or 32-bit floating-point disparity image.
> >
> > **_3dImage** – Output 3-channel floating-point image of the same size as `disparity` . Each element of `_3dImage(x,y)` contains 3D coordinates of the point (x,y) computed from the disparity map.

---

**Q** – $4 \times 4$ perspective transformation matrix that can be obtained with `stereoRectify()`.

**handleMissingValues** – Indicates, whether the function should handle missing values (i.e. points where the disparity was not computed). If `handleMissingValues=true`, then pixels with the minimal disparity that corresponds to the outliers (see `StereoBM::operator()` ) are transformed to 3D points with a very large Z value (currently set to 10000).

**ddepth** – The optional output array depth. If it is `-1`, the output image will have `CV_32F` depth. `ddepth` can also be set to `CV_16S`, `CV_32S` or `CV_32F`.

The function transforms a single-channel disparity map to a 3-channel image representing a 3D surface. That is, for each pixel `(x,y)` andthe corresponding disparity `d=disparity(x,y)` , it computes:

$$[X\ Y\ Z\ W]^\mathsf{T} = \mathtt{Q} * [x\ y\ \mathtt{disparity}(x,y)\ 1]^\mathsf{T}$$
$$\_3d\mathtt{Image}(x,y) = (X/W,\ Y/W,\ Z/W)$$

The matrix `Q` can be an arbitrary $4 \times 4$ matrix (for example, the one computed by `stereoRectify()`). To reproject a sparse set of points $\{(x,y,d),...\}$ to 3D space, use `perspectiveTransform()` .

## RQDecomp3x3

Computes an RQ decomposition of 3x3 matrices.

**C++:** Vec3d **RQDecomp3x3**(InputArray **src**, OutputArray **mtxR**, OutputArray **mtxQ**, OutputArray **Qx**=noArray(), OutputArray **Qy**=noArray(), OutputArray **Qz**=noArray() )

**Python:** cv2.**RQDecomp3x3**(src[, mtxR[, mtxQ[, Qx[, Qy[, Qz ] ] ] ] ]) → retval, mtxR, mtxQ, Qx, Qy, Qz

**C:** void **cvRQDecomp3x3**(const CvMat* **matrixM**, CvMat* **matrixR**, CvMat* **matrixQ**, CvMat* **matrixQx**=NULL, CvMat* **matrixQy**=NULL, CvMat* **matrixQz**=NULL, CvPoint3D64f* **eulerAngles**=NULL )

**Python:** cv.**RQDecomp3x3**(M, R, Q, Qx=None, Qy=None, Qz=None) → eulerAngles

> **Parameters**
>
> > **src** – 3x3 input matrix.
> >
> > **mtxR** – Output 3x3 upper-triangular matrix.
> >
> > **mtxQ** – Output 3x3 orthogonal matrix.
> >
> > **Qx** – Optional output 3x3 rotation matrix around x-axis.
> >
> > **Qy** – Optional output 3x3 rotation matrix around y-axis.
> >
> > **Qz** – Optional output 3x3 rotation matrix around z-axis.

The function computes a RQ decomposition using the given rotations. This function is used in `decomposeProjectionMatrix()` to decompose the left 3x3 submatrix of a projection matrix into a camera and a rotation matrix.

It optionally returns three rotation matrices, one for each axis, and the three Euler angles in degrees (as the return value) that could be used in OpenGL. Note, there is always more than one sequence of rotations about the three principle axes that results in the same orientation of an object, eg. see [Slabaugh]. Returned tree rotation matrices and corresponding three Euler angules are only one of the possible solutions.

## Rodrigues

Converts a rotation matrix to a rotation vector or vice versa.

**C++:** void **Rodrigues**(InputArray **src**, OutputArray **dst**, OutputArray **jacobian**=noArray())

**Python:** cv2.**Rodrigues**(src[, dst[, jacobian ]]) → dst, jacobian

**C:** int **cvRodrigues2**(const CvMat* **src**, CvMat* **dst**, CvMat* **jacobian**=0 )

**Python:** cv.**Rodrigues2**(src, dst, jacobian=0) → None

> Parameters
>> **src** – Input rotation vector (3x1 or 1x3) or rotation matrix (3x3).
>>
>> **dst** – Output rotation matrix (3x3) or rotation vector (3x1 or 1x3), respectively.
>>
>> **jacobian** – Optional output Jacobian matrix, 3x9 or 9x3, which is a matrix of partial derivatives of the output array components with respect to the input array components.

$$\theta \leftarrow norm(r)$$
$$r \leftarrow r/\theta$$
$$R = \cos\theta I + (1 - \cos\theta) r r^T + \sin\theta \begin{bmatrix} 0 & -r_z & r_y \\ r_z & 0 & -r_x \\ -r_y & r_x & 0 \end{bmatrix}$$

Inverse transformation can be also done easily, since

$$\sin(\theta) \begin{bmatrix} 0 & -r_z & r_y \\ r_z & 0 & -r_x \\ -r_y & r_x & 0 \end{bmatrix} = \frac{R - R^T}{2}$$

A rotation vector is a convenient and most compact representation of a rotation matrix (since any rotation matrix has just 3 degrees of freedom). The representation is used in the global 3D geometry optimization procedures like `calibrateCamera()`, `stereoCalibrate()`, or `solvePnP()` .

## StereoBM

class **StereoBM**

Class for computing stereo correspondence using the block matching algorithm.

```
// Block matching stereo correspondence algorithm class StereoBM
{
    enum { NORMALIZED_RESPONSE = CV_STEREO_BM_NORMALIZED_RESPONSE,
        BASIC_PRESET=CV_STEREO_BM_BASIC,
        FISH_EYE_PRESET=CV_STEREO_BM_FISH_EYE,
        NARROW_PRESET=CV_STEREO_BM_NARROW };

    StereoBM();
    // the preset is one of ..._PRESET above.
    // ndisparities is the size of disparity range,
    // in which the optimal disparity at each pixel is searched for.
    // SADWindowSize is the size of averaging window used to match pixel blocks
    //    (larger values mean better robustness to noise, but yield blurry disparity maps)
    StereoBM(int preset, int ndisparities=0, int SADWindowSize=21);
    // separate initialization function
    void init(int preset, int ndisparities=0, int SADWindowSize=21);
    // computes the disparity for the two rectified 8-bit single-channel images.
    // the disparity will be 16-bit signed (fixed-point) or 32-bit floating-point image of the same size as left.
    void operator()( InputArray left, InputArray right, OutputArray disparity, int disptype=CV_16S );
```

```
    Ptr<CvStereoBMState> state;
};
```

The class is a C++ wrapper for the associated functions. In particular, `StereoBM::operator()` is the wrapper for `cvFindStereoCorrespondenceBM()`.

## StereoBM::StereoBM

The constructors.

**C++:** `StereoBM::`**`StereoBM`**`()`

**C++:** `StereoBM::`**`StereoBM`**`(int **preset**, int **ndisparities**=0, int **SADWindowSize**=21)`

**Python:** `cv2.`**`StereoBM`**`([preset[, ndisparities[, SADWindowSize]]]) → <StereoBM object>`

**C:** CvStereoBMState* **`cvCreateStereoBMState`**`(int **preset**=CV_STEREO_BM_BASIC, int **numberOfDisparities**=0 )`

**Python:** `cv.`**`CreateStereoBMState`**`(preset=CV_STEREO_BM_BASIC, numberOfDisparities=0) → CvStereoBMState`

> **Parameters**

>> **preset** – specifies the whole set of algorithm parameters, one of:

>> – BASIC_PRESET - parameters suitable for general cameras

>> – FISH_EYE_PRESET - parameters suitable for wide-angle cameras

>> – NARROW_PRESET - parameters suitable for narrow-angle cameras

>> After constructing the class, you can override any parameters set by the preset.

>> **ndisparities** – the disparity search range. For each pixel algorithm will find the best disparity from 0 (default minimum disparity) to `ndisparities`. The search range can then be shifted by changing the minimum disparity.

>> **SADWindowSize** – the linear size of the blocks compared by the algorithm. The size should be odd (as the block is centered at the current pixel). Larger block size implies smoother, though less accurate disparity map. Smaller block size gives more detailed disparity map, but there is higher chance for algorithm to find a wrong correspondence.

The constructors initialize `StereoBM` state. You can then call `StereoBM::operator()` to compute disparity for a specific stereo pair.

---

**Note:** In the C API you need to deallocate `CvStereoBM` state when it is not needed anymore using `cvReleaseStereoBMState(&stereobm)`.

---

## StereoBM::operator()

Computes disparity using the BM algorithm for a rectified stereo pair.

**C++:** void `StereoBM::`**`operator()`**`(InputArray **left**, InputArray **right**, OutputArray **disparity**, int **disptype**=CV_16S )`

**Python:** `cv2.StereoBM.`**`compute`**`(left, right[, disparity[, disptype]]) → disparity`

**C:** void **`cvFindStereoCorrespondenceBM`**`(const CvArr* **left**, const CvArr* **right**, CvArr* **disparity**, CvStereoBMState* **state**)`

---

**Python:** cv.**FindStereoCorrespondenceBM**(left, right, disparity, state) → None

> **Parameters**
>
>> **left** – Left 8-bit single-channel image.
>>
>> **right** – Right image of the same size and the same type as the left one.
>>
>> **disparity** – Output disparity map. It has the same size as the input images. When `disptype==CV_16S`, the map is a 16-bit signed single-channel image, containing disparity values scaled by 16. To get the true disparity values from such fixed-point representation, you will need to divide each `disp` element by 16. If `disptype==CV_32F`, the disparity map will already contain the real disparity values on output.
>>
>> **disptype** – Type of the output disparity map, `CV_16S` (default) or `CV_32F`.
>>
>> **state** – The pre-initialized `CvStereoBMState` structure in the case of the old API.

The method executes the BM algorithm on a rectified stereo pair. See the `stereo_match.cpp` OpenCV sample on how to prepare images and call the method. Note that the method is not constant, thus you should not use the same `StereoBM` instance from within different threads simultaneously. The function is parallelized with the TBB library.

## StereoSGBM

**class StereoSGBM**

Class for computing stereo correspondence using the semi-global block matching algorithm.

```cpp
class StereoSGBM
{
    StereoSGBM();
    StereoSGBM(int minDisparity, int numDisparities, int SADWindowSize,
               int P1=0, int P2=0, int disp12MaxDiff=0,
               int preFilterCap=0, int uniquenessRatio=0,
               int speckleWindowSize=0, int speckleRange=0,
               bool fullDP=false);
    virtual ~StereoSGBM();

    virtual void operator()(InputArray left, InputArray right, OutputArray disp);

    int minDisparity;
    int numberOfDisparities;
    int SADWindowSize;
    int preFilterCap;
    int uniquenessRatio;
    int P1, P2;
    int speckleWindowSize;
    int speckleRange;
    int disp12MaxDiff;
    bool fullDP;

    ...
};
```

The class implements the modified H. Hirschmuller algorithm [HH08] that differs from the original one as follows:

- By default, the algorithm is single-pass, which means that you consider only 5 directions instead of 8. Set `fullDP=true` to run the full variant of the algorithm but beware that it may consume a lot of memory.

- The algorithm matches blocks, not individual pixels. Though, setting `SADWindowSize=1` reduces the blocks to single pixels.

- Mutual information cost function is not implemented. Instead, a simpler Birchfield-Tomasi sub-pixel metric from [BT98] is used. Though, the color images are supported as well.

- Some pre- and post- processing steps from K. Konolige algorithm `StereoBM::operator()` are included, for example: pre-filtering (`CV_STEREO_BM_XSOBEL` type) and post-filtering (uniqueness check, quadratic interpolation and speckle filtering).

## StereoSGBM::StereoSGBM

**C++:** `StereoSGBM::`**`StereoSGBM`**`()`

**C++:** `StereoSGBM::`**`StereoSGBM`**(int **minDisparity**, int **numDisparities**, int **SADWindowSize**, int **P1**=0, int **P2**=0, int **disp12MaxDiff**=0, int **preFilterCap**=0, int **uniquenessRatio**=0, int **speckleWindowSize**=0, int **speckleRange**=0, bool **fullDP**=false)

**Python:** `cv2.`**`StereoSGBM`**([minDisparity, numDisparities, SADWindowSize[, P1[, P2[, disp12MaxDiff[, preFilterCap[, uniquenessRatio[, speckleWindowSize[, speckleRange[, fullDP]]]]]]]]]) → <StereoSGBM object>
Initializes `StereoSGBM` and sets parameters to custom values.??

   **Parameters**

   **minDisparity** – Minimum possible disparity value. Normally, it is zero but sometimes rectification algorithms can shift images, so this parameter needs to be adjusted accordingly.

   **numDisparities** – Maximum disparity minus minimum disparity. The value is always greater than zero. In the current implementation, this parameter must be divisible by 16.

   **SADWindowSize** – Matched block size. It must be an odd number `>=1`. Normally, it should be somewhere in the `3..11` range.

   **P1** – The first parameter controlling the disparity smoothness. See below.

   **P2** – The second parameter controlling the disparity smoothness. The larger the values are, the smoother the disparity is. `P1` is the penalty on the disparity change by plus or minus 1 between neighbor pixels. `P2` is the penalty on the disparity change by more than 1 between neighbor pixels. The algorithm requires `P2 > P1`. See `stereo_match.cpp` sample where some reasonably good `P1` and `P2` values are shown (like `8*number_of_image_channels*SADWindowSize*SADWindowSize` and `32*number_of_image_channels*SADWindowSize*SADWindowSize`, respectively).

   **disp12MaxDiff** – Maximum allowed difference (in integer pixel units) in the left-right disparity check. Set it to a non-positive value to disable the check.

   **preFilterCap** – Truncation value for the prefiltered image pixels. The algorithm first computes x-derivative at each pixel and clips its value by `[-preFilterCap, preFilterCap]` interval. The result values are passed to the Birchfield-Tomasi pixel cost function.

   **uniquenessRatio** – Margin in percentage by which the best (minimum) computed cost function value should "win" the second best value to consider the found match correct. Normally, a value within the 5-15 range is good enough.

   **speckleWindowSize** – Maximum size of smooth disparity regions to consider their noise speckles and invalidate. Set it to 0 to disable speckle filtering. Otherwise, set it somewhere in the 50-200 range.

   **speckleRange** – Maximum disparity variation within each connected component. If you do speckle filtering, set the parameter to a positive value, multiple of 16. Normally, 16 or 32 is good enough.

> **fullDP** – Set it to `true` to run the full-scale two-pass dynamic programming algorithm. It will consume O(W*H*numDisparities) bytes, which is large for 640x480 stereo and huge for HD-size pictures. By default, it is set to `false` .

The first constructor initializes `StereoSGBM` with all the default parameters. So, you only have to set `StereoSGBM::numberOfDisparities` at minimum. The second constructor enables you to set each parameter to a custom value.

## StereoSGBM::operator ()

**C++:** void StereoSGBM::**operator()**(InputArray **left**, InputArray **right**, OutputArray **disp**)

**Python:** cv2.StereoSGBM.**compute**(left, right[, disp ]) → disp

> Computes disparity using the SGBM algorithm for a rectified stereo pair.

> **Parameters**

>> **left** – Left 8-bit single-channel or 3-channel image.

>> **right** – Right image of the same size and the same type as the left one.

>> **disp** – Output disparity map. It is a 16-bit signed single-channel image of the same size as the input image. It contains disparity values scaled by 16. So, to get the floating-point disparity map, you need to divide each `disp` element by 16.

The method executes the SGBM algorithm on a rectified stereo pair. See `stereo_match.cpp` OpenCV sample on how to prepare images and call the method.

---

**Note:** The method is not constant, so you should not use the same `StereoSGBM` instance from different threads simultaneously.

---

## stereoCalibrate

Calibrates the stereo camera.

**C++:** double **stereoCalibrate**(InputArrayOfArrays **objectPoints**, InputArrayOfArrays **imagePoints1**, InputArrayOfArrays **imagePoints2**, InputOutputArray **cameraMatrix1**, InputOutputArray **distCoeffs1**, InputOutputArray **cameraMatrix2**, InputOutputArray **distCoeffs2**, Size **imageSize**, OutputArray **R**, OutputArray **T**, OutputArray **E**, OutputArray **F**, TermCriteria **criteria**=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 1e-6), int **flags**=CALIB_FIX_INTRINSIC )

**Python:** cv2.**stereoCalibrate**(objectPoints, imagePoints1, imagePoints2, imageSize[, cameraMatrix1[, distCoeffs1[, cameraMatrix2[, distCoeffs2[, R[, T[, E[, F[, criteria[, flags ]]]]]]]]]]) → retval, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, R, T, E, F

**C:** double **cvStereoCalibrate**(const CvMat* **object_points**, const CvMat* **image_points1**, const CvMat* **image_points2**, const CvMat* **npoints**, CvMat* **camera_matrix1**, CvMat* **dist_coeffs1**, CvMat* **camera_matrix2**, CvMat* **dist_coeffs2**, CvSize **image_size**, CvMat* **R**, CvMat* **T**, CvMat* **E**=0, CvMat* **F**=0, CvTermCriteria **term_crit**=cvTermCriteria( CV_TERMCRIT_ITER+CV_TERMCRIT_EPS,30,1e-6), int **flags**=CV_CALIB_FIX_INTRINSIC )

---

**Python:** cv.**StereoCalibrate**(objectPoints, imagePoints1, imagePoints2, pointCounts, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize, R, T, E=None, F=None, term_crit=(CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 30, 1e-6), flags=CV_CALIB_FIX_INTRINSIC) $\rightarrow$ None

> Parameters

>> **objectPoints** – Vector of vectors of the calibration pattern points.

>> **imagePoints1** – Vector of vectors of the projections of the calibration pattern points, observed by the first camera.

>> **imagePoints2** – Vector of vectors of the projections of the calibration pattern points, observed by the second camera.

>> **cameraMatrix1** – Input/output first camera matrix: $\begin{bmatrix} f_x^{(j)} & 0 & c_x^{(j)} \\ 0 & f_y^{(j)} & c_y^{(j)} \\ 0 & 0 & 1 \end{bmatrix}$ , $j = 0, 1$
>> . If any of `CV_CALIB_USE_INTRINSIC_GUESS` , `CV_CALIB_FIX_ASPECT_RATIO` , `CV_CALIB_FIX_INTRINSIC` , or `CV_CALIB_FIX_FOCAL_LENGTH` are specified, some or all of the matrix components must be initialized. See the flags description for details.

>> **distCoeffs1** – Input/output vector of distortion coefficients $(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6]])$ of 4, 5, or 8 elements. The output vector length depends on the flags.

>> **cameraMatrix2** – Input/output second camera matrix. The parameter is similar to `cameraMatrix1` .

>> **distCoeffs2** – Input/output lens distortion coefficients for the second camera. The parameter is similar to `distCoeffs1` .

>> **imageSize** – Size of the image used only to initialize intrinsic camera matrix.

>> **R** – Output rotation matrix between the 1st and the 2nd camera coordinate systems.

>> **T** – Output translation vector between the coordinate systems of the cameras.

>> **E** – Output essential matrix.

>> **F** – Output fundamental matrix.

>> **term_crit** – Termination criteria for the iterative optimization algorithm.

>> **flags** – Different flags that may be zero or a combination of the following values:

>> – **CV_CALIB_FIX_INTRINSIC** Fix `cameraMatrix?` and `distCoeffs?` so that only R, T, E , and F matrices are estimated.

>> – **CV_CALIB_USE_INTRINSIC_GUESS** Optimize some or all of the intrinsic parameters according to the specified flags. Initial values are provided by the user.

>> – **CV_CALIB_FIX_PRINCIPAL_POINT** Fix the principal points during the optimization.

>> – **CV_CALIB_FIX_FOCAL_LENGTH** Fix $f_x^{(j)}$ and $f_y^{(j)}$ .

>> – **CV_CALIB_FIX_ASPECT_RATIO** Optimize $f_y^{(j)}$ . Fix the ratio $f_x^{(j)}/f_y^{(j)}$ .

>> – **CV_CALIB_SAME_FOCAL_LENGTH** Enforce $f_x^{(0)} = f_x^{(1)}$ and $f_y^{(0)} = f_y^{(1)}$ .

>> – **CV_CALIB_ZERO_TANGENT_DIST** Set tangential distortion coefficients for each camera to zeros and fix there.

- **CV_CALIB_FIX_K1,...,CV_CALIB_FIX_K6** Do not change the corresponding radial distortion coefficient during the optimization. If CV_CALIB_USE_INTRINSIC_GUESS is set, the coefficient from the supplied `distCoeffs` matrix is used. Otherwise, it is set to 0.

- **CV_CALIB_RATIONAL_MODEL** Enable coefficients k4, k5, and k6. To provide the backward compatibility, this extra flag should be explicitly specified to make the calibration function use the rational model and return 8 coefficients. If the flag is not set, the function computes and returns only 5 distortion coefficients.

The function estimates transformation between two cameras making a stereo pair. If you have a stereo camera where the relative position and orientation of two cameras is fixed, and if you computed poses of an object relative to the first camera and to the second camera, (R1, T1) and (R2, T2), respectively (this can be done with `solvePnP()` ), then those poses definitely relate to each other. This means that, given ( $R_1$,:math:$T\_1$ ), it should be possible to compute ( $R_2$,:math:$T\_2$ ). You only need to know the position and orientation of the second camera relative to the first camera. This is what the described function does. It computes ( R,:math:$T$ ) so that:

$$R_2 = R * R_1 \quad T_2 = R * T_1 + T,$$

Optionally, it computes the essential matrix E:

$$E = \begin{bmatrix} 0 & -T_2 & T_1 \\ T_2 & 0 & -T_0 \\ -T_1 & T_0 & 0 \end{bmatrix} * R$$

where $T_i$ are components of the translation vector T : $T = [T_0, T_1, T_2]^T$ . And the function can also compute the fundamental matrix F:

$$F = cameraMatrix2^{-T} E cameraMatrix1^{-1}$$

Besides the stereo-related information, the function can also perform a full calibration of each of two cameras. However, due to the high dimensionality of the parameter space and noise in the input data, the function can diverge from the correct solution. If the intrinsic parameters can be estimated with high accuracy for each of the cameras individually (for example, using `calibrateCamera()` ), you are recommended to do so and then pass CV_CALIB_FIX_INTRINSIC flag to the function along with the computed intrinsic parameters. Otherwise, if all the parameters are estimated at once, it makes sense to restrict some parameters, for example, pass CV_CALIB_SAME_FOCAL_LENGTH and CV_CALIB_ZERO_TANGENT_DIST flags, which is usually a reasonable assumption.

Similarly to `calibrateCamera()` , the function minimizes the total re-projection error for all the points in all the available views from both cameras. The function returns the final value of the re-projection error.

## stereoRectify

Computes rectification transforms for each head of a calibrated stereo camera.

**C++:** void **stereoRectify**(InputArray **cameraMatrix1**, InputArray **distCoeffs1**, InputArray **cameraMatrix2**, InputArray **distCoeffs2**, Size **imageSize**, InputArray **R**, InputArray **T**, OutputArray **R1**, OutputArray **R2**, OutputArray **P1**, OutputArray **P2**, OutputArray **Q**, int **flags**=CALIB_ZERO_DISPARITY, double **alpha**=-1, Size **newImageSize**=Size(), Rect* **validPixROI1**=0, Rect* **validPixROI2**=0 )

**C:** void **cvStereoRectify**(const CvMat* **camera_matrix1**, const CvMat* **camera_matrix2**, const CvMat* **dist_coeffs1**, const CvMat* **dist_coeffs2**, CvSize **image_size**, const CvMat* **R**, const CvMat* **T**, CvMat* **R1**, CvMat* **R2**, CvMat* **P1**, CvMat* **P2**, CvMat* **Q**=0, int **flags**=CV_CALIB_ZERO_DISPARITY, double **alpha**=-1, CvSize **new_image_size**=cvSize(0,0), CvRect* **valid_pix_ROI1**=0, CvRect* **valid_pix_ROI2**=0 )

**Python:** cv.**StereoRectify**(cameraMatrix1, cameraMatrix2, distCoeffs1, distCoeffs2, imageSize, R, T, R1, R2, P1, P2, Q=None, flags=CV_CALIB_ZERO_DISPARITY, alpha=-1, newImageSize=(0, 0)) -> (roi1, roi2)

> Parameters

>> **cameraMatrix1** – First camera matrix.

>> **cameraMatrix2** – Second camera matrix.

>> **distCoeffs1** – First camera distortion parameters.

>> **distCoeffs2** – Second camera distortion parameters.

>> **imageSize** – Size of the image used for stereo calibration.

>> **R** – Rotation matrix between the coordinate systems of the first and the second cameras.

>> **T** – Translation vector between coordinate systems of the cameras.

>> **R1** – Output 3x3 rectification transform (rotation matrix) for the first camera.

>> **R2** – Output 3x3 rectification transform (rotation matrix) for the second camera.

>> **P1** – Output 3x4 projection matrix in the new (rectified) coordinate systems for the first camera.

>> **P2** – Output 3x4 projection matrix in the new (rectified) coordinate systems for the second camera.

>> **Q** – Output $4 \times 4$ disparity-to-depth mapping matrix (see `reprojectImageTo3D()` ).

>> **flags** – Operation flags that may be zero or `CV_CALIB_ZERO_DISPARITY` . If the flag is set, the function makes the principal points of each camera have the same pixel coordinates in the rectified views. And if the flag is not set, the function may still shift the images in the horizontal or vertical direction (depending on the orientation of epipolar lines) to maximize the useful image area.

>> **alpha** – Free scaling parameter. If it is -1 or absent, the function performs the default scaling. Otherwise, the parameter should be between 0 and 1. `alpha=0` means that the rectified images are zoomed and shifted so that only valid pixels are visible (no black areas after rectification). `alpha=1` means that the rectified image is decimated and shifted so that all the pixels from the original images from the cameras are retained in the rectified images (no source image pixels are lost). Obviously, any intermediate value yields an intermediate result between those two extreme cases.

>> **newImageSize** – New image resolution after rectification. The same size should be passed to `initUndistortRectifyMap()` (see the `stereo_calib.cpp` sample in OpenCV samples directory). When (0,0) is passed (default), it is set to the original `imageSize` . Setting it to larger value can help you preserve details in the original image, especially when there is a big radial distortion.

>> **validPixROI1** – Optional output rectangles inside the rectified images where all the pixels are valid. If `alpha=0` , the ROIs cover the whole images. Otherwise, they are likely to be smaller (see the picture below).

>> **validPixROI2** – Optional output rectangles inside the rectified images where all the pixels are valid. If `alpha=0` , the ROIs cover the whole images. Otherwise, they are likely to be smaller (see the picture below).

The function computes the rotation matrices for each camera that (virtually) make both camera image planes the same plane. Consequently, this makes all the epipolar lines parallel and thus simplifies the dense stereo correspondence problem. The function takes the matrices computed by `stereoCalibrate()` as input. As output, it provides two

rotation matrices and also two projection matrices in the new coordinates. The function distinguishes the following two cases:

1. **Horizontal stereo**: the first and the second camera views are shifted relative to each other mainly along the x axis (with possible small vertical shift). In the rectified images, the corresponding epipolar lines in the left and right cameras are horizontal and have the same y-coordinate. P1 and P2 look like:

$$\texttt{P1} = \begin{bmatrix} f & 0 & cx_1 & 0 \\ 0 & f & cy & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

$$\texttt{P2} = \begin{bmatrix} f & 0 & cx_2 & T_x * f \\ 0 & f & cy & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix},$$

where $T_x$ is a horizontal shift between the cameras and $cx_1 = cx_2$ if `CV_CALIB_ZERO_DISPARITY` is set.
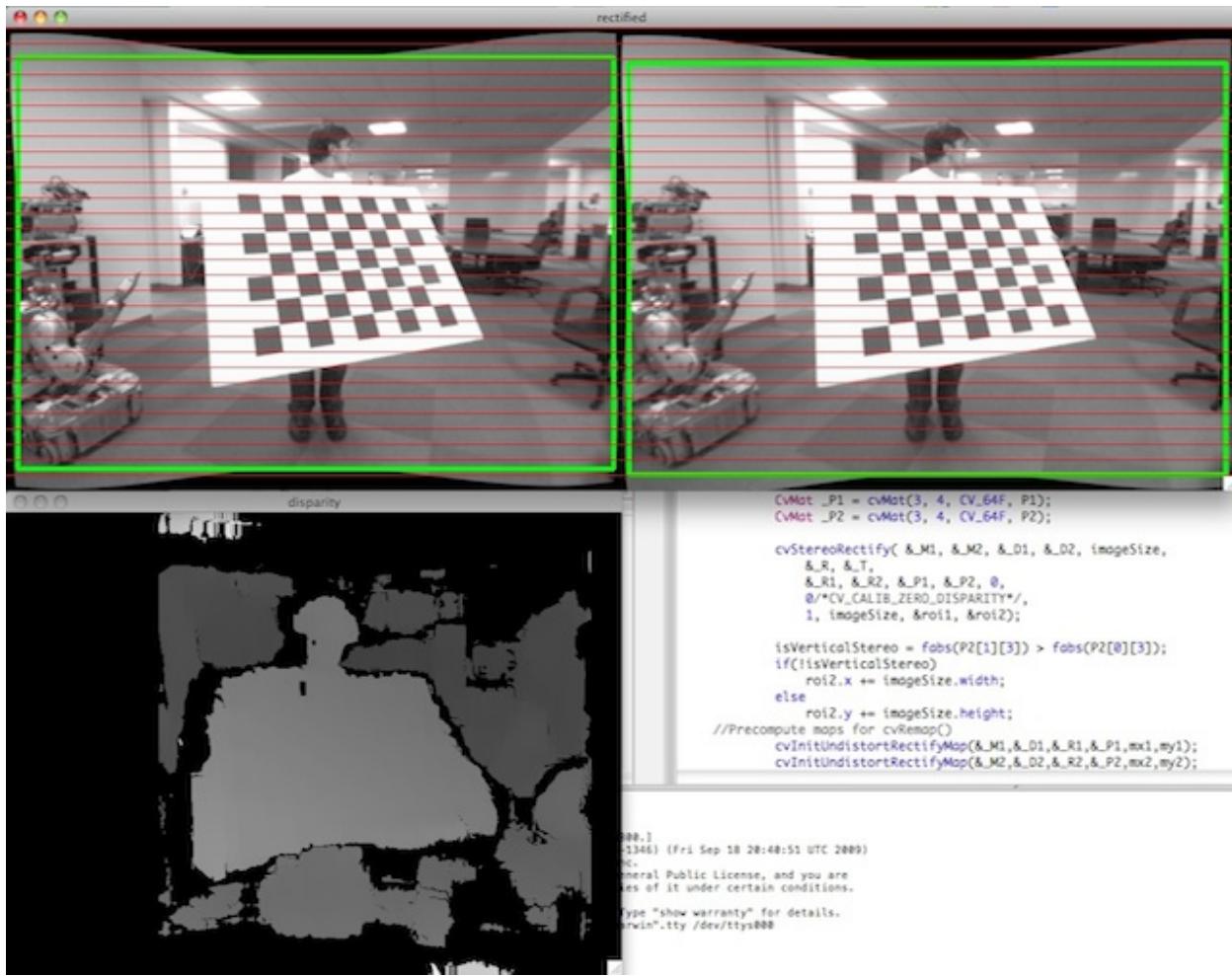
2. **Vertical stereo**: the first and the second camera views are shifted relative to each other mainly in vertical direction (and probably a bit in the horizontal direction too). The epipolar lines in the rectified images are vertical and have the same x-coordinate. P1 and P2 look like:

$$\texttt{P1} = \begin{bmatrix} f & 0 & cx & 0 \\ 0 & f & cy_1 & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

$$\texttt{P2} = \begin{bmatrix} f & 0 & cx & 0 \\ 0 & f & cy_2 & T_y * f \\ 0 & 0 & 1 & 0 \end{bmatrix},$$

where $T_y$ is a vertical shift between the cameras and $cy_1 = cy_2$ if `CALIB_ZERO_DISPARITY` is set.

As you can see, the first three columns of `P1` and `P2` will effectively be the new "rectified" camera matrices. The matrices, together with `R1` and `R2`, can then be passed to `initUndistortRectifyMap()` to initialize the rectification map for each camera.

See below the screenshot from the `stereo_calib.cpp` sample. Some red horizontal lines pass through the corresponding image regions. This means that the images are well rectified, which is what most stereo correspondence algorithms rely on. The green rectangles are `roi1` and `roi2`. You see that their interiors are all valid pixels.

## stereoRectifyUncalibrated

Computes a rectification transform for an uncalibrated stereo camera.

**C++:** bool **stereoRectifyUncalibrated**(InputArray **points1**, InputArray **points2**, InputArray **F**, Size **imgSize**, OutputArray **H1**, OutputArray **H2**, double **threshold**=5 )

**Python:** cv2.**stereoRectifyUncalibrated**(points1, points2, F, imgSize[, H1[, H2[, threshold ] ] ]) → retval, H1, H2

**C:** int **cvStereoRectifyUncalibrated**(const CvMat* **points1**, const CvMat* **points2**, const CvMat* **F**, CvSize **img_size**, CvMat* **H1**, CvMat* **H2**, double **threshold**=5 )

**Python:** cv.**StereoRectifyUncalibrated**(points1, points2, F, imageSize, H1, H2, threshold=5) → None

> **Parameters**
>
>> **points1** – Array of feature points in the first image.
>>
>> **points2** – The corresponding points in the second image. The same formats as in `findFundamentalMat()` are supported.
>>
>> **F** – Input fundamental matrix. It can be computed from the same set of point pairs using `findFundamentalMat()` .
>>
>> **imgSize** – Size of the image.

**H1** – Output rectification homography matrix for the first image.

**H2** – Output rectification homography matrix for the second image.

**threshold** – Optional threshold used to filter out the outliers. If the parameter is greater than zero, all the point pairs that do not comply with the epipolar geometry (that is, the points for which $|\text{points2[i]}^T * F * \text{points1[i]}| > \text{threshold}$) are rejected prior to computing the homographies. Otherwise, all the points are considered inliers.

The function computes the rectification transformations without knowing intrinsic parameters of the cameras and their relative position in the space, which explains the suffix "uncalibrated". Another related difference from `stereoRectify()` is that the function outputs not the rectification transformations in the object (3D) space, but the planar perspective transformations encoded by the homography matrices H1 and H2 . The function implements the algorithm [Hartley99].

---

**Note:** While the algorithm does not need to know the intrinsic parameters of the cameras, it heavily depends on the epipolar geometry. Therefore, if the camera lenses have a significant distortion, it would be better to correct it before computing the fundamental matrix and calling this function. For example, distortion coefficients can be estimated for each head of stereo camera separately by using `calibrateCamera()` . Then, the images can be corrected using `undistort()` , or just the point coordinates can be corrected with `undistortPoints()` .

## triangulatePoints

Reconstructs points by triangulation.

**C++:** void **triangulatePoints**(InputArray **projMatr1**, InputArray **projMatr2**, InputArray **projPoints1**, InputArray **projPoints2**, OutputArray **points4D**)

**Python:** cv2.**triangulatePoints**(projMatr1, projMatr2, projPoints1, projPoints2[, points4D]) → points4D

**C:** void **cvTriangulatePoints**(CvMat* **projMatr1**, CvMat* **projMatr2**, CvMat* **projPoints1**, CvMat* **projPoints2**, CvMat* **points4D**)

> Parameters

> > **projMatr1** – 3x4 projection matrix of the first camera.

> > **projMatr2** – 3x4 projection matrix of the second camera.

> > **projPoints1** – 2xN array of feature points in the first image. In case of c++ version it can be also a vector of feature points or two-channel matrix of size 1xN or Nx1.

> > **projPoints2** – 2xN array of corresponding points in the second image. In case of c++ version it can be also a vector of feature points or two-channel matrix of size 1xN or Nx1.

> > **points4D** – 4xN array of reconstructed points in homogeneous coordinates.

The function reconstructs 3-dimensional points (in homogeneous coordinates) by using their observations with a stereo camera. Projections matrices can be obtained from `stereoRectify()`.

**See Also:**

`reprojectImageTo3D()`

# FEATURES2D. 2D FEATURES FRAMEWORK

## 7.1 Feature Detection and Description

### FAST

Detects corners using the FAST algorithm

**C++:** void **FAST**(InputArray **image**, vector<KeyPoint>& **keypoints**, int **threshold**, bool **nonmaxSupression**=true )

**C++:** void **FASTX**(InputArray **image**, vector<KeyPoint>& **keypoints**, int **threshold**, bool **nonmaxSupression**, int **type**)

> **Parameters**
>
>> **image** – grayscale image where keypoints (corners) are detected.
>>
>> **keypoints** – keypoints detected on the image.
>>
>> **threshold** – threshold on difference between intensity of the central pixel and pixels of a circle around this pixel.
>>
>> **nonmaxSupression** – if true, non-maximum suppression is applied to detected corners (keypoints).
>>
>> **type** – one of the three neighborhoods as defined in the paper: FastFeatureDetector::TYPE_9_16, FastFeatureDetector::TYPE_7_12, FastFeatureDetector::TYPE_5_8

Detects corners using the FAST algorithm by [Rosten06].

### MSER

**class MSER** : **public FeatureDetector**

Maximally stable extremal region extractor.

```
class MSER : public CvMSERParams
{
public:
    // default constructor
    MSER();
```

```
    // constructor that initializes all the algorithm parameters
    MSER( int _delta, int _min_area, int _max_area,
          float _max_variation, float _min_diversity,
          int _max_evolution, double _area_threshold,
          double _min_margin, int _edge_blur_size );
    // runs the extractor on the specified image; returns the MSERs,
    // each encoded as a contour (vector<Point>, see findContours)
    // the optional mask marks the area where MSERs are searched for
    void operator()( const Mat& image, vector<vector<Point> >& msers, const Mat& mask ) const;
};
```

The class encapsulates all the parameters of the MSER extraction algorithm (see http://en.wikipedia.org/wiki/Maximally_stable_extremal_regions). Also see http://opencv.willowgarage.com/wiki/documentation/cpp/features2d/MSER for useful comments and parameters description.

## ORB

**class ORB** : **public Feature2D**

Class implementing the ORB (*oriented BRIEF*) keypoint detector and descriptor extractor, described in [RRKB11]. The algorithm uses FAST in pyramids to detect stable keypoints, selects the strongest features using FAST or Harris response, finds their orientation using first-order moments and computes the descriptors using BRIEF (where the coordinates of random point pairs (or k-tuples) are rotated according to the measured orientation).

## ORB::ORB

The ORB constructor

**C++:** ORB::**ORB**(int **nfeatures**=500, float **scaleFactor**=1.2f, int **nlevels**=8, int **edgeThreshold**=31, int **firstLevel**=0, int **WTA_K**=2, int **scoreType**=ORB::HARRIS_SCORE, int **patchSize**=31)

> **Parameters**
>
>> **nfeatures** – The maximum number of features to retain.
>>
>> **scaleFactor** – Pyramid decimation ratio, greater than 1. `scaleFactor==2` means the classical pyramid, where each next level has 4x less pixels than the previous, but such a big scale factor will degrade feature matching scores dramatically. On the other hand, too close to 1 scale factor will mean that to cover certain scale range you will need more pyramid levels and so the speed will suffer.
>>
>> **nlevels** – The number of pyramid levels. The smallest level will have linear size equal to `input_image_linear_size/pow(scaleFactor, nlevels)`.
>>
>> **edgeThreshold** – This is size of the border where the features are not detected. It should roughly match the `patchSize` parameter.
>>
>> **firstLevel** – It should be 0 in the current implementation.
>>
>> **WTA_K** – The number of points that produce each element of the oriented BRIEF descriptor. The default value 2 means the BRIEF where we take a random point pair and compare their brightnesses, so we get 0/1 response. Other possible values are 3 and 4. For example, 3 means that we take 3 random points (of course, those point coordinates are random, but they are generated from the pre-defined seed, so each element of BRIEF descriptor is computed deterministically from the pixel rectangle), find point of maximum brightness and output index of the winner (0, 1 or 2). Such output will occupy 2 bits, and therefore it will need

a special variant of Hamming distance, denoted as `NORM_HAMMING2` (2 bits per bin). When `WTA_K=4`, we take 4 random points to compute each bin (that will also occupy 2 bits with possible values 0, 1, 2 or 3).

**scoreType** – The default HARRIS_SCORE means that Harris algorithm is used to rank features (the score is written to `KeyPoint::score` and is used to retain best `nfeatures` features); FAST_SCORE is alternative value of the parameter that produces slightly less stable keypoints, but it is a little faster to compute.

**patchSize** – size of the patch used by the oriented BRIEF descriptor. Of course, on smaller pyramid layers the perceived image area covered by a feature will be larger.

## ORB::operator()

Finds keypoints in an image and computes their descriptors

**C++:** void ORB::**operator()**(InputArray **image**, InputArray **mask**, vector<KeyPoint>& **keypoints**, OutputArray **descriptors**, bool **useProvidedKeypoints**=false ) const

> **Parameters**

>> **image** – The input 8-bit grayscale image.

>> **mask** – The operation mask.

>> **keypoints** – The output vector of keypoints.

>> **descriptors** – The output descriptors. Pass `cv::noArray()` if you do not need it.

>> **useProvidedKeypoints** – If it is true, then the method will use the provided vector of keypoints instead of detecting them.

## FREAK

**class FREAK** : **public DescriptorExtractor**

Class implementing the FREAK (*Fast Retina Keypoint*) keypoint descriptor, described in [AOV12]. The algorithm propose a novel keypoint descriptor inspired by the human visual system and more precisely the retina, coined Fast Retina Key- point (FREAK). A cascade of binary strings is computed by efficiently comparing image intensities over a retinal sampling pattern. FREAKs are in general faster to compute with lower memory load and also more robust than SIFT, SURF or BRISK. They are competitive alternatives to existing keypoints in particular for embedded applications.

## FREAK::FREAK

The FREAK constructor

**C++:** FREAK::**FREAK**(bool **orientationNormalized**=true, bool **scaleNormalized**=true, float **patternScale**=22.0f, int **nOctaves**=4, const vector<int>& **selectedPairs**=vector<int>() )

> **Parameters**

>> **orientationNormalized** – Enable orientation normalization.

>> **scaleNormalized** – Enable scale normalization.

>> **patternScale** – Scaling of the description pattern.

>> **nOctaves** – Number of octaves covered by the detected keypoints.

>> **selectedPairs** – (Optional) user defined selected pairs indexes,

## FREAK::selectPairs

Select the 512 best description pair indexes from an input (grayscale) image set. FREAK is available with a set of pairs learned off-line. Researchers can run a training process to learn their own set of pair. For more details read section 4.2 in: A. Alahi, R. Ortiz, and P. Vandergheynst. FREAK: Fast Retina Keypoint. In IEEE Conference on Computer Vision and Pattern Recognition, 2012.

We notice that for keypoint matching applications, image content has little effect on the selected pairs unless very specific what does matter is the detector type (blobs, corners,...) and the options used (scale/rotation invariance,...). Reduce corrThresh if not enough pairs are selected (43 points –> 903 possible pairs)

**C++:** vector<int> FREAK::**selectPairs**(const vector<Mat>& **images**, vector<vector<KeyPoint>>& **keypoints**, const double **corrThresh**=0.7, bool **verbose**=true)

> **Parameters**
>
> > **images** – Grayscale image input set.
> >
> > **keypoints** – Set of detected keypoints
> >
> > **corrThresh** – Correlation threshold.
> >
> > **verbose** – Prints pair selection informations.

## 7.2 Common Interfaces of Feature Detectors

Feature detectors in OpenCV have wrappers with a common interface that enables you to easily switch between different algorithms solving the same problem. All objects that implement keypoint detectors inherit the `FeatureDetector` interface.

## KeyPoint

class **KeyPoint**
Data structure for salient point detectors.

Point2f **pt**
coordinates of the keypoint

float **size**
diameter of the meaningful keypoint neighborhood

float **angle**
computed orientation of the keypoint (-1 if not applicable). Its possible values are in a range [0,360) degrees. It is measured relative to image coordinate system (y-axis is directed downward), ie in clockwise.

float **response**
the response by which the most strong keypoints have been selected. Can be used for further sorting or subsampling

int **octave**
octave (pyramid layer) from which the keypoint has been extracted

int **class_id**
object id that can be used to clustered keypoints by an object they belong to

## KeyPoint::KeyPoint

The keypoint constructors

**C++:** KeyPoint::**KeyPoint**()

**C++:** KeyPoint::**KeyPoint**(Point2f **_pt**, float **_size**, float **_angle**=-1, float **_response**=0, int **_octave**=0, int **_class_id**=-1)

**C++:** KeyPoint::**KeyPoint**(float **x**, float **y**, float **_size**, float **_angle**=-1, float **_response**=0, int **_octave**=0, int **_class_id**=-1)

**Python:** cv2.**KeyPoint**([x, y, _size[, _angle[, _response[, _octave[, _class_id ]]]]]) → <KeyPoint object>

> **Parameters**
>
> > **x** – x-coordinate of the keypoint
> >
> > **y** – y-coordinate of the keypoint
> >
> > **_pt** – x & y coordinates of the keypoint
> >
> > **_size** – keypoint diameter
> >
> > **_angle** – keypoint orientation
> >
> > **_response** – keypoint detector response on the keypoint (that is, strength of the keypoint)
> >
> > **_octave** – pyramid octave in which the keypoint has been detected
> >
> > **_class_id** – object id

## FeatureDetector

class **FeatureDetector** : public **Algorithm**

Abstract base class for 2D image feature detectors.

```
class CV_EXPORTS FeatureDetector
{
public:
    virtual ~FeatureDetector();

    void detect( const Mat& image, vector<KeyPoint>& keypoints,
                 const Mat& mask=Mat() ) const;

    void detect( const vector<Mat>& images,
                 vector<vector<KeyPoint> >& keypoints,
                 const vector<Mat>& masks=vector<Mat>() ) const;

    virtual void read(const FileNode&);
    virtual void write(FileStorage&) const;

    static Ptr<FeatureDetector> create( const string& detectorType );

protected:
...
};
```

## FeatureDetector::detect

Detects keypoints in an image (first variant) or image set (second variant).

**C++:** void FeatureDetector::**detect**(const Mat& **image**, vector<KeyPoint>& **keypoints**, const Mat& **mask**=Mat() ) const

**C++:** void FeatureDetector::**detect**(const vector<Mat>& **images**, vector<vector<KeyPoint>>& **keypoints**, const vector<Mat>& **masks**=vector<Mat>() ) const

> **Parameters**

>> **image** – Image.

>> **images** – Image set.

>> **keypoints** – The detected keypoints. In the second variant of the method keypoints[i] is a set of keypoints detected in images[i] .

>> **mask** – Mask specifying where to look for keypoints (optional). It must be a 8-bit integer matrix with non-zero values in the region of interest.

>> **masks** – Masks for each input image specifying where to look for keypoints (optional). masks[i] is a mask for images[i].

## FeatureDetector::create

Creates a feature detector by its name.

**C++:** Ptr<FeatureDetector> FeatureDetector::**create**(const string& **detectorType**)

> **Parameters**

>> **detectorType** – Feature detector type.

The following detector types are supported:

- "FAST" – `FastFeatureDetector`

- "STAR" – `StarFeatureDetector`

- "SIFT" – `SIFT` (nonfree module)

- "SURF" – `SURF` (nonfree module)

- "ORB" – `ORB`

- "MSER" – `MSER`

- "GFTT" – `GoodFeaturesToTrackDetector`

- "HARRIS" – `GoodFeaturesToTrackDetector` with Harris detector enabled

- "Dense" – `DenseFeatureDetector`

- "SimpleBlob" – `SimpleBlobDetector`

Also a combined format is supported: feature detector adapter name ( "Grid" – `GridAdaptedFeatureDetector`, "Pyramid" – `PyramidAdaptedFeatureDetector` ) + feature detector name (see above), for example: "GridFAST", "PyramidSTAR" .

## FastFeatureDetector

**class FastFeatureDetector** : **public FeatureDetector**

Wrapping class for feature detection using the FAST() method.

```
class FastFeatureDetector : public FeatureDetector
{
public:
    FastFeatureDetector( int threshold=1, bool nonmaxSuppression=true, type=FastFeatureDetector::TYPE_9_16 );
    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

## GoodFeaturesToTrackDetector

**class GoodFeaturesToTrackDetector** : **public FeatureDetector**

Wrapping class for feature detection using the goodFeaturesToTrack() function.

```
class GoodFeaturesToTrackDetector : public FeatureDetector
{
public:
    class Params
    {
    public:
        Params( int maxCorners=1000, double qualityLevel=0.01,
                double minDistance=1., int blockSize=3,
                bool useHarrisDetector=false, double k=0.04 );
        void read( const FileNode& fn );
        void write( FileStorage& fs ) const;

        int maxCorners;
        double qualityLevel;
        double minDistance;
        int blockSize;
        bool useHarrisDetector;
        double k;
    };

    GoodFeaturesToTrackDetector( const GoodFeaturesToTrackDetector::Params& params=
                                        GoodFeaturesToTrackDetector::Params() );
    GoodFeaturesToTrackDetector( int maxCorners, double qualityLevel,
                                 double minDistance, int blockSize=3,
                                 bool useHarrisDetector=false, double k=0.04 );
    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

## MserFeatureDetector

**class MserFeatureDetector** : **public FeatureDetector**

Wrapping class for feature detection using the MSER class.

```
class MserFeatureDetector : public FeatureDetector
{
public:
    MserFeatureDetector( CvMSERParams params=cvMSERParams() );
    MserFeatureDetector( int delta, int minArea, int maxArea,
                         double maxVariation, double minDiversity,
                         int maxEvolution, double areaThreshold,
                         double minMargin, int edgeBlurSize );
    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

## StarFeatureDetector

### class StarFeatureDetector : public FeatureDetector

Wrapping class for feature detection using the StarDetector class.

```
class StarFeatureDetector : public FeatureDetector
{
public:
    StarFeatureDetector( int maxSize=16, int responseThreshold=30,
                         int lineThresholdProjected = 10,
                         int lineThresholdBinarized=8, int suppressNonmaxSize=5 );
    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

## DenseFeatureDetector

### class DenseFeatureDetector : public FeatureDetector

Class for generation of image features which are distributed densely and regularly over the image.

```
    class DenseFeatureDetector : public FeatureDetector
    {
    public:
            DenseFeatureDetector( float initFeatureScale=1.f, int featureScaleLevels=1,
                         float featureScaleMul=0.1f,
                         int initXyStep=6, int initImgBound=0,
                         bool varyXyStepWithScale=true,
                         bool varyImgBoundWithScale=false );
    protected:
    ...
};
```

The detector generates several levels (in the amount of featureScaleLevels) of features. Features of each level are located in the nodes of a regular grid over the image (excluding the image boundary of given size). The level parameters (a feature scale, a node size, a size of boundary) are multiplied by featureScaleMul with level index growing depending on input flags, viz.:

- Feature scale is multiplied always.

- The grid node size is multiplied if `varyXyStepWithScale` is `true`.

- Size of image boundary is multiplied if `varyImgBoundWithScale` is `true`.

## SimpleBlobDetector

**class** `SimpleBlobDetector` : **public** `FeatureDetector`

Class for extracting blobs from an image.

```
class SimpleBlobDetector : public FeatureDetector
{
public:
struct Params
{
    Params();
    float thresholdStep;
    float minThreshold;
    float maxThreshold;
    size_t minRepeatability;
    float minDistBetweenBlobs;

    bool filterByColor;
    uchar blobColor;

    bool filterByArea;
    float minArea, maxArea;

    bool filterByCircularity;
    float minCircularity, maxCircularity;

    bool filterByInertia;
    float minInertiaRatio, maxInertiaRatio;

    bool filterByConvexity;
    float minConvexity, maxConvexity;
};

SimpleBlobDetector(const SimpleBlobDetector::Params &parameters = SimpleBlobDetector::Params());

protected:
    ...
};
```

The class implements a simple algorithm for extracting blobs from an image:

1. Convert the source image to binary images by applying thresholding with several thresholds from `minThreshold` (inclusive) to `maxThreshold` (exclusive) with distance `thresholdStep` between neighboring thresholds.

2. Extract connected components from every binary image by `findContours()` and calculate their centers.

3. Group centers from several binary images by their coordinates. Close centers form one group that corresponds to one blob, which is controlled by the `minDistBetweenBlobs` parameter.

4. From the groups, estimate final centers of blobs and their radiuses and return as locations and sizes of keypoints.

This class performs several filtrations of returned blobs. You should set `filterBy*` to true/false to turn on/off corresponding filtration. Available filtrations:

- **By color**. This filter compares the intensity of a binary image at the center of a blob to `blobColor`. If they differ, the blob is filtered out. Use `blobColor = 0` to extract dark blobs and `blobColor = 255` to extract light blobs.

- **By area**. Extracted blobs have an area between `minArea` (inclusive) and `maxArea` (exclusive).

- **By circularity**. Extracted blobs have circularity ($\frac{4*\pi*Area}{perimeter*perimeter}$) between `minCircularity` (inclusive) and `maxCircularity` (exclusive).

- **By ratio of the minimum inertia to maximum inertia**. Extracted blobs have this ratio between `minInertiaRatio` (inclusive) and `maxInertiaRatio` (exclusive).

- **By convexity**. Extracted blobs have convexity (area / area of blob convex hull) between `minConvexity` (inclusive) and `maxConvexity` (exclusive).

Default values of parameters are tuned to extract dark circular blobs.

## GridAdaptedFeatureDetector

**class GridAdaptedFeatureDetector** : **public FeatureDetector**

Class adapting a detector to partition the source image into a grid and detect points in each cell.

```
class GridAdaptedFeatureDetector : public FeatureDetector
{
public:
    /*
     * detector           Detector that will be adapted.
     * maxTotalKeypoints  Maximum count of keypoints detected on the image.
     *                    Only the strongest keypoints will be kept.
     * gridRows           Grid row count.
     * gridCols           Grid column count.
     */
    GridAdaptedFeatureDetector( const Ptr<FeatureDetector>& detector,
                                int maxTotalKeypoints, int gridRows=4,
                                int gridCols=4 );
    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

## PyramidAdaptedFeatureDetector

**class PyramidAdaptedFeatureDetector** : **public FeatureDetector**

Class adapting a detector to detect points over multiple levels of a Gaussian pyramid. Consider using this class for detectors that are not inherently scaled.

```
class PyramidAdaptedFeatureDetector : public FeatureDetector
{
public:
    PyramidAdaptedFeatureDetector( const Ptr<FeatureDetector>& detector,
                                   int levels=2 );
    virtual void read( const FileNode& fn );
```

```cpp
    virtual void write( FileStorage& fs ) const;
protected:
    ...
};
```

# DynamicAdaptedFeatureDetector

**class DynamicAdaptedFeatureDetector : public FeatureDetector**

Adaptively adjusting detector that iteratively detects features until the desired number is found.

```cpp
class DynamicAdaptedFeatureDetector: public FeatureDetector
{
public:
    DynamicAdaptedFeatureDetector( const Ptr<AdjusterAdapter>& adjuster,
        int min_features=400, int max_features=500, int max_iters=5 );
    ...
};
```

If the detector is persisted, it "remembers" the parameters used for the last detection. In this case, the detector may be used for consistent numbers of keypoints in a set of temporally related images, such as video streams or panorama series.

DynamicAdaptedFeatureDetector uses another detector, such as FAST or SURF, to do the dirty work, with the help of AdjusterAdapter. If the detected number of features is not large enough, AdjusterAdapter adjusts the detection parameters so that the next detection results in a bigger or smaller number of features. This is repeated until either the number of desired features are found or the parameters are maxed out.

Adapters can be easily implemented for any detector via the AdjusterAdapter interface.

Beware that this is not thread-safe since the adjustment of parameters requires modification of the feature detector class instance.

Example of creating DynamicAdaptedFeatureDetector :

```cpp
//sample usage:
//will create a detector that attempts to find
//100 - 110 FAST Keypoints, and will at most run
//FAST feature detection 10 times until that
//number of keypoints are found
Ptr<FeatureDetector> detector(new DynamicAdaptedFeatureDetector (100, 110, 10,
                              new FastAdjuster(20,true)));
```

# DynamicAdaptedFeatureDetector::DynamicAdaptedFeatureDetector

The constructor

**C++:** DynamicAdaptedFeatureDetector::**DynamicAdaptedFeatureDetector**(const
Ptr<AdjusterAdapter>&
**adjuster**, int
**min_features**=400,
int **max_features**=500,
int **max_iters**=5 )

>        **Parameters**

>                **adjuster** – AdjusterAdapter that detects features and adjusts parameters.

---

**min_features** – Minimum desired number of features.

**max_features** – Maximum desired number of features.

**max_iters** – Maximum number of times to try adjusting the feature detector parameters. For `FastAdjuster` , this number can be high, but with `Star` or `Surf` many iterations can be time-consuming. At each iteration the detector is rerun.

## AdjusterAdapter

**class AdjusterAdapter : public FeatureDetector**

Class providing an interface for adjusting parameters of a feature detector. This interface is used by `DynamicAdaptedFeatureDetector` . It is a wrapper for `FeatureDetector` that enables adjusting parameters after feature detection.

```cpp
class AdjusterAdapter: public FeatureDetector
{
public:
    virtual ~AdjusterAdapter() {}
    virtual void tooFew(int min, int n_detected) = 0;
    virtual void tooMany(int max, int n_detected) = 0;
    virtual bool good() const = 0;
    virtual Ptr<AdjusterAdapter> clone() const = 0;
    static Ptr<AdjusterAdapter> create( const string& detectorType );
};
```

See `FastAdjuster`, `StarAdjuster`, and `SurfAdjuster` for concrete implementations.

## AdjusterAdapter::tooFew

Adjusts the detector parameters to detect more features.

**C++:** void AdjusterAdapter::**tooFew**(int **min**, int **n_detected**)

> **Parameters**
>
> > **min** – Minimum desired number of features.
> >
> > **n_detected** – Number of features detected during the latest run.

Example:

```cpp
void FastAdjuster::tooFew(int min, int n_detected)
{
        thresh_--;
}
```

## AdjusterAdapter::tooMany

Adjusts the detector parameters to detect less features.

**C++:** void AdjusterAdapter::**tooMany**(int **max**, int **n_detected**)

> **Parameters**
>
> > **max** – Maximum desired number of features.
> >
> > **n_detected** – Number of features detected during the latest run.

Example:

```
void FastAdjuster::tooMany(int min, int n_detected)
{
        thresh_++;
}
```

## AdjusterAdapter::good

Returns false if the detector parameters cannot be adjusted any more.

**C++:** bool AdjusterAdapter::**good**() const

Example:

```
bool FastAdjuster::good() const
{
        return (thresh_ > 1) && (thresh_ < 200);
}
```

## AdjusterAdapter::create

Creates an adjuster adapter by name

**C++:** Ptr<AdjusterAdapter> AdjusterAdapter::**create**(const string& **detectorType**)
Creates an adjuster adapter by name detectorType. The detector name is the same as in FeatureDetector::create(), but now supports "FAST", "STAR", and "SURF" only.

## FastAdjuster

**class FastAdjuster** : **public AdjusterAdapter**

AdjusterAdapter for FastFeatureDetector. This class decreases or increases the threshold value by 1.

```
class FastAdjuster FastAdjuster: public AdjusterAdapter
{
public:
        FastAdjuster(int init_thresh = 20, bool nonmax = true);
        ...
};
```

## StarAdjuster

**class StarAdjuster** : **public AdjusterAdapter**

AdjusterAdapter for StarFeatureDetector. This class adjusts the responseThreshhold of StarFeatureDetector.

```
class StarAdjuster: public AdjusterAdapter
{
        StarAdjuster(double initial_thresh = 30.0);
        ...
};
```

# 7.3 Common Interfaces of Descriptor Extractors

Extractors of keypoint descriptors in OpenCV have wrappers with a common interface that enables you to easily switch between different algorithms solving the same problem. This section is devoted to computing descriptors represented as vectors in a multidimensional space. All objects that implement the `vector` descriptor extractors inherit the `DescriptorExtractor` interface.

## DescriptorExtractor

class **DescriptorExtractor** : public **Algorithm**

Abstract base class for computing descriptors for image keypoints.

```
class CV_EXPORTS DescriptorExtractor
{
public:
    virtual ~DescriptorExtractor();

    void compute( const Mat& image, vector<KeyPoint>& keypoints,
                  Mat& descriptors ) const;
    void compute( const vector<Mat>& images, vector<vector<KeyPoint> >& keypoints,
                  vector<Mat>& descriptors ) const;

    virtual void read( const FileNode& );
    virtual void write( FileStorage& ) const;

    virtual int descriptorSize() const = 0;
    virtual int descriptorType() const = 0;

    static Ptr<DescriptorExtractor> create( const string& descriptorExtractorType );

protected:
    ...
};
```

In this interface, a keypoint descriptor can be represented as a dense, fixed-dimension vector of a basic type. Most descriptors follow this pattern as it simplifies computing distances between descriptors. Therefore, a collection of descriptors is represented as `Mat` , where each row is a keypoint descriptor.

## DescriptorExtractor::compute

Computes the descriptors for a set of keypoints detected in an image (first variant) or image set (second variant).

**C++:** void DescriptorExtractor::**compute**(const Mat& **image**, vector<KeyPoint>& **keypoints**, Mat& **descriptors**) const

**C++:** void DescriptorExtractor::**compute**(const vector<Mat>& **images**, vector<vector<KeyPoint>>& **keypoints**, vector<Mat>& **descriptors**) const

> **Parameters**
>
>> **image** – Image.
>>
>> **images** – Image set.
>>
>> **keypoints** – Input collection of keypoints. Keypoints for which a descriptor cannot be computed are removed. Sometimes new keypoints can be added, for example: SIFT duplicates keypoint with several dominant orientations (for each orientation).

> **descriptors** – Computed descriptors. In the second variant of the method `descriptors[i]` are descriptors computed for a `keypoints[i]'`. Row ''j is the `keypoints` (or `keypoints[i]`) is the descriptor for keypoint j-th keypoint.

## DescriptorExtractor::create

Creates a descriptor extractor by name.

**C++:** Ptr<DescriptorExtractor> DescriptorExtractor::**create**(const string& **descriptorExtractorType**)

> **Parameters**
>
> > **descriptorExtractorType** – Descriptor extractor type.

The current implementation supports the following types of a descriptor extractor:

- `"SIFT"` – SIFT

- `"SURF"` – SURF

- `"ORB"` – ORB

- `"BRIEF"` – BriefDescriptorExtractor

A combined format is also supported: descriptor extractor adapter name ( `"Opponent"` – OpponentColorDescriptorExtractor ) + descriptor extractor name (see above), for example: `"OpponentSIFT"` .

## OpponentColorDescriptorExtractor

class **OpponentColorDescriptorExtractor** : **public** **DescriptorExtractor**

Class adapting a descriptor extractor to compute descriptors in the Opponent Color Space (refer to Van de Sande et al., CGIV 2008 *Color Descriptors for Object Category Recognition*). Input RGB image is transformed in the Opponent Color Space. Then, an unadapted descriptor extractor (set in the constructor) computes descriptors on each of three channels and concatenates them into a single color descriptor.

```
class OpponentColorDescriptorExtractor : public DescriptorExtractor
{
public:
    OpponentColorDescriptorExtractor( const Ptr<DescriptorExtractor>& dextractor );

    virtual void read( const FileNode& );
    virtual void write( FileStorage& ) const;
    virtual int descriptorSize() const;
    virtual int descriptorType() const;
protected:
    ...
};
```

## BriefDescriptorExtractor

class **BriefDescriptorExtractor** : **public** **DescriptorExtractor**

Class for computing BRIEF descriptors described in a paper of Calonder M., Lepetit V., Strecha C., Fua P. *BRIEF: Binary Robust Independent Elementary Features* , 11th European Conference on Computer Vision (ECCV), Heraklion, Crete. LNCS Springer, September 2010.

```
class BriefDescriptorExtractor : public DescriptorExtractor
{
public:
    static const int PATCH_SIZE = 48;
    static const int KERNEL_SIZE = 9;

    // bytes is a length of descriptor in bytes. It can be equal 16, 32 or 64 bytes.
    BriefDescriptorExtractor( int bytes = 32 );

    virtual void read( const FileNode& );
    virtual void write( FileStorage& ) const;
    virtual int descriptorSize() const;
    virtual int descriptorType() const;
protected:
    ...
};
```

# 7.4 Common Interfaces of Descriptor Matchers

Matchers of keypoint descriptors in OpenCV have wrappers with a common interface that enables you to easily switch between different algorithms solving the same problem. This section is devoted to matching descriptors that are represented as vectors in a multidimensional space. All objects that implement vector descriptor matchers inherit the DescriptorMatcher interface.

## DMatch

**struct DMatch**

Class for matching keypoint descriptors: query descriptor index, train descriptor index, train image index, and distance between descriptors.

```
struct DMatch
{
    DMatch() : queryIdx(-1), trainIdx(-1), imgIdx(-1),
               distance(std::numeric_limits<float>::max()) {}
    DMatch( int _queryIdx, int _trainIdx, float _distance ) :
            queryIdx(_queryIdx), trainIdx(_trainIdx), imgIdx(-1),
            distance(_distance) {}
    DMatch( int _queryIdx, int _trainIdx, int _imgIdx, float _distance ) :
            queryIdx(_queryIdx), trainIdx(_trainIdx), imgIdx(_imgIdx),
            distance(_distance) {}

    int queryIdx; // query descriptor index
    int trainIdx; // train descriptor index
    int imgIdx;   // train image index

    float distance;

    // less is better
    bool operator<( const DMatch &m ) const;
};
```

## DescriptorMatcher

**class DescriptorMatcher** : **public Algorithm**

Abstract base class for matching keypoint descriptors. It has two groups of match methods: for matching descriptors of an image with another image or with an image set.

```
class DescriptorMatcher
{
public:
    virtual ~DescriptorMatcher();

    virtual void add( const vector<Mat>& descriptors );

    const vector<Mat>& getTrainDescriptors() const;
    virtual void clear();
    bool empty() const;
    virtual bool isMaskSupported() const = 0;

    virtual void train();

    /*
     * Group of methods to match descriptors from an image pair.
     */
    void match( const Mat& queryDescriptors, const Mat& trainDescriptors,
                vector<DMatch>& matches, const Mat& mask=Mat() ) const;
    void knnMatch( const Mat& queryDescriptors, const Mat& trainDescriptors,
                   vector<vector<DMatch> >& matches, int k,
                   const Mat& mask=Mat(), bool compactResult=false ) const;
    void radiusMatch( const Mat& queryDescriptors, const Mat& trainDescriptors,
                      vector<vector<DMatch> >& matches, float maxDistance,
                      const Mat& mask=Mat(), bool compactResult=false ) const;
    /*
     * Group of methods to match descriptors from one image to an image set.
     */
    void match( const Mat& queryDescriptors, vector<DMatch>& matches,
                const vector<Mat>& masks=vector<Mat>() );
    void knnMatch( const Mat& queryDescriptors, vector<vector<DMatch> >& matches,
                   int k, const vector<Mat>& masks=vector<Mat>(),
                   bool compactResult=false );
    void radiusMatch( const Mat& queryDescriptors, vector<vector<DMatch> >& matches,
                      float maxDistance, const vector<Mat>& masks=vector<Mat>(),
                      bool compactResult=false );

    virtual void read( const FileNode& );
    virtual void write( FileStorage& ) const;

    virtual Ptr<DescriptorMatcher> clone( bool emptyTrainData=false ) const = 0;

    static Ptr<DescriptorMatcher> create( const string& descriptorMatcherType );

protected:
    vector<Mat> trainDescCollection;
    ...
};
```

## DescriptorMatcher::add

Adds descriptors to train a descriptor collection. If the collection `trainDescCollectionis` is not empty, the new descriptors are added to existing train descriptors.

**C++:** void DescriptorMatcher::**add**(const vector<Mat>& **descriptors**)

>   **Parameters**
>
>>   **descriptors** – Descriptors to add. Each `descriptors[i]` is a set of descriptors from the same train image.

## DescriptorMatcher::getTrainDescriptors

Returns a constant link to the train descriptor collection `trainDescCollection` .

**C++:** const vector<Mat>& DescriptorMatcher::**getTrainDescriptors**() const

## DescriptorMatcher::clear

Clears the train descriptor collection.

**C++:** void DescriptorMatcher::**clear**()

## DescriptorMatcher::empty

Returns true if there are no train descriptors in the collection.

**C++:** bool DescriptorMatcher::**empty**() const

## DescriptorMatcher::isMaskSupported

Returns true if the descriptor matcher supports masking permissible matches.

**C++:** bool DescriptorMatcher::**isMaskSupported**()

## DescriptorMatcher::train

Trains a descriptor matcher

**C++:** void DescriptorMatcher::**train**()

Trains a descriptor matcher (for example, the flann index). In all methods to match, the method `train()` is run every time before matching. Some descriptor matchers (for example, `BruteForceMatcher`) have an empty implementation of this method. Other matchers really train their inner structures (for example, `FlannBasedMatcher` trains `flann::Index` ).

## DescriptorMatcher::match

Finds the best match for each descriptor from a query set.

**C++:** void DescriptorMatcher::**match**(const Mat& **queryDescriptors**, const Mat& **trainDescriptors**, vector<DMatch>& **matches**, const Mat& **mask**=Mat() ) const

---

**C++:** void DescriptorMatcher::**match**(const Mat& **queryDescriptors**, vector<DMatch>& **matches**, const vector<Mat>& **masks**=vector<Mat>() )

> Parameters

>> **queryDescriptors** – Query set of descriptors.

>> **trainDescriptors** – Train set of descriptors. This set is not added to the train descriptors collection stored in the class object.

>> **matches** – Matches. If a query descriptor is masked out in `mask` , no match is added for this descriptor. So, `matches` size may be smaller than the query descriptors count.

>> **mask** – Mask specifying permissible matches between an input query and train matrices of descriptors.

>> **masks** – Set of masks. Each `masks[i]` specifies permissible matches between the input query descriptors and stored train descriptors from the i-th image `trainDescCollection[i]`.

In the first variant of this method, the train descriptors are passed as an input argument. In the second variant of the method, train descriptors collection that was set by `DescriptorMatcher::add` is used. Optional mask (or masks) can be passed to specify which query and training descriptors can be matched. Namely, `queryDescriptors[i]` can be matched with `trainDescriptors[j]` only if `mask.at<uchar>(i,j)` is non-zero.

## DescriptorMatcher::knnMatch

Finds the k best matches for each descriptor from a query set.

**C++:** void DescriptorMatcher::**knnMatch**(const Mat& **queryDescriptors**, const Mat& **trainDescriptors**, vector<vector<DMatch>>& **matches**, int **k**, const Mat& **mask**=Mat(), bool **compactResult**=false ) const

**C++:** void DescriptorMatcher::**knnMatch**(const Mat& **queryDescriptors**, vector<vector<DMatch>>& **matches**, int **k**, const vector<Mat>& **masks**=vector<Mat>(), bool **compactResult**=false )

> Parameters

>> **queryDescriptors** – Query set of descriptors.

>> **trainDescriptors** – Train set of descriptors. This set is not added to the train descriptors collection stored in the class object.

>> **mask** – Mask specifying permissible matches between an input query and train matrices of descriptors.

>> **masks** – Set of masks. Each `masks[i]` specifies permissible matches between the input query descriptors and stored train descriptors from the i-th image `trainDescCollection[i]`.

>> **matches** – Matches. Each `matches[i]` is k or less matches for the same query descriptor.

>> **k** – Count of best matches found per each query descriptor or less if a query descriptor has less than k possible matches in total.

>> **compactResult** – Parameter used when the mask (or masks) is not empty. If `compactResult` is false, the `matches` vector has the same size as `queryDescriptors` rows. If `compactResult` is true, the `matches` vector does not contain matches for fully masked-out query descriptors.

These extended variants of `DescriptorMatcher::match()` methods find several best matches for each query descriptor. The matches are returned in the distance increasing order. See `DescriptorMatcher::match()` for the details about query and train descriptors.

## DescriptorMatcher::radiusMatch

For each query descriptor, finds the training descriptors not farther than the specified distance.

**C++:** void DescriptorMatcher::**radiusMatch**(const Mat& **queryDescriptors**, const Mat& **trainDescriptors**, vector<vector<DMatch>>& **matches**, float **maxDistance**, const Mat& **mask**=Mat(), bool **compactResult**=false ) const

**C++:** void DescriptorMatcher::**radiusMatch**(const Mat& **queryDescriptors**, vector<vector<DMatch>>& **matches**, float **maxDistance**, const vector<Mat>& **masks**=vector<Mat>(), bool **compactResult**=false )

> **Parameters**
>
>> **queryDescriptors** – Query set of descriptors.
>>
>> **trainDescriptors** – Train set of descriptors. This set is not added to the train descriptors collection stored in the class object.
>>
>> **mask** – Mask specifying permissible matches between an input query and train matrices of descriptors.
>>
>> **masks** – Set of masks. Each `masks[i]` specifies permissible matches between the input query descriptors and stored train descriptors from the i-th image `trainDescCollection[i]`.
>>
>> **matches** – Found matches.
>>
>> **compactResult** – Parameter used when the mask (or masks) is not empty. If `compactResult` is false, the `matches` vector has the same size as `queryDescriptors` rows. If `compactResult` is true, the `matches` vector does not contain matches for fully masked-out query descriptors.
>>
>> **maxDistance** – Threshold for the distance between matched descriptors.

For each query descriptor, the methods find such training descriptors that the distance between the query descriptor and the training descriptor is equal or smaller than `maxDistance`. Found matches are returned in the distance increasing order.

## DescriptorMatcher::clone

Clones the matcher.

**C++:** Ptr<DescriptorMatcher> DescriptorMatcher::**clone**(bool **emptyTrainData**=false )

> **Parameters**
>
>> **emptyTrainData** – If `emptyTrainData` is false, the method creates a deep copy of the object, that is, copies both parameters and train data. If `emptyTrainData` is true, the method creates an object copy with the current parameters but with empty train data.

## DescriptorMatcher::create

Creates a descriptor matcher of a given type with the default parameters (using default constructor).

**C++:** Ptr<DescriptorMatcher> DescriptorMatcher::**create**(const string& **descriptorMatcherType**)

> **Parameters**
>
>> **descriptorMatcherType** – Descriptor matcher type. Now the following matcher types are supported:
>>
>> – `BruteForce` (it uses `L2` )
>>
>> – `BruteForce-L1`
>>
>> – `BruteForce-Hamming`
>>
>> – `BruteForce-Hamming(2)`
>>
>> – `FlannBased`

## BFMatcher

**class BFMatcher** : **public DescriptorMatcher**

Brute-force descriptor matcher. For each descriptor in the first set, this matcher finds the closest descriptor in the second set by trying each one. This descriptor matcher supports masking permissible matches of descriptor sets.

## BFMatcher::BFMatcher

Brute-force matcher constructor.

**C++:** BFMatcher::**BFMatcher**(int **normType**, bool **crossCheck**=false )

> **Parameters**
>
>> **normType** – One of `NORM_L1`, `NORM_L2`, `NORM_HAMMING`, `NORM_HAMMING2`. L1 and L2 norms are preferable choices for SIFT and SURF descriptors, `NORM_HAMMING` should be used with ORB and BRIEF, `NORM_HAMMING2` should be used with ORB when `WTA_K==3` or 4 (see ORB::ORB constructor description).
>>
>> **crossCheck** – If it is false, this is will be default BFMatcher behaviour when it finds the k nearest neighbors for each query descriptor. If `crossCheck==true`, then the `knnMatch()` method with k=1 will only return pairs `(i,j)` such that for `i-th` query descriptor the `j-th` descriptor in the matcher's collection is the nearest and vice versa, i.e. the `BFMathcher` will only return consistent pairs. Such technique usually produces best results with minimal number of outliers when there are enough matches. This is alternative to the ratio test, used by D. Lowe in SIFT paper.

## FlannBasedMatcher

**class FlannBasedMatcher** : **public DescriptorMatcher**

Flann-based descriptor matcher. This matcher trains `flann::Index_` on a train descriptor collection and calls its nearest search methods to find the best matches. So, this matcher may be faster when matching a large train collection than the brute force matcher. `FlannBasedMatcher` does not support masking permissible matches of descriptor sets because `flann::Index` does not support this.

```
class FlannBasedMatcher : public DescriptorMatcher
{
public:
    FlannBasedMatcher(
      const Ptr<flann::IndexParams>& indexParams=new flann::KDTreeIndexParams(),
      const Ptr<flann::SearchParams>& searchParams=new flann::SearchParams() );

    virtual void add( const vector<Mat>& descriptors );
    virtual void clear();

    virtual void train();
    virtual bool isMaskSupported() const;

    virtual Ptr<DescriptorMatcher> clone( bool emptyTrainData=false ) const;
protected:
    ...
};
```

# 7.5 Common Interfaces of Generic Descriptor Matchers

Matchers of keypoint descriptors in OpenCV have wrappers with a common interface that enables you to easily switch between different algorithms solving the same problem. This section is devoted to matching descriptors that cannot be represented as vectors in a multidimensional space. GenericDescriptorMatcher is a more generic interface for descriptors. It does not make any assumptions about descriptor representation. Every descriptor with the DescriptorExtractor interface has a wrapper with the GenericDescriptorMatcher interface (see VectorDescriptorMatcher ). There are descriptors such as the One-way descriptor and Ferns that have the GenericDescriptorMatcher interface implemented but do not support DescriptorExtractor.

## GenericDescriptorMatcher

class **GenericDescriptorMatcher**

Abstract interface for extracting and matching a keypoint descriptor. There are also DescriptorExtractor and DescriptorMatcher for these purposes but their interfaces are intended for descriptors represented as vectors in a multidimensional space. GenericDescriptorMatcher is a more generic interface for descriptors. DescriptorMatcher and GenericDescriptorMatcher have two groups of match methods: for matching keypoints of an image with another image or with an image set.

```
class GenericDescriptorMatcher
{
public:
    GenericDescriptorMatcher();
    virtual ~GenericDescriptorMatcher();

    virtual void add( const vector<Mat>& images,
                      vector<vector<KeyPoint> >& keypoints );

    const vector<Mat>& getTrainImages() const;
    const vector<vector<KeyPoint> >& getTrainKeypoints() const;
    virtual void clear();

    virtual void train() = 0;

    virtual bool isMaskSupported() = 0;
```

```cpp
    void classify( const Mat& queryImage,
                   vector<KeyPoint>& queryKeypoints,
                   const Mat& trainImage,
                   vector<KeyPoint>& trainKeypoints ) const;
    void classify( const Mat& queryImage,
                   vector<KeyPoint>& queryKeypoints );

    /*
     * Group of methods to match keypoints from an image pair.
     */
    void match( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                const Mat& trainImage, vector<KeyPoint>& trainKeypoints,
                vector<DMatch>& matches, const Mat& mask=Mat() ) const;
    void knnMatch( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                   const Mat& trainImage, vector<KeyPoint>& trainKeypoints,
                   vector<vector<DMatch> >& matches, int k,
                   const Mat& mask=Mat(), bool compactResult=false ) const;
    void radiusMatch( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                      const Mat& trainImage, vector<KeyPoint>& trainKeypoints,
                      vector<vector<DMatch> >& matches, float maxDistance,
                      const Mat& mask=Mat(), bool compactResult=false ) const;
    /*
     * Group of methods to match keypoints from one image to an image set.
     */
    void match( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                vector<DMatch>& matches, const vector<Mat>& masks=vector<Mat>() );
    void knnMatch( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                   vector<vector<DMatch> >& matches, int k,
                   const vector<Mat>& masks=vector<Mat>(), bool compactResult=false );
    void radiusMatch( const Mat& queryImage, vector<KeyPoint>& queryKeypoints,
                      vector<vector<DMatch> >& matches, float maxDistance,
                      const vector<Mat>& masks=vector<Mat>(), bool compactResult=false );

    virtual void read( const FileNode& );
    virtual void write( FileStorage& ) const;

    virtual Ptr<GenericDescriptorMatcher> clone( bool emptyTrainData=false ) const = 0;

protected:
    ...
};
```

## GenericDescriptorMatcher::add

Adds images and their keypoints to the training collection stored in the class instance.

**C++:** void `GenericDescriptorMatcher::`**add**(const vector<Mat>& **images**, vector<vector<KeyPoint>>& **keypoints**)

>   **Parameters**

>>      **images** – Image collection.

>>      **keypoints** – Point collection. It is assumed that `keypoints[i]` are keypoints detected in the image `images[i]` .

## GenericDescriptorMatcher::getTrainImages

Returns a train image collection.

**C++:** const vector<Mat>& GenericDescriptorMatcher::**getTrainImages**() const

## GenericDescriptorMatcher::getTrainKeypoints

Returns a train keypoints collection.

**C++:** const vector<vector<KeyPoint>>& GenericDescriptorMatcher::**getTrainKeypoints**() const

## GenericDescriptorMatcher::clear

Clears a train collection (images and keypoints).

**C++:** void GenericDescriptorMatcher::**clear**()

## GenericDescriptorMatcher::train

Trains descriptor matcher

**C++:** void GenericDescriptorMatcher::**train**()

Prepares descriptor matcher, for example, creates a tree-based structure, to extract descriptors or to optimize descriptors matching.

## GenericDescriptorMatcher::isMaskSupported

Returns `true` if a generic descriptor matcher supports masking permissible matches.

**C++:** bool GenericDescriptorMatcher::**isMaskSupported**()

## GenericDescriptorMatcher::classify

Classifies keypoints from a query set.

**C++:** void GenericDescriptorMatcher::**classify**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, const Mat& **trainImage**, vector<KeyPoint>& **trainKeypoints**) const

**C++:** void GenericDescriptorMatcher::**classify**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**)

> **Parameters**
>> **queryImage** – Query image.
>>
>> **queryKeypoints** – Keypoints from a query image.
>>
>> **trainImage** – Train image.
>>
>> **trainKeypoints** – Keypoints from a train image.

The method classifies each keypoint from a query set. The first variant of the method takes a train image and its keypoints as an input argument. The second variant uses the internally stored training collection that can be built using the `GenericDescriptorMatcher::add` method.

The methods do the following:

1. Call the `GenericDescriptorMatcher::match` method to find correspondence between the query set and the training set.

2. Set the `class_id` field of each keypoint from the query set to `class_id` of the corresponding keypoint from the training set.

## GenericDescriptorMatcher::match

Finds the best match in the training set for each keypoint from the query set.

**C++:** void GenericDescriptorMatcher::**match**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, const Mat& **trainImage**, vector<KeyPoint>& **trainKeypoints**, vector<DMatch>& **matches**, const Mat& **mask**=Mat() ) const

**C++:** void GenericDescriptorMatcher::**match**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, vector<DMatch>& **matches**, const vector<Mat>& **masks**=vector<Mat>() )

> **Parameters**
>
> > **queryImage** – Query image.
> >
> > **queryKeypoints** – Keypoints detected in `queryImage` .
> >
> > **trainImage** – Train image. It is not added to a train image collection stored in the class object.
> >
> > **trainKeypoints** – Keypoints detected in `trainImage` . They are not added to a train points collection stored in the class object.
> >
> > **matches** – Matches. If a query descriptor (keypoint) is masked out in `mask` , match is added for this descriptor. So, `matches` size may be smaller than the query keypoints count.
> >
> > **mask** – Mask specifying permissible matches between an input query and train keypoints.
> >
> > **masks** – Set of masks. Each `masks[i]` specifies permissible matches between input query keypoints and stored train keypoints from the i-th image.

The methods find the best match for each query keypoint. In the first variant of the method, a train image and its keypoints are the input arguments. In the second variant, query keypoints are matched to the internally stored training collection that can be built using the `GenericDescriptorMatcher::add` method. Optional mask (or masks) can be passed to specify which query and training descriptors can be matched. Namely, `queryKeypoints[i]` can be matched with `trainKeypoints[j]` only if `mask.at<uchar>(i,j)` is non-zero.

## GenericDescriptorMatcher::knnMatch

Finds the `k` best matches for each query keypoint.

**C++:** void GenericDescriptorMatcher::**knnMatch**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, const Mat& **trainImage**, vector<KeyPoint>& **trainKeypoints**, vector<vector<DMatch>>& **matches**, int **k**, const Mat& **mask**=Mat(), bool **compactResult**=false ) const

**C++:** void GenericDescriptorMatcher::**knnMatch**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, vector<vector<DMatch>>& **matches**, int **k**, const vector<Mat>& **masks**=vector<Mat>(), bool **compactResult**=false )

The methods are extended variants of `GenericDescriptorMatch::match`. The parameters are similar, and the semantics is similar to `DescriptorMatcher::knnMatch`. But this class does not require explicitly computed keypoint descriptors.

## GenericDescriptorMatcher::radiusMatch

For each query keypoint, finds the training keypoints not farther than the specified distance.

**C++:** void GenericDescriptorMatcher::**radiusMatch**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, const Mat& **trainImage**, vector<KeyPoint>& **trainKeypoints**, vector<vector<DMatch>>& **matches**, float **maxDistance**, const Mat& **mask**=Mat(), bool **compactResult**=false ) const

**C++:** void GenericDescriptorMatcher::**radiusMatch**(const Mat& **queryImage**, vector<KeyPoint>& **queryKeypoints**, vector<vector<DMatch>>& **matches**, float **maxDistance**, const vector<Mat>& **masks**=vector<Mat>(), bool **compactResult**=false )

The methods are similar to `DescriptorMatcher::radius`. But this class does not require explicitly computed keypoint descriptors.

## GenericDescriptorMatcher::read

Reads a matcher object from a file node.

**C++:** void GenericDescriptorMatcher::**read**(const FileNode& **fn**)

## GenericDescriptorMatcher::write

Writes a match object to a file storage.

**C++:** void GenericDescriptorMatcher::**write**(FileStorage& **fs**) const

## GenericDescriptorMatcher::clone

Clones the matcher.

**C++:** Ptr<GenericDescriptorMatcher> GenericDescriptorMatcher::**clone**(bool **emptyTrainData**=false ) const

   **Parameters**

> **emptyTrainData** – If `emptyTrainData` is false, the method creates a deep copy of the
> object, that is, copies both parameters and train data. If `emptyTrainData` is true, the method
> creates an object copy with the current parameters but with empty train data.

## VectorDescriptorMatcher

class `VectorDescriptorMatcher` : public `GenericDescriptorMatcher`

Class used for matching descriptors that can be described as vectors in a finite-dimensional space.

```
class CV_EXPORTS VectorDescriptorMatcher : public GenericDescriptorMatcher
{
public:
    VectorDescriptorMatcher( const Ptr<DescriptorExtractor>& extractor, const Ptr<DescriptorMatcher>& matcher );
    virtual ~VectorDescriptorMatcher();

    virtual void add( const vector<Mat>& imgCollection,
                      vector<vector<KeyPoint> >& pointCollection );
    virtual void clear();
    virtual void train();
    virtual bool isMaskSupported();

    virtual void read( const FileNode& fn );
    virtual void write( FileStorage& fs ) const;

    virtual Ptr<GenericDescriptorMatcher> clone( bool emptyTrainData=false ) const;

protected:
    ...
};
```

Example:

```
VectorDescriptorMatcher matcher( new SurfDescriptorExtractor,
                                 new BruteForceMatcher<L2<float> > );
```

# 7.6 Drawing Function of Keypoints and Matches

## drawMatches

Draws the found matches of keypoints from two images.

**C++:** void **drawMatches**(const Mat& **img1**, const vector<KeyPoint>& **keypoints1**, const Mat& **img2**, const vector<KeyPoint>& **keypoints2**, const vector<DMatch>& **matches1to2**, Mat& **outImg**, const Scalar& **matchColor**=Scalar::all(-1), const Scalar& **singlePointColor**=Scalar::all(-1), const vector<char>& **matchesMask**=vector<char>(), int **flags**=DrawMatchesFlags::DEFAULT )

**C++:** void **drawMatches**(const Mat& **img1**, const vector<KeyPoint>& **keypoints1**, const Mat& **img2**, const vector<KeyPoint>& **keypoints2**, const vector<vector<DMatch>>& **matches1to2**, Mat& **outImg**, const Scalar& **matchColor**=Scalar::all(-1), const Scalar& **singlePointColor**=Scalar::all(-1), const vector<vector<char>>& **matchesMask**=vector<vector<char> >(), int **flags**=DrawMatchesFlags::DEFAULT )

> **Parameters**

>> **img1** – First source image.

>> **keypoints1** – Keypoints from the first source image.

>> **img2** – Second source image.

>> **keypoints2** – Keypoints from the second source image.

>> **matches1to2** – Matches from the first image to the second one, which means that `keypoints1[i]` has a corresponding point in `keypoints2[matches[i]]` .

>> **outImg** – Output image. Its content depends on the `flags` value defining what is drawn in the output image. See possible `flags` bit values below.

>> **matchColor** – Color of matches (lines and connected keypoints). If `matchColor==Scalar::all(-1)` , the color is generated randomly.

>> **singlePointColor** – Color of single keypoints (circles), which means that keypoints do not have the matches. If `singlePointColor==Scalar::all(-1)` , the color is generated randomly.

>> **matchesMask** – Mask determining which matches are drawn. If the mask is empty, all matches are drawn.

>> **flags** – Flags setting drawing features. Possible `flags` bit values are defined by `DrawMatchesFlags`.

This function draws matches of keypoints from two images in the output image. Match is a line connecting two keypoints (circles). The structure `DrawMatchesFlags` is defined as follows:

```
struct DrawMatchesFlags
{
    enum
    {
        DEFAULT = 0, // Output image matrix will be created (Mat::create),
                     // i.e. existing memory of output image may be reused.
                     // Two source images, matches, and single keypoints
                     // will be drawn.
                     // For each keypoint, only the center point will be
                     // drawn (without a circle around the keypoint with the
                     // keypoint size and orientation).
        DRAW_OVER_OUTIMG = 1, // Output image matrix will not be
                     // created (using Mat::create). Matches will be drawn
                     // on existing content of output image.
        NOT_DRAW_SINGLE_POINTS = 2, // Single keypoints will not be drawn.
        DRAW_RICH_KEYPOINTS = 4 // For each keypoint, the circle around
                     // keypoint with keypoint size and orientation will
                     // be drawn.
    };
};
```

## drawKeypoints

Draws keypoints.

**C++:** void **drawKeypoints**(const Mat& **image**, const vector<KeyPoint>& **keypoints**, Mat& **outImage**, const Scalar& **color**=Scalar::all(-1), int **flags**=DrawMatchesFlags::DEFAULT )

>> **Parameters**

>>> **image** – Source image.

**keypoints** – Keypoints from the source image.

**outImage** – Output image. Its content depends on the `flags` value defining what is drawn in the output image. See possible `flags` bit values below.

**color** – Color of keypoints.

**flags** – Flags setting drawing features. Possible `flags` bit values are defined by DrawMatchesFlags. See details above in `drawMatches()` .

# 7.7 Object Categorization

This section describes approaches based on local 2D features and used to categorize objects.

## BOWTrainer

**class** `BOWTrainer`

Abstract base class for training the *bag of visual words* vocabulary from a set of descriptors. For details, see, for example, *Visual Categorization with Bags of Keypoints* by Gabriella Csurka, Christopher R. Dance, Lixin Fan, Jutta Willamowski, Cedric Bray, 2004.

```cpp
class BOWTrainer
{
public:
    BOWTrainer(){}
    virtual ~BOWTrainer(){}

    void add( const Mat& descriptors );
    const vector<Mat>& getDescriptors() const;
    int descripotorsCount() const;

    virtual void clear();

    virtual Mat cluster() const = 0;
    virtual Mat cluster( const Mat& descriptors ) const = 0;

protected:
    ...
};
```

## BOWTrainer::add

Adds descriptors to a training set.

**C++:** void BOWTrainer::**add**(const Mat& **descriptors**)

        **Parameters**

                **descriptors** – Descriptors to add to a training set. Each row of the `descriptors` matrix is a descriptor.

The training set is clustered using `clustermethod` to construct the vocabulary.

## BOWTrainer::getDescriptors

Returns a training set of descriptors.

**C++:** const vector<Mat>& BOWTrainer::**getDescriptors**() const

## BOWTrainer::descripotorsCount

Returns the count of all descriptors stored in the training set.

**C++:** int BOWTrainer::**descripotorsCount**() const

## BOWTrainer::cluster

Clusters train descriptors.

**C++:** Mat BOWTrainer::**cluster**() const

**C++:** Mat BOWTrainer::**cluster**(const Mat& **descriptors**) const

> **Parameters**
>
> > **descriptors** – Descriptors to cluster. Each row of the `descriptors` matrix is a descriptor. Descriptors are not added to the inner train descriptor set.

The vocabulary consists of cluster centers. So, this method returns the vocabulary. In the first variant of the method, train descriptors stored in the object are clustered. In the second variant, input descriptors are clustered.

## BOWKMeansTrainer

**class BOWKMeansTrainer** : **public BOWTrainer**

`kmeans()` -based class to train visual vocabulary using the *bag of visual words* approach.

```
class BOWKMeansTrainer : public BOWTrainer
{
public:
    BOWKMeansTrainer( int clusterCount, const TermCriteria& termcrit=TermCriteria(),
                      int attempts=3, int flags=KMEANS_PP_CENTERS );
    virtual ~BOWKMeansTrainer(){}

    // Returns trained vocabulary (i.e. cluster centers).
    virtual Mat cluster() const;
    virtual Mat cluster( const Mat& descriptors ) const;

protected:
    ...
};
```

## BOWKMeansTrainer::BOWKMeansTrainer

The constructor.

**C++:** BOWKMeansTrainer::**BOWKMeansTrainer**(int **clusterCount**, const TermCriteria& **term-crit**=TermCriteria(), int **attempts**=3, int **flags**=KMEANS_PP_CENTERS )

>   See kmeans() function parameters.

# BOWImgDescriptorExtractor

**class BOWImgDescriptorExtractor**

Class to compute an image descriptor using the *bag of visual words*. Such a computation consists of the following steps:

1. Compute descriptors for a given image and its keypoints set.

2. Find the nearest visual words from the vocabulary for each keypoint descriptor.

3. Compute the bag-of-words image descriptor as is a normalized histogram of vocabulary words encountered in the image. The i-th bin of the histogram is a frequency of i-th word of the vocabulary in the given image.

The class declaration is the following:

```cpp
class BOWImgDescriptorExtractor
{
public:
    BOWImgDescriptorExtractor( const Ptr<DescriptorExtractor>& dextractor,
                               const Ptr<DescriptorMatcher>& dmatcher );
    virtual ~BOWImgDescriptorExtractor(){}

    void setVocabulary( const Mat& vocabulary );
    const Mat& getVocabulary() const;
    void compute( const Mat& image, vector<KeyPoint>& keypoints,
                  Mat& imgDescriptor,
                  vector<vector<int> >* pointIdxsOfClusters=0,
                  Mat* descriptors=0 );
    int descriptorSize() const;
    int descriptorType() const;

protected:
    ...
};
```

# BOWImgDescriptorExtractor::BOWImgDescriptorExtractor

The constructor.

**C++:** BOWImgDescriptorExtractor::**BOWImgDescriptorExtractor**(const Ptr<DescriptorExtractor>& **dextractor**, const Ptr<DescriptorMatcher>& **dmatcher**)

>   **Parameters**

>> **dextractor** – Descriptor extractor that is used to compute descriptors for an input image and its keypoints.

>> **dmatcher** – Descriptor matcher that is used to find the nearest word of the trained vocabulary for each keypoint descriptor of the image.

## BOWImgDescriptorExtractor::setVocabulary

Sets a visual vocabulary.

**C++:** void BOWImgDescriptorExtractor::**setVocabulary**(const Mat& **vocabulary**)

> **Parameters**
>
> > **vocabulary** – Vocabulary (can be trained using the inheritor of `BOWTrainer` ). Each row of the vocabulary is a visual word (cluster center).

## BOWImgDescriptorExtractor::getVocabulary

Returns the set vocabulary.

**C++:** const Mat& BOWImgDescriptorExtractor::**getVocabulary**() const

## BOWImgDescriptorExtractor::compute

Computes an image descriptor using the set visual vocabulary.

**C++:** void BOWImgDescriptorExtractor::**compute**(const Mat& **image**, vector<KeyPoint>& **keypoints**, Mat& **imgDescriptor**, vector<vector<int>>* **pointIdxsOfClusters**=0, Mat* **descriptors**=0 )

> **Parameters**
>
> > **image** – Image, for which the descriptor is computed.
> >
> > **keypoints** – Keypoints detected in the input image.
> >
> > **imgDescriptor** – Computed output image descriptor.
> >
> > **pointIdxsOfClusters** – Indices of keypoints that belong to the cluster. This means that `pointIdxsOfClusters[i]` are keypoint indices that belong to the `i` -th cluster (word of vocabulary) returned if it is non-zero.
> >
> > **descriptors** – Descriptors of the image keypoints that are returned if they are non-zero.

## BOWImgDescriptorExtractor::descriptorSize

Returns an image descriptor size if the vocabulary is set. Otherwise, it returns 0.

**C++:** int BOWImgDescriptorExtractor::**descriptorSize**() const

## BOWImgDescriptorExtractor::descriptorType

Returns an image descriptor type.

**C++:** int BOWImgDescriptorExtractor::**descriptorType**() const

# OBJDETECT. OBJECT DETECTION

## 8.1 Cascade Classification

### Haar Feature-based Cascade Classifier for Object Detection
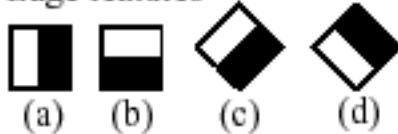
The object detector described below has been initially proposed by Paul Viola [Viola01] and improved by Rainer Lienhart [Lienhart02].

First, a classifier (namely a *cascade of boosted classifiers working with haar-like features*) is trained with a few hundred sample views of a particular object (i.e., a face or a car), called positive examples, that are scaled to the same size (say, 20x20), and negative examples - arbitrary images of the same size.
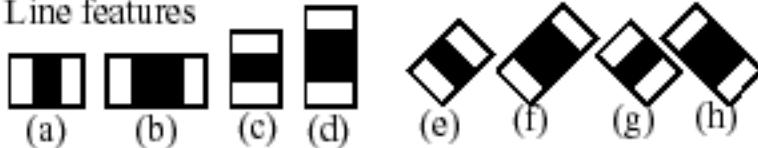
After a classifier is trained, it can be applied to a region of interest (of the same size as used during the training) in an input image. The classifier outputs a "1" if the region is likely to show the object (i.e., face/car), and "0" otherwise. To search for the object in the whole image one can move the search window across the image and check every location using the classifier. The classifier is designed so that it can be easily "resized" in order to be able to find the objects of interest at different sizes, which is more efficient than resizing the image itself. So, to find an object of an unknown size in the image the scan procedure should be done several times at different scales.

The word "cascade" in the classifier name means that the resultant classifier consists of several simpler classifiers (*stages*) that are applied subsequently to a region of interest until at some stage the candidate is rejected or all the stages are passed. The word "boosted" means that the classifiers at every stage of the cascade are complex themselves and they are built out of basic classifiers using one of four different `boosting` techniques (weighted voting). Currently Discrete Adaboost, Real Adaboost, Gentle Adaboost and Logitboost are supported. The basic classifiers are decision-tree classifiers with at least 2 leaves. Haar-like features are the input to the basic classifiers, and are calculated as described below. The current algorithm uses the following Haar-like features:
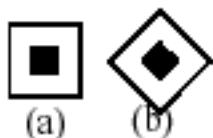
The feature used in a particular classifier is specified by its shape (1a, 2b etc.), position within the region of interest and the scale (this scale is not the same as the scale used at the detection stage, though these two scales are multiplied). For example, in the case of the third line feature (2c) the response is calculated as the difference between the sum of image pixels under the rectangle covering the whole feature (including the two white stripes and the black stripe in the middle) and the sum of the image pixels under the black stripe multiplied by 3 in order to compensate for the differences in the size of areas. The sums of pixel values over a rectangular regions are calculated rapidly using integral images (see below and the `integral()` description).

To see the object detector at work, have a look at the facedetect demo: http://code.opencv.org/projects/opencv/repository/revisions/master/entry/samples/cpp/facedetect.cpp

The following reference is for the detection part only. There is a separate application called `opencv_traincascade` that can train a cascade of boosted classifiers from a set of samples.

**Note:** In the new C++ interface it is also possible to use LBP (local binary pattern) features in addition to Haar-like features.

## FeatureEvaluator

**class FeatureEvaluator**

Base class for computing feature values in cascade classifiers.

```cpp
class CV_EXPORTS FeatureEvaluator
{
public:
    enum { HAAR = 0, LBP = 1 }; // supported feature types
    virtual ~FeatureEvaluator(); // destructor
    virtual bool read(const FileNode& node);
    virtual Ptr<FeatureEvaluator> clone() const;
    virtual int getFeatureType() const;

    virtual bool setImage(const Mat& img, Size origWinSize);
    virtual bool setWindow(Point p);

    virtual double calcOrd(int featureIdx) const;
```

```
    virtual int calcCat(int featureIdx) const;

    static Ptr<FeatureEvaluator> create(int type);
};
```

## FeatureEvaluator::read

Reads parameters of features from the `FileStorage` node.

**C++:** bool FeatureEvaluator::**read**(const FileNode& **node**)

>   **Parameters**

>>   **node** – File node from which the feature parameters are read.

## FeatureEvaluator::clone

Returns a full copy of the feature evaluator.

**C++:** Ptr<FeatureEvaluator> FeatureEvaluator::**clone**() const

## FeatureEvaluator::getFeatureType

Returns the feature type (HAAR or LBP for now).

**C++:** int FeatureEvaluator::**getFeatureType**() const

## FeatureEvaluator::setImage

Assigns an image to feature evaluator.

**C++:** bool FeatureEvaluator::**setImage**(const Mat& **img**, Size **origWinSize**)

>   **Parameters**

>>   **img** – Matrix of the type `CV_8UC1` containing an image where the features are computed.

>>   **origWinSize** – Size of training images.

The method assigns an image, where the features will be computed, to the feature evaluator.

## FeatureEvaluator::setWindow

Assigns a window in the current image where the features will be computed.

**C++:** bool FeatureEvaluator::**setWindow**(Point **p**)

>   **Parameters**

>>   **p** – Upper left point of the window where the features are computed. Size of the window is equal to the size of training images.

## FeatureEvaluator::calcOrd

Computes the value of an ordered (numerical) feature.

**C++:** double FeatureEvaluator::**calcOrd**(int **featureIdx**) const

> **Parameters**
>
> > **featureIdx** – Index of the feature whose value is computed.

The function returns the computed value of an ordered feature.

## FeatureEvaluator::calcCat

Computes the value of a categorical feature.

**C++:** int FeatureEvaluator::**calcCat**(int **featureIdx**) const

> **Parameters**
>
> > **featureIdx** – Index of the feature whose value is computed.

The function returns the computed label of a categorical feature, which is the value from [0,... (number of categories - 1)].

## FeatureEvaluator::create

Constructs the feature evaluator.

**C++:** Ptr<FeatureEvaluator> FeatureEvaluator::**create**(int **type**)

> **Parameters**
>
> > **type** – Type of features evaluated by cascade (HAAR or LBP for now).

## CascadeClassifier

class **CascadeClassifier**

Cascade classifier class for object detection.

## CascadeClassifier::CascadeClassifier

Loads a classifier from a file.

**C++:** CascadeClassifier::**CascadeClassifier**(const string& **filename**)

**Python:** cv2.**CascadeClassifier**($\left[$filename$\right]$) → <CascadeClassifier object>

> **Parameters**
>
> > **filename** – Name of the file from which the classifier is loaded.

## CascadeClassifier::empty

Checks whether the classifier has been loaded.

**C++:** bool CascadeClassifier::**empty**() const

**Python:** cv2.CascadeClassifier.**empty**() → retval

## CascadeClassifier::load

Loads a classifier from a file.

**C++:** bool CascadeClassifier::**load**(const string& **filename**)

**Python:** cv2.CascadeClassifier.**load**(filename) → retval

> **Parameters**
>
>> **filename** – Name of the file from which the classifier is loaded. The file may contain an old HAAR classifier trained by the haartraining application or a new cascade classifier trained by the traincascade application.

## CascadeClassifier::read

Reads a classifier from a FileStorage node.

**C++:** bool CascadeClassifier::**read**(const FileNode& **node**)

---

**Note:** The file may contain a new cascade classifier (trained traincascade application) only.

---

## CascadeClassifier::detectMultiScale

Detects objects of different sizes in the input image. The detected objects are returned as a list of rectangles.

**C++:** void CascadeClassifier::**detectMultiScale**(const Mat& **image**, vector<Rect>& **objects**, double **scaleFactor**=1.1, int **minNeighbors**=3, int **flags**=0, Size **minSize**=Size(), Size **maxSize**=Size())

**Python:** cv2.CascadeClassifier.**detectMultiScale**(image[, scaleFactor[, minNeighbors[, flags[, minSize[, maxSize ] ] ] ] ]) → objects

**Python:** cv2.CascadeClassifier.**detectMultiScale**(image, rejectLevels, levelWeights[, scaleFactor[, minNeighbors[, flags[, minSize[, maxSize[, outputRejectLevels ] ] ] ] ] ]) → objects

**C:** CvSeq* **cvHaarDetectObjects**(const CvArr* **image**, CvHaarClassifierCascade* **cascade**, CvMemStorage* **storage**, double **scale_factor**=1.1, int **min_neighbors**=3, int **flags**=0, CvSize **min_size**=cvSize(0,0), CvSize **max_size**=cvSize(0,0) )

**Python:** cv.**HaarDetectObjects**(image, cascade, storage, scale_factor=1.1, min_neighbors=3, flags=0, min_size=(0, 0)) → detectedObjects

> **Parameters**
>
>> **cascade** – Haar classifier cascade (OpenCV 1.x API only). It can be loaded from XML or YAML file using `Load()`. When the cascade is not needed anymore, release it using `cvReleaseHaarClassifierCascade(&cascade)`.

---

image – Matrix of the type CV_8U containing an image where objects are detected.

objects – Vector of rectangles where each rectangle contains the detected object.

scaleFactor – Parameter specifying how much the image size is reduced at each image scale.

minNeighbors – Parameter specifying how many neighbors each candidate rectangle should have to retain it.

flags – Parameter with the same meaning for an old cascade as in the function cvHaarDetectObjects. It is not used for a new cascade.

minSize – Minimum possible object size. Objects smaller than that are ignored.

maxSize – Maximum possible object size. Objects larger than that are ignored.

The function is parallelized with the TBB library.

## CascadeClassifier::setImage

Sets an image for detection.

**C++:** bool CascadeClassifier::**setImage**(Ptr<FeatureEvaluator>& **feval**, const Mat& **image**)

**C:** void **cvSetImagesForHaarClassifierCascade**(CvHaarClassifierCascade* **cascade**, const CvArr* **sum**, const CvArr* **sqsum**, const CvArr* **tilted_sum**, double **scale**)

Parameters

cascade – Haar classifier cascade (OpenCV 1.x API only). See CascadeClassifier::detectMultiScale() for more information.

feval – Pointer to the feature evaluator used for computing features.

image – Matrix of the type CV_8UC1 containing an image where the features are computed.

The function is automatically called by CascadeClassifier::detectMultiScale() at every image scale. But if you want to test various locations manually using CascadeClassifier::runAt(), you need to call the function before, so that the integral images are computed.

---

**Note:** in the old API you need to supply integral images (that can be obtained using Integral()) instead of the original image.

---

## CascadeClassifier::runAt

Runs the detector at the specified point.

**C++:** int CascadeClassifier::**runAt**(Ptr<FeatureEvaluator>& **feval**, Point **pt**, double& **weight**)

**C:** int **cvRunHaarClassifierCascade**(const CvHaarClassifierCascade* **cascade**, CvPoint **pt**, int **start_stage**=0 )

Parameters

cascade – Haar classifier cascade (OpenCV 1.x API only). See CascadeClassifier::detectMultiScale() for more information.

feval – Feature evaluator used for computing features.

> **pt** – Upper left point of the window where the features are computed. Size of the window is equal to the size of training images.

The function returns 1 if the cascade classifier detects an object in the given location. Otherwise, it returns negated index of the stage at which the candidate has been rejected.

Use `CascadeClassifier::setImage()` to set the image for the detector to work with.

## groupRectangles

Groups the object candidate rectangles.

**C++:** void **groupRectangles**(vector<Rect>& **rectList**, int **groupThreshold**, double **eps**=0.2)

**C++:** void **groupRectangles**(vector<Rect>& **rectList**, vector<int>& **weights**, int **groupThreshold**, double **eps**=0.2)

**Python:** cv2.**groupRectangles**(rectList, groupThreshold[, eps]) → rectList, weights

> **Parameters**

>> **rectList** – Input/output vector of rectangles. Output vector includes retained and grouped rectangles. (The Python list is not modified in place.)

>> **groupThreshold** – Minimum possible number of rectangles minus 1. The threshold is used in a group of rectangles to retain it.

>> **eps** – Relative difference between sides of the rectangles to merge them into a group.

The function is a wrapper for the generic function `partition()`. It clusters all the input rectangles using the rectangle equivalence criteria that combines rectangles with similar sizes and similar locations. The similarity is defined by `eps`. When `eps=0`, no clustering is done at all. If $eps \rightarrow +\inf$, all the rectangles are put in one cluster. Then, the small clusters containing less than or equal to `groupThreshold` rectangles are rejected. In each other cluster, the average rectangle is computed and put into the output rectangle list.

# 8.2 Latent SVM

## Discriminatively Trained Part Based Models for Object Detection

The object detector described below has been initially proposed by P.F. Felzenszwalb in [Felzenszwalb2010]. It is based on a Dalal-Triggs detector that uses a single filter on histogram of oriented gradients (HOG) features to represent an object category. This detector uses a sliding window approach, where a filter is applied at all positions and scales of an image. The first innovation is enriching the Dalal-Triggs model using a star-structured part-based model defined by a "root" filter (analogous to the Dalal-Triggs filter) plus a set of parts filters and associated deformation models. The score of one of star models at a particular position and scale within an image is the score of the root filter at the given location plus the sum over parts of the maximum, over placements of that part, of the part filter score on its location minus a deformation cost easuring the deviation of the part from its ideal location relative to the root. Both root and part filter scores are defined by the dot product between a filter (a set of weights) and a subwindow of a feature pyramid computed from the input image. Another improvement is a representation of the class of models by a mixture of star models. The score of a mixture model at a particular position and scale is the maximum over components, of the score of that component model at the given location.

In OpenCV there are C implementation of Latent SVM and C++ wrapper of it. C version is the structure `CvObjectDetection` and a set of functions working with this structure (see `cvLoadLatentSvmDetector()`, `cvReleaseLatentSvmDetector()`, `cvLatentSvmDetectObjects()`). C++ version is the class `LatentSvmDetector` and has slightly different functionality in contrast with C version - it supports loading and detection of several models.

There are two examples of Latent SVM usage: `samples/c/latentsvmdetect.cpp` and `samples/cpp/latentsvm_multidetect.cpp`.

## CvLSVMFilterPosition

**struct `CvLSVMFilterPosition`**
    Structure describes the position of the filter in the feature pyramid.

    unsigned int **`l`**
        level in the feature pyramid

    unsigned int **`x`**
        x-coordinate in level l

    unsigned int **`y`**
        y-coordinate in level l

## CvLSVMFilterObject

**struct `CvLSVMFilterObject`**
    Description of the filter, which corresponds to the part of the object.

    CvLSVMFilterPosition **`V`**
        ideal (penalty = 0) position of the partial filter from the root filter position ($V_i$ in the paper)

    float **`fineFunction[4]`**
        vector describes penalty function ($d_i$ in the paper) pf[0] * x + pf[1] * y + pf[2] * x^2 + pf[3] * y^2

    int **`sizeX`**

    int **`sizeY`**
        Rectangular map (sizeX x sizeY), every cell stores feature vector (dimension = p)

    int **`numFeatures`**
        number of features

    float* **`H`**
        matrix of feature vectors to set and get feature vectors (i,j) used formula H[(j * sizeX + i) * p + k], where k - component of feature vector in cell (i, j)

## CvLatentSvmDetector

**struct `CvLatentSvmDetector`**
    Structure contains internal representation of trained Latent SVM detector.

    int **`num_filters`**
        total number of filters (root plus part) in model

    int **`num_components`**
        number of components in model

    int* **`num_part_filters`**
        array containing number of part filters for each component

    CvLSVMFilterObject** **`filters`**
        root and part filters for all model components

float* **b**
    biases for all model components

float **score_threshold**
    confidence level threshold

## CvObjectDetection

**struct CvObjectDetection**
    Structure contains the bounding box and confidence level for detected object.

    CvRect **rect**
        bounding box for a detected object

    float **score**
        confidence level

## cvLoadLatentSvmDetector

Loads trained detector from a file.

**C++:** CvLatentSvmDetector* **cvLoadLatentSvmDetector**(const char* **filename**)

> **Parameters**

> > **filename** – Name of the file containing the description of a trained detector

## cvReleaseLatentSvmDetector

Release memory allocated for CvLatentSvmDetector structure.

**C++:** void **cvReleaseLatentSvmDetector**(CvLatentSvmDetector** **detector**)

> **Parameters**

> > **detector** – CvLatentSvmDetector structure to be released

## cvLatentSvmDetectObjects

Find rectangular regions in the given image that are likely to contain objects and corresponding confidence levels.

**C++:** CvSeq* **cvLatentSvmDetectObjects**(IplImage* **image**, CvLatentSvmDetector* **detector**, CvMem-Storage* **storage**, float **overlap_threshold**=0.5f, int **numThreads**=-1 )

> **Parameters**

> > **image** – image

> > **detector** – LatentSVM detector in internal representation

> > **storage** – Memory storage to store the resultant sequence of the object candidate rectangles

> > **overlap_threshold** – Threshold for the non-maximum suppression algorithm

> > **numThreads** – Number of threads used in parallel version of the algorithm

## LatentSvmDetector

**class LatentSvmDetector**

This is a C++ wrapping class of Latent SVM. It contains internal representation of several trained Latent SVM detectors (models) and a set of methods to load the detectors and detect objects using them.

## LatentSvmDetector::ObjectDetection

**struct LatentSvmDetector::ObjectDetection**
> Structure contains the detection information.

> Rect **rect**
>> bounding box for a detected object

> float **score**
>> confidence level

> int **classID**
>> class (model or detector) ID that detect an object

## LatentSvmDetector::LatentSvmDetector

Two types of constructors.

**C++:** LatentSvmDetector::**LatentSvmDetector**()

**C++:** LatentSvmDetector::**LatentSvmDetector**(const vector<string>& **filenames**, const vector<string>& **classNames**=vector<string>())

> **Parameters**

>> **filenames** – A set of filenames storing the trained detectors (models). Each file contains one model. See examples of such files here /opencv_extra/testdata/cv/latentsvmdetector/models_VOC2007/.

>> **classNames** – A set of trained models names. If it's empty then the name of each model will be constructed from the name of file containing the model. E.g. the model stored in "/home/user/cat.xml" will get the name "cat".

## LatentSvmDetector::~LatentSvmDetector

Destructor.

**C++:** LatentSvmDetector::**~LatentSvmDetector**()

## LatentSvmDetector::~clear

Clear all trained models and their names stored in an class object.

**C++:** void LatentSvmDetector::**clear**()

## LatentSvmDetector::load

Load the trained models from given `.xml` files and return `true` if at least one model was loaded.

**C++:** bool LatentSvmDetector::**load**(const vector<string>& **filenames**, const vector<string>& **class-Names**=vector<string>() )

> **Parameters**
>
> > **filenames** – A set of filenames storing the trained detectors (models). Each file contains one model. See examples of such files here /opencv_extra/testdata/cv/latentsvmdetector/models_VOC2007/.
> >
> > **classNames** – A set of trained models names. If it's empty then the name of each model will be constructed from the name of file containing the model. E.g. the model stored in "/home/user/cat.xml" will get the name "cat".

## LatentSvmDetector::detect

Find rectangular regions in the given image that are likely to contain objects of loaded classes (models) and corresponding confidence levels.

**C++:** void LatentSvmDetector::**detect**(const Mat& **image**, vector<ObjectDetection>& **objectDetections**, float **overlapThreshold**=0.5f, int **numThreads**=-1 )

> **Parameters**
>
> > **image** – An image.
> >
> > **objectDetections** – The detections: rectangulars, scores and class IDs.
> >
> > **overlapThreshold** – Threshold for the non-maximum suppression algorithm.
> >
> > **numThreads** – Number of threads used in parallel version of the algorithm.

## LatentSvmDetector::getClassNames

Return the class (model) names that were passed in constructor or method `load` or extracted from models filenames in those methods.

**C++:** const vector<string>& LatentSvmDetector::**getClassNames**() const

## LatentSvmDetector::getClassCount

Return a count of loaded models (classes).

**C++:** size_t LatentSvmDetector::**getClassCount**() const

# ML. MACHINE LEARNING

The Machine Learning Library (MLL) is a set of classes and functions for statistical classification, regression, and clustering of data.

Most of the classification and regression algorithms are implemented as C++ classes. As the algorithms have different sets of features (like an ability to handle missing measurements or categorical input variables), there is a little common ground between the classes. This common ground is defined by the class *CvStatModel* that all the other ML classes are derived from.

## 9.1 Statistical Models

### CvStatModel

**class CvStatModel**

Base class for statistical models in ML.

```
class CvStatModel
{
public:
    /* CvStatModel(); */
    /* CvStatModel( const Mat& train_data ... ); */

    virtual ~CvStatModel();

    virtual void clear()=0;

    /* virtual bool train( const Mat& train_data, [int tflag,] ..., const
        Mat& responses, ...,
     [const Mat& var_idx,] ..., [const Mat& sample_idx,] ...
     [const Mat& var_type,] ..., [const Mat& missing_mask,]
        <misc_training_alg_params> ... )=0;
      */

    /* virtual float predict( const Mat& sample ... ) const=0; */

    virtual void save( const char* filename, const char* name=0 )=0;
    virtual void load( const char* filename, const char* name=0 )=0;

    virtual void write( CvFileStorage* storage, const char* name )=0;
    virtual void read( CvFileStorage* storage, CvFileNode* node )=0;
};
```

In this declaration, some methods are commented off. These are methods for which there is no unified API (with the exception of the default constructor). However, there are many similarities in the syntax and semantics that are briefly described below in this section, as if they are part of the base class.

## CvStatModel::CvStatModel

The default constructor.

**C++:** `CvStatModel::`**`CvStatModel`**`()`

Each statistical model class in ML has a default constructor without parameters. This constructor is useful for a two-stage model construction, when the default constructor is followed by `CvStatModel::train()` or `CvStatModel::load()`.

## CvStatModel::CvStatModel(...)

The training constructor.

**C++:** `CvStatModel::`**`CvStatModel`**`()`

Most ML classes provide a single-step constructor and train constructors. This constructor is equivalent to the default constructor, followed by the `CvStatModel::train()` method with the parameters that are passed to the constructor.

## CvStatModel::~CvStatModel

The virtual destructor.

**C++:** `CvStatModel::`**`~CvStatModel`**`()`

The destructor of the base class is declared as virtual. So, it is safe to write the following code:

```
CvStatModel* model;
if( use_svm )
    model = new CvSVM(... /* SVM params */);
else
    model = new CvDTree(... /* Decision tree params */);
...
delete model;
```

Normally, the destructor of each derived class does nothing. But in this instance, it calls the overridden method `CvStatModel::clear()` that deallocates all the memory.

## CvStatModel::clear

Deallocates memory and resets the model state.

**C++:** `void CvStatModel::`**`clear`**`()`

The method `clear` does the same job as the destructor: it deallocates all the memory occupied by the class members. But the object itself is not destructed and can be reused further. This method is called from the destructor, from the `CvStatModel::train()` methods of the derived classes, from the methods `CvStatModel::load()`, `CvStatModel::read()`, or even explicitly by the user.

## CvStatModel::save

Saves the model to a file.

**C++:** void CvStatModel::**save**(const char* **filename**, const char* **name**=0 )

**Python:** cv2.StatModel.**save**(filename[, name ]) → None

The method save saves the complete model state to the specified XML or YAML file with the specified name or default name (which depends on a particular class). *Data persistence* functionality from CxCore is used.

## CvStatModel::load

Loads the model from a file.

**C++:** void CvStatModel::**load**(const char* **filename**, const char* **name**=0 )

**Python:** cv2.StatModel.**load**(filename[, name ]) → None

The method load loads the complete model state with the specified name (or default model-dependent name) from the specified XML or YAML file. The previous model state is cleared by CvStatModel::clear().

## CvStatModel::write

Writes the model to the file storage.

**C++:** void CvStatModel::**write**(CvFileStorage* **storage**, const char* **name**)

The method write stores the complete model state in the file storage with the specified name or default name (which depends on the particular class). The method is called by CvStatModel::save().

## CvStatModel::read

Reads the model from the file storage.

**C++:** void CvStatModel::**read**(CvFileStorage* **storage**, CvFileNode* **node**)

The method read restores the complete model state from the specified node of the file storage. Use the function GetFileNodeByName() to locate the node.

The previous model state is cleared by CvStatModel::clear().

## CvStatModel::train

Trains the model.

**bool CvStatModel::train( const Mat& train_data, [int tflag,] ..., const Mat& responses, ...,**     **[const Ma**

The method trains the statistical model using a set of input feature vectors and the corresponding output values (responses). Both input and output vectors/values are passed as matrices. By default, the input feature vectors are stored as train_data rows, that is, all the components (features) of a training vector are stored continuously. However, some algorithms can handle the transposed representation when all values of each particular feature (component/input variable) over the whole input set are stored continuously. If both layouts are supported, the method includes the tflag parameter that specifies the orientation as follows:

- tflag=CV_ROW_SAMPLE The feature vectors are stored as rows.

- tflag=CV_COL_SAMPLE The feature vectors are stored as columns.

The `train_data` must have the `CV_32FC1` (32-bit floating-point, single-channel) format. Responses are usually stored in a 1D vector (a row or a column) of `CV_32SC1` (only in the classification problem) or `CV_32FC1` format, one value per input vector. Although, some algorithms, like various flavors of neural nets, take vector responses.

For classification problems, the responses are discrete class labels. For regression problems, the responses are values of the function to be approximated. Some algorithms can deal only with classification problems, some - only with regression problems, and some can deal with both problems. In the latter case, the type of output variable is either passed as a separate parameter or as the last element of the `var_type` vector:

- `CV_VAR_CATEGORICAL` The output values are discrete class labels.

- `CV_VAR_ORDERED(=CV_VAR_NUMERICAL)` The output values are ordered. This means that two different values can be compared as numbers, and this is a regression problem.

Types of input variables can be also specified using `var_type`. Most algorithms can handle only ordered input variables.

Many ML models may be trained on a selected feature subset, and/or on a selected sample subset of the training set. To make it easier for you, the method `train` usually includes the `var_idx` and `sample_idx` parameters. The former parameter identifies variables (features) of interest, and the latter one identifies samples of interest. Both vectors are either integer (`CV_32SC1`) vectors (lists of 0-based indices) or 8-bit (`CV_8UC1`) masks of active variables/samples. You may pass `NULL` pointers instead of either of the arguments, meaning that all of the variables/samples are used for training.

Additionally, some algorithms can handle missing measurements, that is, when certain features of certain training samples have unknown values (for example, they forgot to measure a temperature of patient A on Monday). The parameter `missing_mask`, an 8-bit matrix of the same size as `train_data`, is used to mark the missed values (non-zero elements of the mask).

Usually, the previous model state is cleared by `CvStatModel::clear()` before running the training procedure. However, some algorithms may optionally update the model state with the new training data, instead of resetting it.

### CvStatModel::predict

Predicts the response for a sample.

```
float CvStatModel::predict( const Mat& sample[, <prediction_params>] ) const
```

The method is used to predict the response for a new sample. In case of a classification, the method returns the class label. In case of a regression, the method returns the output function value. The input sample must have as many components as the `train_data` passed to `train` contains. If the `var_idx` parameter is passed to `train`, it is remembered and then is used to extract only the necessary components from the input sample in the method `predict`.

The suffix `const` means that prediction does not affect the internal model state, so the method can be safely called from within different threads.

## 9.2 Normal Bayes Classifier

This simple classification model assumes that feature vectors from each class are normally distributed (though, not necessarily independently distributed). So, the whole data distribution function is assumed to be a Gaussian mixture, one component per class. Using the training data the algorithm estimates mean vectors and covariance matrices for every class, and then it uses them for prediction.

### CvNormalBayesClassifier

```
class CvNormalBayesClassifier : public CvStatModel
```

Bayes classifier for normally distributed data.

## CvNormalBayesClassifier::CvNormalBayesClassifier

Default and training constructors.

**C++:** CvNormalBayesClassifier::**CvNormalBayesClassifier**()

**C++:** CvNormalBayesClassifier::**CvNormalBayesClassifier**(const Mat& **trainData**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat() )

**C++:** CvNormalBayesClassifier::**CvNormalBayesClassifier**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0 )

**Python:** cv2.**NormalBayesClassifier**([trainData, responses[, varIdx[, sampleIdx ] ] ]) → <NormalBayesClassifier object>

The constructors follow conventions of `CvStatModel::CvStatModel()`. See `CvStatModel::train()` for parameters descriptions.

## CvNormalBayesClassifier::train

Trains the model.

**C++:** bool CvNormalBayesClassifier::**train**(const Mat& **trainData**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), bool **update**=false )

**C++:** bool CvNormalBayesClassifier::**train**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, bool **update**=false )

**Python:** cv2.NormalBayesClassifier.**train**(trainData, responses[, varIdx[, sampleIdx[, update ] ] ]) → retval

> **Parameters**
>
> > **update** – Identifies whether the model should be trained from scratch (`update=false`) or should be updated using the new training data (`update=true`).

The method trains the Normal Bayes classifier. It follows the conventions of the generic `CvStatModel::train()` approach with the following limitations:

- Only `CV_ROW_SAMPLE` data layout is supported.

- Input variables are all ordered.

- Output variable is categorical , which means that elements of `responses` must be integer numbers, though the vector may have the `CV_32FC1` type.

- Missing measurements are not supported.

## CvNormalBayesClassifier::predict

Predicts the response for sample(s).

**C++:** float CvNormalBayesClassifier::**predict**(const Mat& **samples**, Mat* **results**=0 ) const

**C++:** float CvNormalBayesClassifier::**predict**(const CvMat* **samples**, CvMat* **results**=0 ) const

**Python:** cv2.NormalBayesClassifier.**predict**(samples) → retval, results

The method estimates the most probable classes for input vectors. Input vectors (one or more) are stored as rows of the matrix `samples`. In case of multiple input vectors, there should be one output vector `results`. The predicted class for a single input vector is returned by the method.

The function is parallelized with the TBB library.

# 9.3 K-Nearest Neighbors

The algorithm caches all training samples and predicts the response for a new sample by analyzing a certain number (**K**) of the nearest neighbors of the sample using voting, calculating weighted sum, and so on. The method is sometimes referred to as "learning by example" because for prediction it looks for the feature vector with a known response that is closest to the given vector.

## CvKNearest

class **CvKNearest** : public **CvStatModel**

The class implements K-Nearest Neighbors model as described in the beginning of this section.

## CvKNearest::CvKNearest

Default and training constructors.

**C++:** CvKNearest::**CvKNearest**()

**C++:** CvKNearest::**CvKNearest**(const Mat& **trainData**, const Mat& **responses**, const Mat& **sampleIdx**=Mat(), bool **isRegression**=false, int **max_k**=32 )

**C++:** CvKNearest::**CvKNearest**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **sampleIdx**=0, bool **isRegression**=false, int **max_k**=32 )

See `CvKNearest::train()` for additional parameters descriptions.

## CvKNearest::train

Trains the model.

**C++:** bool CvKNearest::**train**(const Mat& **trainData**, const Mat& **responses**, const Mat& **sampleIdx**=Mat(), bool **isRegression**=false, int **maxK**=32, bool **updateBase**=false )

**C++:** bool CvKNearest::**train**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **sampleIdx**=0, bool **is_regression**=false, int **maxK**=32, bool **updateBase**=false )

**Python:** cv2.KNearest.**train**(trainData, responses[, sampleIdx[, isRegression[, maxK[, updateBase ]]]]) → retval

> **Parameters**

>> **isRegression** – Type of the problem: `true` for regression and `false` for classification.

>> **maxK** – Number of maximum neighbors that may be passed to the method `CvKNearest::find_nearest()`.

>>>>**updateBase** – Specifies whether the model is trained from scratch (`update_base=false`), or it is updated using the new training data (`update_base=true`). In the latter case, the parameter `maxK` must not be larger than the original value.

The method trains the K-Nearest model. It follows the conventions of the generic `CvStatModel::train()` approach with the following limitations:

- Only `CV_ROW_SAMPLE` data layout is supported.

- Input variables are all ordered.

- Output variables can be either categorical ( `is_regression=false` ) or ordered ( `is_regression=true` ).

- Variable subsets (`var_idx`) and missing measurements are not supported.

## CvKNearest::find_nearest

Finds the neighbors and predicts responses for input vectors.

**C++:** float CvKNearest::**find_nearest**(const Mat& **samples**, int **k**, Mat\* **results**=0, const float\*\* **neighbors**=0, Mat\* **neighborResponses**=0, Mat\* **dist**=0 ) const

**C++:** float CvKNearest::**find_nearest**(const Mat& **samples**, int **k**, Mat& **results**, Mat& **neighborResponses**, Mat& **dists**) const

**C++:** float CvKNearest::**find_nearest**(const CvMat\* **samples**, int **k**, CvMat\* **results**=0, const float\*\* **neighbors**=0, CvMat\* **neighborResponses**=0, CvMat\* **dist**=0 ) const

**Python:** cv2.KNearest.**find_nearest**(samples, k[, results[, neighborResponses[, dists ] ] ]) → retval, results, neighborResponses, dists

>**Parameters**

>>**samples** – Input samples stored by rows. It is a single-precision floating-point matrix of $number\_of\_samples \times number\_of\_features$ size.

>>**k** – Number of used nearest neighbors. It must satisfy constraint: $k \leq$ CvKNearest::get_max_k().

>>**results** – Vector with results of prediction (regression or classification) for each input sample. It is a single-precision floating-point vector with `number_of_samples` elements.

>>**neighbors** – Optional output pointers to the neighbor vectors themselves. It is an array of `k*samples->rows` pointers.

>>**neighborResponses** – Optional output values for corresponding `neighbors`. It is a single-precision floating-point matrix of $number\_of\_samples \times k$ size.

>>**dist** – Optional output distances from the input vectors to the corresponding `neighbors`. It is a single-precision floating-point matrix of $number\_of\_samples \times k$ size.

For each input vector (a row of the matrix `samples`), the method finds the k nearest neighbors. In case of regression, the predicted result is a mean value of the particular vector's neighbor responses. In case of classification, the class is determined by voting.

For each input vector, the neighbors are sorted by their distances to the vector.

In case of C++ interface you can use output pointers to empty matrices and the function will allocate memory itself.

If only a single input vector is passed, all output matrices are optional and the predicted value is returned by the method.

The function is parallelized with the TBB library.

## CvKNearest::get_max_k

Returns the number of maximum neighbors that may be passed to the method `CvKNearest::find_nearest()`.

**C++:** int CvKNearest::**get_max_k**() const

## CvKNearest::get_var_count

Returns the number of used features (variables count).

**C++:** int CvKNearest::**get_var_count**() const

## CvKNearest::get_sample_count

Returns the total number of train samples.

**C++:** int CvKNearest::**get_sample_count**() const

## CvKNearest::is_regression

Returns type of the problem: `true` for regression and `false` for classification.

**C++:** bool CvKNearest::**is_regression**() const

The sample below (currently using the obsolete `CvMat` structures) demonstrates the use of the k-nearest classifier for 2D point classification:

```c
#include "ml.h"
#include "highgui.h"

int main( int argc, char** argv )
{
    const int K = 10;
    int i, j, k, accuracy;
    float response;
    int train_sample_count = 100;
    CvRNG rng_state = cvRNG(-1);
    CvMat* trainData = cvCreateMat( train_sample_count, 2, CV_32FC1 );
    CvMat* trainClasses = cvCreateMat( train_sample_count, 1, CV_32FC1 );
    IplImage* img = cvCreateImage( cvSize( 500, 500 ), 8, 3 );
    float _sample[2];
    CvMat sample = cvMat( 1, 2, CV_32FC1, _sample );
    cvZero( img );

    CvMat trainData1, trainData2, trainClasses1, trainClasses2;

    // form the training samples
    cvGetRows( trainData, &trainData1, 0, train_sample_count/2 );
    cvRandArr( &rng_state, &trainData1, CV_RAND_NORMAL, cvScalar(200,200), cvScalar(50,50) );

    cvGetRows( trainData, &trainData2, train_sample_count/2, train_sample_count );
    cvRandArr( &rng_state, &trainData2, CV_RAND_NORMAL, cvScalar(300,300), cvScalar(50,50) );

    cvGetRows( trainClasses, &trainClasses1, 0, train_sample_count/2 );
    cvSet( &trainClasses1, cvScalar(1) );
```

```
cvGetRows( trainClasses, &trainClasses2, train_sample_count/2, train_sample_count );
cvSet( &trainClasses2, cvScalar(2) );

// learn classifier
CvKNearest knn( trainData, trainClasses, 0, false, K );
CvMat* nearests = cvCreateMat( 1, K, CV_32FC1);

for( i = 0; i < img->height; i++ )
{
    for( j = 0; j < img->width; j++ )
    {
        sample.data.fl[0] = (float)j;
        sample.data.fl[1] = (float)i;

        // estimate the response and get the neighbors' labels
        response = knn.find_nearest(&sample,K,0,0,nearests,0);

        // compute the number of neighbors representing the majority
        for( k = 0, accuracy = 0; k < K; k++ )
        {
            if( nearests->data.fl[k] == response)
                accuracy++;
        }
        // highlight the pixel depending on the accuracy (or confidence)
        cvSet2D( img, i, j, response == 1 ?
            (accuracy > 5 ? CV_RGB(180,0,0) : CV_RGB(180,120,0)) :
            (accuracy > 5 ? CV_RGB(0,180,0) : CV_RGB(120,120,0)) );
    }
}

// display the original training samples
for( i = 0; i < train_sample_count/2; i++ )
{
    CvPoint pt;
    pt.x = cvRound(trainData1.data.fl[i*2]);
    pt.y = cvRound(trainData1.data.fl[i*2+1]);
    cvCircle( img, pt, 2, CV_RGB(255,0,0), CV_FILLED );
    pt.x = cvRound(trainData2.data.fl[i*2]);
    pt.y = cvRound(trainData2.data.fl[i*2+1]);
    cvCircle( img, pt, 2, CV_RGB(0,255,0), CV_FILLED );
}

cvNamedWindow( "classifier result", 1 );
cvShowImage( "classifier result", img );
cvWaitKey(0);

cvReleaseMat( &trainClasses );
cvReleaseMat( &trainData );
return 0;
}
```

# 9.4 Support Vector Machines

Originally, support vector machines (SVM) was a technique for building an optimal binary (2-class) classifier. Later the technique was extended to regression and clustering problems. SVM is a partial case of kernel-based methods. It maps feature vectors into a higher-dimensional space using a kernel function and builds an optimal linear discriminat-

ing function in this space or an optimal hyper-plane that fits into the training data. In case of SVM, the kernel is not defined explicitly. Instead, a distance between any 2 points in the hyper-space needs to be defined.

The solution is optimal, which means that the margin between the separating hyper-plane and the nearest feature vectors from both classes (in case of 2-class classifier) is maximal. The feature vectors that are the closest to the hyper-plane are called *support vectors*, which means that the position of other vectors does not affect the hyper-plane (the decision function).

SVM implementation in OpenCV is based on [LibSVM].

## CvParamGrid

**struct `CvParamGrid`**

>The structure represents the logarithmic grid range of statmodel parameters. It is used for optimizing statmodel accuracy by varying model parameters, the accuracy estimate being computed by cross-validation.

>double `CvParamGrid::`**`min_val`**
>>Minimum value of the statmodel parameter.

>double `CvParamGrid::`**`max_val`**
>>Maximum value of the statmodel parameter.

>double `CvParamGrid::`**`step`**
>>Logarithmic step for iterating the statmodel parameter.

The grid determines the following iteration sequence of the statmodel parameter values:

$$(min\_val, min\_val * step, min\_val * step^2, \ldots, min\_val * step^n),$$

where $n$ is the maximal index satisfying

$$min\_val * step^n < max\_val$$

The grid is logarithmic, so `step` must always be greater then 1.

## CvParamGrid::CvParamGrid

The constructors.

**C++:** `CvParamGrid::`**`CvParamGrid`**`()`

**C++:** `CvParamGrid::`**`CvParamGrid`**`(`double **min_val**, double **max_val**, double **log_step**`)`

The full constructor initializes corresponding members. The default constructor creates a dummy grid:

```
CvParamGrid::CvParamGrid()
{
    min_val = max_val = step = 0;
}
```

## CvParamGrid::check

Checks validness of the grid.

**C++:** bool `CvParamGrid::`**`check`**`()`

Returns `true` if the grid is valid and `false` otherwise. The grid is valid if and only if:

- Lower bound of the grid is less then the upper one.

- Lower bound of the grid is positive.

- Grid step is greater then 1.

## CvSVMParams

**struct** `CvSVMParams`

SVM training parameters.

The structure must be initialized and passed to the training method of `CvSVM`.

## CvSVMParams::CvSVMParams

The constructors.

**C++:** `CvSVMParams::`**`CvSVMParams`**`()`

**C++:** `CvSVMParams::`**`CvSVMParams`**`(int **svm_type**, int **kernel_type**, double **degree**, double **gamma**, double **coef0**, double **Cvalue**, double **nu**, double **p**, CvMat* **class_weights**, CvTermCriteria **term_crit**)`

>   **Parameters**

>>      **svm_type** – Type of a SVM formulation. Possible values are:

>>      – **CvSVM::C_SVC** C-Support Vector Classification. n-class classification ($n \geq 2$), allows imperfect separation of classes with penalty multiplier C for outliers.

>>      – **CvSVM::NU_SVC** $\nu$-Support Vector Classification. n-class classification with possible imperfect separation. Parameter $\nu$ (in the range 0..1, the larger the value, the smoother the decision boundary) is used instead of C.

>>      – **CvSVM::ONE_CLASS** Distribution Estimation (One-class SVM). All the training data are from the same class, SVM builds a boundary that separates the class from the rest of the feature space.

>>      – **CvSVM::EPS_SVR** $\epsilon$-Support Vector Regression. The distance between feature vectors from the training set and the fitting hyper-plane must be less than p. For outliers the penalty multiplier C is used.

>>      – **CvSVM::NU_SVR** $\nu$-Support Vector Regression. $\nu$ is used instead of p.

>>      See [LibSVM] for details.

>>      **kernel_type** – Type of a SVM kernel. Possible values are:

>>      – **CvSVM::LINEAR** Linear kernel. No mapping is done, linear discrimination (or regression) is done in the original feature space. It is the fastest option. $K(x_i, x_j) = x_i^\mathsf{T} x_j$.

>>      – **CvSVM::POLY** Polynomial kernel: $K(x_i, x_j) = (\gamma x_i^\mathsf{T} x_j + coef0)^{degree}, \gamma > 0$.

>>      – **CvSVM::RBF** Radial basis function (RBF), a good choice in most cases. $K(x_i, x_j) = e^{-\gamma ||x_i - x_j||^2}, \gamma > 0$.

>>      – **CvSVM::SIGMOID** Sigmoid kernel: $K(x_i, x_j) = \tanh(\gamma x_i^\mathsf{T} x_j + coef0)$.

>>      **degree** – Parameter `degree` of a kernel function (POLY).

>>      **gamma** – Parameter $\gamma$ of a kernel function (POLY / RBF / SIGMOID).

>>      **coef0** – Parameter `coef0` of a kernel function (POLY / SIGMOID).

> > **Cvalue** – Parameter `C` of a SVM optimization problem (C_SVC / EPS_SVR / NU_SVR).
> >
> > **nu** – Parameter $\nu$ of a SVM optimization problem (NU_SVC / ONE_CLASS / NU_SVR).
> >
> > **p** – Parameter $\epsilon$ of a SVM optimization problem (EPS_SVR).
> >
> > **class_weights** – Optional weights in the C_SVC problem , assigned to particular classes. They are multiplied by `C` so the parameter `C` of class #i becomes $class\_weights_i * C$. Thus these weights affect the misclassification penalty for different classes. The larger weight, the larger penalty on misclassification of data from the corresponding class.
> >
> > **term_crit** – Termination criteria of the iterative SVM training procedure which solves a partial case of constrained quadratic optimization problem. You can specify tolerance and/or the maximum number of iterations.

The default constructor initialize the structure with following values:

```
CvSVMParams::CvSVMParams() :
    svm_type(CvSVM::C_SVC), kernel_type(CvSVM::RBF), degree(0),
    gamma(1), coef0(0), C(1), nu(0), p(0), class_weights(0)
{
    term_crit = cvTermCriteria( CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 1000, FLT_EPSILON );
}
```

## CvSVM

class **CvSVM** : public **CvStatModel**

Support Vector Machines.

## CvSVM::CvSVM

Default and training constructors.

**C++:** CvSVM::**CvSVM**()

**C++:** CvSVM::**CvSVM**(const Mat& **trainData**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), CvSVMParams **params**=CvSVMParams() )

**C++:** CvSVM::**CvSVM**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, CvSVMParams **params**=CvSVMParams() )

**Python:** cv2.**SVM**([trainData, responses[, varIdx[, sampleIdx[, params]]]]) → <SVM object>

The constructors follow conventions of `CvStatModel::CvStatModel()`. See `CvStatModel::train()` for parameters descriptions.

## CvSVM::train

Trains an SVM.

**C++:** bool CvSVM::**train**(const Mat& **trainData**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), CvSVMParams **params**=CvSVMParams() )

**C++:** bool CvSVM::**train**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, CvSVMParams **params**=CvSVMParams() )

**Python:** cv2.SVM.**train**(trainData, responses[, varIdx[, sampleIdx[, params]]]) → retval

The method trains the SVM model. It follows the conventions of the generic `CvStatModel::train()` approach with the following limitations:

- Only the `CV_ROW_SAMPLE` data layout is supported.

- Input variables are all ordered.

- Output variables can be either categorical (`params.svm_type=CvSVM::C_SVC` or `params.svm_type=CvSVM::NU_SVC`), or ordered (`params.svm_type=CvSVM::EPS_SVR` or `params.svm_type=CvSVM::NU_SVR`), or not required at all (`params.svm_type=CvSVM::ONE_CLASS`).

- Missing measurements are not supported.

All the other parameters are gathered in the `CvSVMParams` structure.

## CvSVM::train_auto

Trains an SVM with optimal parameters.

**C++:** bool CvSVM::**train_auto**(const Mat& **trainData**, const Mat& **responses**, const Mat& **varIdx**, const Mat& **sampleIdx**, CvSVMParams **params**, int **k_fold**=10, CvParamGrid **Cgrid**=CvSVM::get_default_grid(CvSVM::C), CvParamGrid **gammaGrid**=CvSVM::get_default_grid(CvSVM::GAMMA), CvParamGrid **pGrid**=CvSVM::get_default_grid(CvSVM::P), CvParamGrid **nuGrid**=CvSVM::get_default_grid(CvSVM::NU), CvParamGrid **coeffGrid**=CvSVM::get_default_grid(CvSVM::COEF), CvParamGrid **degreeGrid**=CvSVM::get_default_grid(CvSVM::DEGREE), bool **balanced**=false)

**C++:** bool CvSVM::**train_auto**(const CvMat* **trainData**, const CvMat* **responses**, const CvMat* **varIdx**, const CvMat* **sampleIdx**, CvSVMParams **params**, int **kfold**=10, CvParamGrid **Cgrid**=get_default_grid(CvSVM::C), CvParamGrid **gammaGrid**=get_default_grid(CvSVM::GAMMA), CvParamGrid **pGrid**=get_default_grid(CvSVM::P), CvParamGrid **nuGrid**=get_default_grid(CvSVM::NU), CvParamGrid **coeffGrid**=get_default_grid(CvSVM::COEF), CvParamGrid **degreeGrid**=get_default_grid(CvSVM::DEGREE), bool **balanced**=false )

**Python:** cv2.SVM.**train_auto**(trainData, responses, varIdx, sampleIdx, params[, k_fold[, Cgrid[, gammaGrid[, pGrid[, nuGrid[, coeffGrid[, degreeGrid[, balanced ]]]]]]]]) → retval

> **Parameters**
>
> > **k_fold** – Cross-validation parameter. The training set is divided into `k_fold` subsets. One subset is used to test the model, the others form the train set. So, the SVM algorithm is executed `k_fold` times.
> >
> > **\*Grid** – Iteration grid for the corresponding SVM parameter.
> >
> > **balanced** – If `true` and the problem is 2-class classification then the method creates more balanced cross-validation subsets that is proportions between classes in subsets are close to such proportion in the whole train dataset.

The method trains the SVM model automatically by choosing the optimal parameters C, gamma, p, nu, coef0, degree from `CvSVMParams`. Parameters are considered optimal when the cross-validation estimate of the test set error is minimal.

If there is no need to optimize a parameter, the corresponding grid step should be set to any value less than or equal to 1. For example, to avoid optimization in gamma, set `gamma_grid.step = 0`, `gamma_grid.min_val`, `gamma_grid.max_val` as arbitrary numbers. In this case, the value `params.gamma` is taken for gamma.

And, finally, if the optimization in a parameter is required but the corresponding grid is unknown, you may call the function `CvSVM::get_default_grid()`. To generate a grid, for example, for `gamma`, call `CvSVM::get_default_grid(CvSVM::GAMMA)`.

This function works for the classification (`params.svm_type=CvSVM::C_SVC` or `params.svm_type=CvSVM::NU_SVC`) as well as for the regression (`params.svm_type=CvSVM::EPS_SVR` or `params.svm_type=CvSVM::NU_SVR`). If `params.svm_type=CvSVM::ONE_CLASS`, no optimization is made and the usual SVM with parameters specified in `params` is executed.

## CvSVM::predict

Predicts the response for input sample(s).

**C++:** float CvSVM::**predict**(const Mat& **sample**, bool **returnDFVal**=false ) `const`

**C++:** float CvSVM::**predict**(const CvMat* **sample**, bool **returnDFVal**=false ) `const`

**C++:** float CvSVM::**predict**(const CvMat* **samples**, CvMat* **results**) `const`

**Python:** cv2.SVM.**predict**(sample[, returnDFVal ]) → retval

> **Parameters**
>
>> **sample** – Input sample for prediction.
>>
>> **samples** – Input samples for prediction.
>>
>> **returnDFVal** – Specifies a type of the return value. If `true` and the problem is 2-class classification then the method returns the decision function value that is signed distance to the margin, else the function returns a class label (classification) or estimated function value (regression).
>>
>> **results** – Output prediction responses for corresponding samples.

If you pass one sample then prediction result is returned. If you want to get responses for several samples then you should pass the `results` matrix where prediction results will be stored.

The function is parallelized with the TBB library.

## CvSVM::get_default_grid

Generates a grid for SVM parameters.

**C++:** CvParamGrid CvSVM::**get_default_grid**(int **param_id**)

> **Parameters**
>
>> **param_id** – SVM parameters IDs that must be one of the following:
>>
>> – **CvSVM::C**
>>
>> – **CvSVM::GAMMA**
>>
>> – **CvSVM::P**
>>
>> – **CvSVM::NU**
>>
>> – **CvSVM::COEF**
>>
>> – **CvSVM::DEGREE**
>>
>> The grid is generated for the parameter with this ID.

The function generates a grid for the specified parameter of the SVM algorithm. The grid may be passed to the function `CvSVM::train_auto()`.

## CvSVM::get_params

Returns the current SVM parameters.

**C++:** CvSVMParams CvSVM::**get_params**() const

This function may be used to get the optimal parameters obtained while automatically training `CvSVM::train_auto()`.

## CvSVM::get_support_vector

Retrieves a number of support vectors and the particular vector.

**C++:** int CvSVM::**get_support_vector_count**() const

**C++:** const float* CvSVM::**get_support_vector**(int **i**) const

**Python:** cv2.SVM.**get_support_vector_count**() → retval

> **Parameters**
>
> > **i** – Index of the particular support vector.

The methods can be used to retrieve a set of support vectors.

## CvSVM::get_var_count

Returns the number of used features (variables count).

**C++:** int CvSVM::**get_var_count**() const

**Python:** cv2.SVM.**get_var_count**() → retval

# 9.5 Decision Trees

The ML classes discussed in this section implement Classification and Regression Tree algorithms described in [Breiman84].

The class `CvDTree` represents a single decision tree that may be used alone or as a base class in tree ensembles (see *Boosting* and *Random Trees* ).

A decision tree is a binary tree (tree where each non-leaf node has two child nodes). It can be used either for classification or for regression. For classification, each tree leaf is marked with a class label; multiple leaves may have the same label. For regression, a constant is also assigned to each tree leaf, so the approximation function is piecewise constant.

## Predicting with Decision Trees

To reach a leaf node and to obtain a response for the input feature vector, the prediction procedure starts with the root node. From each non-leaf node the procedure goes to the left (selects the left child node as the next observed node) or to the right based on the value of a certain variable whose index is stored in the observed node. The following variables are possible:

- **Ordered variables.** The variable value is compared with a threshold that is also stored in the node. If the value is less than the threshold, the procedure goes to the left. Otherwise, it goes to the right. For example, if the weight is less than 1 kilogram, the procedure goes to the left, else to the right.

- **Categorical variables.** A discrete variable value is tested to see whether it belongs to a certain subset of values (also stored in the node) from a limited set of values the variable could take. If it does, the procedure goes to the left. Otherwise, it goes to the right. For example, if the color is green or red, go to the left, else to the right.

So, in each node, a pair of entities (`variable_index`, `decision_rule (threshold/subset)`) is used. This pair is called a *split* (split on the variable `variable_index`). Once a leaf node is reached, the value assigned to this node is used as the output of the prediction procedure.

Sometimes, certain features of the input vector are missed (for example, in the darkness it is difficult to determine the object color), and the prediction procedure may get stuck in the certain node (in the mentioned example, if the node is split by color). To avoid such situations, decision trees use so-called *surrogate splits*. That is, in addition to the best "primary" split, every tree node may also be split to one or more other variables with nearly the same results.

## Training Decision Trees

The tree is built recursively, starting from the root node. All training data (feature vectors and responses) is used to split the root node. In each node the optimum decision rule (the best "primary" split) is found based on some criteria. In machine learning, `gini` "purity" criteria are used for classification, and sum of squared errors is used for regression. Then, if necessary, the surrogate splits are found. They resemble the results of the primary split on the training data. All the data is divided using the primary and the surrogate splits (like it is done in the prediction procedure) between the left and the right child node. Then, the procedure recursively splits both left and right nodes. At each node the recursive procedure may stop (that is, stop splitting the node further) in one of the following cases:

- Depth of the constructed tree branch has reached the specified maximum value.

- Number of training samples in the node is less than the specified threshold when it is not statistically representative to split the node further.

- All the samples in the node belong to the same class or, in case of regression, the variation is too small.

- The best found split does not give any noticeable improvement compared to a random choice.

When the tree is built, it may be pruned using a cross-validation procedure, if necessary. That is, some branches of the tree that may lead to the model overfitting are cut off. Normally, this procedure is only applied to standalone decision trees. Usually tree ensembles build trees that are small enough and use their own protection schemes against overfitting.

## Variable Importance

Besides the prediction that is an obvious use of decision trees, the tree can be also used for various data analyses. One of the key properties of the constructed decision tree algorithms is an ability to compute the importance (relative decisive power) of each variable. For example, in a spam filter that uses a set of words occurred in the message as a feature vector, the variable importance rating can be used to determine the most "spam-indicating" words and thus help keep the dictionary size reasonable.

Importance of each variable is computed over all the splits on this variable in the tree, primary and surrogate ones. Thus, to compute variable importance correctly, the surrogate splits must be enabled in the training parameters, even if there is no missing data.

## CvDTreeSplit

**struct CvDTreeSplit**

    The structure represents a possible decision tree node split. It has public members:

    int **var_idx**

        Index of variable on which the split is created.

    int **inversed**

        If it is not null then inverse split rule is used that is left and right branches are exchanged in the rule expressions below.

    float **quality**

        The split quality, a positive number. It is used to choose the best primary split, then to choose and sort the surrogate splits. After the tree is constructed, it is also used to compute variable importance.

    CvDTreeSplit* **next**

        Pointer to the next split in the node list of splits.

    int[] **subset**

        Bit array indicating the value subset in case of split on a categorical variable. The rule is:

```
if var_value in subset
  then next_node <- left
  else next_node <- right
```

    float ord::**c**

        The threshold value in case of split on an ordered variable. The rule is:

```
if var_value < ord.c
  then next_node<-left
  else next_node<-right
```

    int ord::**split_point**

        Used internally by the training algorithm.

## CvDTreeNode

**struct CvDTreeNode**

    The structure represents a node in a decision tree. It has public members:

    int **class_idx**

        Class index normalized to 0..class_count-1 range and assigned to the node. It is used internally in classification trees and tree ensembles.

    int **Tn**

        Tree index in a ordered sequence of pruned trees. The indices are used during and after the pruning procedure. The root node has the maximum value Tn of the whole tree, child nodes have Tn less than or equal to the parent's Tn, and nodes with $Tn \leq CvDTree :: pruned\_tree\_idx$ are not used at prediction stage (the corresponding branches are considered as cut-off), even if they have not been physically deleted from the tree at the pruning stage.

    double **value**

        Value at the node: a class label in case of classification or estimated function value in case of regression.

    CvDTreeNode* **parent**

        Pointer to the parent node.

    CvDTreeNode* **left**

        Pointer to the left child node.

CvDTreeNode* `right`
    Pointer to the right child node.

CvDTreeSplit* `split`
    Pointer to the first (primary) split in the node list of splits.

int `sample_count`
    The number of samples that fall into the node at the training stage. It is used to resolve the difficult cases - when the variable for the primary split is missing and all the variables for other surrogate splits are missing too. In this case the sample is directed to the left if `left->sample_count > right->sample_count` and to the right otherwise.

int `depth`
    Depth of the node. The root node depth is 0, the child nodes depth is the parent's depth + 1.

Other numerous fields of `CvDTreeNode` are used internally at the training stage.

## CvDTreeParams

**struct `CvDTreeParams`**

The structure contains all the decision tree training parameters. You can initialize it by default constructor and then override any parameters directly before training, or the structure may be fully initialized using the advanced variant of the constructor.

## CvDTreeParams::CvDTreeParams

The constructors.

**C++:** `CvDTreeParams::`**`CvDTreeParams`**`()`

**C++:** `CvDTreeParams::`**`CvDTreeParams`**`(int` **`max_depth`**`, int` **`min_sample_count`**`, float` **`regression_accuracy`**`, bool` **`use_surrogates`**`, int` **`max_categories`**`, int` **`cv_folds`**`, bool` **`use_1se_rule`**`, bool` **`truncate_pruned_tree`**`, const float*` **`priors`**`)`

> Parameters

>> **max_depth** – The maximum possible depth of the tree. That is the training algorithms attempts to split a node while its depth is less than `max_depth`. The actual depth may be smaller if the other termination criteria are met (see the outline of the training procedure in the beginning of the section), and/or if the tree is pruned.

>> **min_sample_count** – If the number of samples in a node is less than this parameter then the node will not be split.

>> **regression_accuracy** – Termination criteria for regression trees. If all absolute differences between an estimated value in a node and values of train samples in this node are less than this parameter then the node will not be split.

>> **use_surrogates** – If true then surrogate splits will be built. These splits allow to work with missing data and compute variable importance correctly.

>> **max_categories** – Cluster possible values of a categorical variable into $K \leq$ `max_categories` clusters to find a suboptimal split. If a discrete variable, on which the training procedure tries to make a split, takes more than `max_categories` values, the precise best subset estimation may take a very long time because the algorithm is exponential. Instead, many decision trees engines (including ML) try to find sub-optimal split in this case by clustering all the samples into `max_categories` clusters that is some categories

are merged together. The clustering is applied only in n>2-class classification problems for categorical variables with `N > max_categories` possible values. In case of regression and 2-class classification the optimal split can be found efficiently without employing clustering, thus the parameter is not used in these cases.

**cv_folds** – If `cv_folds > 1` then prune a tree with K-fold cross-validation where K is equal to `cv_folds`.

**use_1se_rule** – If true then a pruning will be harsher. This will make a tree more compact and more resistant to the training data noise but a bit less accurate.

**truncate_pruned_tree** – If true then pruned branches are physically removed from the tree. Otherwise they are retained and it is possible to get results from the original unpruned (or pruned less aggressively) tree by decreasing `CvDTree::pruned_tree_idx` parameter.

**priors** – The array of a priori class probabilities, sorted by the class label value. The parameter can be used to tune the decision tree preferences toward a certain class. For example, if you want to detect some rare anomaly occurrence, the training base will likely contain much more normal cases than anomalies, so a very good classification performance will be achieved just by considering every case as normal. To avoid this, the priors can be specified, where the anomaly probability is artificially increased (up to 0.5 or even greater), so the weight of the misclassified anomalies becomes much bigger, and the tree is adjusted properly. You can also think about this parameter as weights of prediction categories which determine relative weights that you give to misclassification. That is, if the weight of the first category is 1 and the weight of the second category is 10, then each mistake in predicting the second category is equivalent to making 10 mistakes in predicting the first category.

The default constructor initializes all the parameters with the default values tuned for the standalone classification tree:

```
CvDTreeParams() : max_categories(10), max_depth(INT_MAX), min_sample_count(10),
    cv_folds(10), use_surrogates(true), use_1se_rule(true),
    truncate_pruned_tree(true), regression_accuracy(0.01f), priors(0)
{}
```

## CvDTreeTrainData

**struct `CvDTreeTrainData`**

Decision tree training data and shared data for tree ensembles. The structure is mostly used internally for storing both standalone trees and tree ensembles efficiently. Basically, it contains the following types of information:

1. Training parameters, an instance of `CvDTreeParams`.

2. Training data preprocessed to find the best splits more efficiently. For tree ensembles, this preprocessed data is reused by all trees. Additionally, the training data characteristics shared by all trees in the ensemble are stored here: variable types, the number of classes, a class label compression map, and so on.

3. Buffers, memory storages for tree nodes, splits, and other elements of the constructed trees.

There are two ways of using this structure. In simple cases (for example, a standalone tree or the ready-to-use "black box" tree ensemble from machine learning, like *Random Trees* or *Boosting* ), there is no need to care or even to know about the structure. You just construct the needed statistical model, train it, and use it. The `CvDTreeTrainData` structure is constructed and used internally. However, for custom tree algorithms or another sophisticated cases, the structure may be constructed and used explicitly. The scheme is the following:

1. The structure is initialized using the default constructor, followed by `set_data`, or it is built using the full form of constructor. The parameter `_shared` must be set to `true`.

2. One or more trees are trained using this data (see the special form of the method `CvDTree::train()`).

3. The structure is released as soon as all the trees using it are released.

## CvDTree

class **CvDTree** : public **CvStatModel**

The class implements a decision tree as described in the beginning of this section.

## CvDTree::train

Trains a decision tree.

**C++:** bool CvDTree::**train**(const Mat& **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **varType**=Mat(), const Mat& **missingDataMask**=Mat(), CvDTreeParams **params**=CvDTreeParams() )

**C++:** bool CvDTree::**train**(const CvMat* **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **varType**=0, const CvMat* **missingDataMask**=0, CvDTreeParams **params**=CvDTreeParams() )

**C++:** bool CvDTree::**train**(CvMLData* **trainData**, CvDTreeParams **params**=CvDTreeParams() )

**C++:** bool CvDTree::**train**(CvDTreeTrainData* **trainData**, const CvMat* **subsampleIdx**)

**Python:** cv2.DTree.**train**(trainData, tflag, responses[, varIdx[, sampleIdx[, varType[, missingDataMask[, params]]]]]) → retval

There are four train methods in CvDTree:

- The **first two** methods follow the generic CvStatModel::train() conventions. It is the most complete form. Both data layouts (tflag=CV_ROW_SAMPLE and tflag=CV_COL_SAMPLE) are supported, as well as sample and variable subsets, missing measurements, arbitrary combinations of input and output variable types, and so on. The last parameter contains all of the necessary training parameters (see the CvDTreeParams description).

- The **third** method uses CvMLData to pass training data to a decision tree.

- The **last** method train is mostly used for building tree ensembles. It takes the pre-constructed CvDTreeTrainData instance and an optional subset of the training set. The indices in subsampleIdx are counted relatively to the _sample_idx , passed to the CvDTreeTrainData constructor. For example, if _sample_idx=[1, 5, 7, 100], then subsampleIdx=[0,3] means that the samples [1, 100] of the original training set are used.

The function is parallelized with the TBB library.

## CvDTree::predict

Returns the leaf node of a decision tree corresponding to the input vector.

**C++:** CvDTreeNode* CvDTree::**predict**(const Mat& **sample**, const Mat& **missingDataMask**=Mat(), bool **preprocessedInput**=false ) const

**C++:** CvDTreeNode* CvDTree::**predict**(const CvMat* **sample**, const CvMat* **missingDataMask**=0, bool **preprocessedInput**=false ) const

**Python:** cv2.DTree.**predict**(sample[, missingDataMask[, preprocessedInput]]) → retval

> **Parameters**

> > **sample** – Sample for prediction.

> **missingDataMask** – Optional input missing measurement mask.

> **preprocessedInput** – This parameter is normally set to `false`, implying a regular input. If it is `true`, the method assumes that all the values of the discrete input variables have been already normalized to $0$ to $num\_of\_categories_i - 1$ ranges since the decision tree uses such normalized representation internally. It is useful for faster prediction with tree ensembles. For ordered input variables, the flag is not used.

The method traverses the decision tree and returns the reached leaf node as output. The prediction result, either the class label or the estimated function value, may be retrieved as the `value` field of the `CvDTreeNode` structure, for example: `dtree->predict(sample,mask)->value`.

## CvDTree::calc_error

Returns error of the decision tree.

**C++:** float CvDTree::**calc_error**(CvMLData* **trainData**, int **type**, std::vector<float>* **resp**=0 )

> **Parameters**

>> **trainData** – Data for the decision tree.

>> **type** – Type of error. Possible values are:

>>> – **CV_TRAIN_ERROR** Error on train samples.

>>> – **CV_TEST_ERROR** Error on test samples.

>> **resp** – If it is not null then size of this vector will be set to the number of samples and each element will be set to result of prediction on the corresponding sample.

The method calculates error of the decision tree. In case of classification it is the percentage of incorrectly classified samples and in case of regression it is the mean of squared errors on samples.

## CvDTree::getVarImportance

Returns the variable importance array.

**C++:** Mat CvDTree::**getVarImportance**()

**C++:** const CvMat* CvDTree::**get_var_importance**()

**Python:** cv2.DTree.**getVarImportance**() → retval

## CvDTree::get_root

Returns the root of the decision tree.

**C++:** const CvDTreeNode* CvDTree::**get_root**() const

## CvDTree::get_pruned_tree_idx

Returns the `CvDTree::pruned_tree_idx` parameter.

**C++:** int CvDTree::**get_pruned_tree_idx**() const

The parameter `DTree::pruned_tree_idx` is used to prune a decision tree. See the `CvDTreeNode::Tn` parameter.

## CvDTree::get_data

Returns used train data of the decision tree.

**C++:** CvDTreeTrainData* CvDTree::**get_data**() const

Example: building a tree for classifying mushrooms. See the mushroom.cpp sample that demonstrates how to build and use the decision tree.

# 9.6 Boosting

A common machine learning task is supervised learning. In supervised learning, the goal is to learn the functional relationship $F : y = F(x)$ between the input $x$ and the output $y$. Predicting the qualitative output is called *classification*, while predicting the quantitative output is called *regression*.

Boosting is a powerful learning concept that provides a solution to the supervised classification learning task. It combines the performance of many "weak" classifiers to produce a powerful committee [HTF01]. A weak classifier is only required to be better than chance, and thus can be very simple and computationally inexpensive. However, many of them smartly combine results to a strong classifier that often outperforms most "monolithic" strong classifiers such as SVMs and Neural Networks.

Decision trees are the most popular weak classifiers used in boosting schemes. Often the simplest decision trees with only a single split node per tree (called stumps ) are sufficient.

The boosted model is based on N training examples $(x_i, y_i)1N$ with $x_i \in R^K$ and $y_i \in -1, +1$. $x_i$ is a K -component vector. Each component encodes a feature relevant to the learning task at hand. The desired two-class output is encoded as -1 and +1.

Different variants of boosting are known as Discrete Adaboost, Real AdaBoost, LogitBoost, and Gentle AdaBoost [FHT98]. All of them are very similar in their overall structure. Therefore, this chapter focuses only on the standard two-class Discrete AdaBoost algorithm, outlined below. Initially the same weight is assigned to each sample (step 2). Then, a weak classifier $f_{m(x)}$ is trained on the weighted training data (step 3a). Its weighted training error and scaling factor $c_m$ is computed (step 3b). The weights are increased for training samples that have been misclassified (step 3c). All weights are then normalized, and the process of finding the next weak classifier continues for another M -1 times. The final classifier $F(x)$ is the sign of the weighted sum over the individual weak classifiers (step 4).

**Two-class Discrete AdaBoost Algorithm**

1. Set N examples $(x_i, y_i)1N$ with $x_i \in R^K, y_i \in -1, +1$.

2. Assign weights as $w_i = 1/N, i = 1, ..., N$.

3. Repeat for $m = 1, 2, ..., M$:

   3.1. Fit the classifier $f_m(x) \in -1, 1$, using weights $w_i$ on the training data.

   3.2. Compute $err_m = E_w[1_{(y \neq f_m(x))}], c_m = log((1 - err_m)/err_m)$.

   3.3. Set $w_i \Leftarrow w_i exp[c_m 1_{(y_i \neq f_m(x_i))}], i = 1, 2, ..., N$, and renormalize so that $\Sigma i w_i = 1$.

4. Classify new samples $x$ using the formula: $sign(\Sigma m = 1 M c_m f_m(x))$.

---

**Note:** Similar to the classical boosting methods, the current implementation supports two-class classifiers only. For M > 2 classes, there is the **AdaBoost.MH** algorithm (described in [FHT98]) that reduces the problem to the two-class problem, yet with a much larger training set.

---

To reduce computation time for boosted models without substantially losing accuracy, the influence trimming technique can be employed. As the training algorithm proceeds and the number of trees in the ensemble is increased, a larger number of the training samples are classified correctly and with increasing confidence, thereby those samples

receive smaller weights on the subsequent iterations. Examples with a very low relative weight have a small impact on the weak classifier training. Thus, such examples may be excluded during the weak classifier training without having much effect on the induced classifier. This process is controlled with the `weight_trim_rate` parameter. Only examples with the summary fraction `weight_trim_rate` of the total weight mass are used in the weak classifier training. Note that the weights for **all** training examples are recomputed at each training iteration. Examples deleted at a particular iteration may be used again for learning some of the weak classifiers further [FHT98].

## CvBoostParams

**struct CvBoostParams** : **public CvDTreeParams**

>   Boosting training parameters.

>   There is one structure member that you can set directly:

int **split_criteria**

>   Splitting criteria used to choose optimal splits during a weak tree construction. Possible values are:

>   >   • **CvBoost::DEFAULT** Use the default for the particular boosting method, see below.

>   >   • **CvBoost::GINI** Use Gini index. This is default option for Real AdaBoost; may be also used for Discrete AdaBoost.

>   >   • **CvBoost::MISCLASS** Use misclassification rate. This is default option for Discrete AdaBoost; may be also used for Real AdaBoost.

>   >   • **CvBoost::SQERR** Use least squares criteria. This is default and the only option for LogitBoost and Gentle AdaBoost.

The structure is derived from `CvDTreeParams` but not all of the decision tree parameters are supported. In particular, cross-validation is not supported.

All parameters are public. You can initialize them by a constructor and then override some of them directly if you want.

## CvBoostParams::CvBoostParams

The constructors.

**C++:** CvBoostParams::**CvBoostParams**()

**C++:** CvBoostParams::**CvBoostParams**(int **boost_type**, int **weak_count**, double **weight_trim_rate**, int **max_depth**, bool **use_surrogates**, const float* **priors**)

>   **Parameters**

>   >   **boost_type** – Type of the boosting algorithm. Possible values are:

>   >   >   – **CvBoost::DISCRETE** Discrete AdaBoost.

>   >   >   – **CvBoost::REAL** Real AdaBoost. It is a technique that utilizes confidence-rated predictions and works well with categorical data.

>   >   >   – **CvBoost::LOGIT** LogitBoost. It can produce good regression fits.

>   >   >   – **CvBoost::GENTLE** Gentle AdaBoost. It puts less weight on outlier data points and for that reason is often good with regression data.

>   >   Gentle AdaBoost and Real AdaBoost are often the preferable choices.

>   >   **weak_count** – The number of weak classifiers.

> **weight_trim_rate** – A threshold between 0 and 1 used to save computational time. Samples with summary weight $\leq 1 - weight\_trim\_rate$ do not participate in the *next* iteration of training. Set this parameter to 0 to turn off this functionality.

See `CvDTreeParams::CvDTreeParams()` for description of other parameters.

Default parameters are:

```
CvBoostParams::CvBoostParams()
{
    boost_type = CvBoost::REAL;
    weak_count = 100;
    weight_trim_rate = 0.95;
    cv_folds = 0;
    max_depth = 1;
}
```

## CvBoostTree

**class CvBoostTree** : **public CvDTree**

The weak tree classifier, a component of the boosted tree classifier `CvBoost`, is a derivative of `CvDTree`. Normally, there is no need to use the weak classifiers directly. However, they can be accessed as elements of the sequence `CvBoost::weak`, retrieved by `CvBoost::get_weak_predictors()`.

---

**Note:** In case of LogitBoost and Gentle AdaBoost, each weak predictor is a regression tree, rather than a classification tree. Even in case of Discrete AdaBoost and Real AdaBoost, the `CvBoostTree::predict` return value (`CvDTreeNode::value`) is not an output class label. A negative value "votes" for class #0, a positive value - for class #1. The votes are weighted. The weight of each individual tree may be increased or decreased using the method `CvBoostTree::scale`.

---

## CvBoost

**class CvBoost** : **public CvStatModel**

Boosted tree classifier derived from `CvStatModel`.

## CvBoost::CvBoost

Default and training constructors.

**C++:** CvBoost::**CvBoost**()

**C++:** CvBoost::**CvBoost**(const Mat& **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **varType**=Mat(), const Mat& **missingDataMask**=Mat(), CvBoostParams **params**=CvBoostParams() )

**C++:** CvBoost::**CvBoost**(const CvMat* **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **varType**=0, const CvMat* **missingDataMask**=0, CvBoostParams **params**=CvBoostParams() )

**Python:** cv2.**Boost**([trainData, tflag, responses[, varIdx[, sampleIdx[, varType[, missingDataMask[, params ]]]]]]) → <Boost object>

The constructors follow conventions of `CvStatModel::CvStatModel()`. See `CvStatModel::train()` for parameters descriptions.

---

## CvBoost::train

Trains a boosted tree classifier.

**C++:** bool CvBoost::**train**(const Mat& **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **varType**=Mat(), const Mat& **missingDataMask**=Mat(), CvBoostParams **params**=CvBoostParams(), bool **update**=false )

**C++:** bool CvBoost::**train**(const CvMat* **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **varType**=0, const CvMat* **missingDataMask**=0, CvBoostParams **params**=CvBoostParams(), bool **update**=false )

**C++:** bool CvBoost::**train**(CvMLData* **data**, CvBoostParams **params**=CvBoostParams(), bool **update**=false )

**Python:** cv2.Boost.**train**(trainData, tflag, responses[, varIdx[, sampleIdx[, varType[, missingDataMask[, params[, update ] ] ] ] ] ]) → retval

> **Parameters**
>
> > **update** – Specifies whether the classifier needs to be updated (`true`, the new weak tree classifiers added to the existing ensemble) or the classifier needs to be rebuilt from scratch (`false`).

The train method follows the common template of `CvStatModel::train()`. The responses must be categorical, which means that boosted trees cannot be built for regression, and there should be two classes.

## CvBoost::predict

Predicts a response for an input sample.

**C++:** float CvBoost::**predict**(const Mat& **sample**, const Mat& **missing**=Mat(), const Range& **slice**=Range::all(), bool **rawMode**=false, bool **returnSum**=false ) const

**C++:** float CvBoost::**predict**(const CvMat* **sample**, const CvMat* **missing**=0, CvMat* **weak_responses**=0, CvSlice **slice**=CV_WHOLE_SEQ, bool **raw_mode**=false, bool **return_sum**=false ) const

**Python:** cv2.Boost.**predict**(sample[, missing[, slice[, rawMode[, returnSum ] ] ] ]) → retval

> **Parameters**
>
> > **sample** – Input sample.
> >
> > **missing** – Optional mask of missing measurements. To handle missing measurements, the weak classifiers must include surrogate splits (see `CvDTreeParams::use_surrogates`).
> >
> > **weak_responses** – Optional output parameter, a floating-point vector with responses of each individual weak classifier. The number of elements in the vector must be equal to the slice length.
> >
> > **slice** – Continuous subset of the sequence of weak classifiers to be used for prediction. By default, all the weak classifiers are used.
> >
> > **raw_mode** – Normally, it should be set to `false`.
> >
> > **return_sum** – If `true` then return sum of votes instead of the class label.

The method runs the sample through the trees in the ensemble and returns the output class label based on the weighted voting.

## CvBoost::prune

Removes the specified weak classifiers.

**C++:** void CvBoost::**prune**(CvSlice **slice**)

**Python:** cv2.Boost.**prune**(slice) → None

> **Parameters**
>
> > **slice** – Continuous subset of the sequence of weak classifiers to be removed.

The method removes the specified weak classifiers from the sequence.

---

**Note:** Do not confuse this method with the pruning of individual decision trees, which is currently not supported.

---

## CvBoost::calc_error

Returns error of the boosted tree classifier.

**C++:** float CvBoost::**calc_error**(CvMLData* **_data**, int **type**, std::vector<float>* **resp**=0 )

The method is identical to `CvDTree::calc_error()` but uses the boosted tree classifier as predictor.

## CvBoost::get_weak_predictors

Returns the sequence of weak tree classifiers.

**C++:** CvSeq* CvBoost::**get_weak_predictors**()

The method returns the sequence of weak classifiers. Each element of the sequence is a pointer to the `CvBoostTree` class or to some of its derivatives.

## CvBoost::get_params

Returns current parameters of the boosted tree classifier.

**C++:** const CvBoostParams& CvBoost::**get_params**() const

## CvBoost::get_data

Returns used train data of the boosted tree classifier.

**C++:** const CvDTreeTrainData* CvBoost::**get_data**() const

# 9.7 Gradient Boosted Trees

Gradient Boosted Trees (GBT) is a generalized boosting algorithm introduced by Jerome Friedman: http://www.salfordsystems.com/doc/GreedyFuncApproxSS.pdf . In contrast to the AdaBoost.M1 algorithm, GBT can deal with both multiclass classification and regression problems. Moreover, it can use any differential loss function, some popular ones are implemented. Decision trees (`CvDTree`) usage as base learners allows to process ordered and categorical variables.

---

## Training the GBT model

Gradient Boosted Trees model represents an ensemble of single regression trees built in a greedy fashion. Training procedure is an iterative process similar to the numerical optimization via the gradient descent method. Summary loss on the training set depends only on the current model predictions for the training samples, in other words $\sum_{i=1}^{N} L(y_i, F(x_i)) \equiv \mathcal{L}(F(x_1), F(x_2), ..., F(x_N)) \equiv \mathcal{L}(F)$. And the $\mathcal{L}(F)$ gradient can be computed as follows:

$$grad(\mathcal{L}(F)) = \left( \frac{\partial L(y_1, F(x_1))}{\partial F(x_1)}, \frac{\partial L(y_2, F(x_2))}{\partial F(x_2)}, ..., \frac{\partial L(y_N, F(x_N))}{\partial F(x_N)} \right).$$

At every training step, a single regression tree is built to predict an antigradient vector components. Step length is computed corresponding to the loss function and separately for every region determined by the tree leaf. It can be eliminated by changing values of the leaves directly.

See below the main scheme of the training process:

1. Find the best constant model.

2. For $i$ in $[1, M]$:

    (a) Compute the antigradient.

    (b) Grow a regression tree to predict antigradient components.

    (c) Change values in the tree leaves.

    (d) Add the tree to the model.

The following loss functions are implemented for regression problems:

- Squared loss (`CvGBTrees::SQUARED_LOSS`): $L(y, f(x)) = \frac{1}{2}(y - f(x))^2$

- Absolute loss (`CvGBTrees::ABSOLUTE_LOSS`): $L(y, f(x)) = |y - f(x)|$

- Huber loss (`CvGBTrees::HUBER_LOSS`): $L(y, f(x)) = \begin{cases} \delta \cdot \left( |y - f(x)| - \dfrac{\delta}{2} \right) & : |y - f(x)| > \delta \\ \dfrac{1}{2} \cdot (y - f(x))^2 & : |y - f(x)| \le \delta \end{cases}$,

    where $\delta$ is the $\alpha$-quantile estimation of the $|y - f(x)|$. In the current implementation $\alpha = 0.2$.

The following loss functions are implemented for classification problems:

- Deviance or cross-entropy loss (`CvGBTrees::DEVIANCE_LOSS`): K functions are built, one function for each output class, and $L(y, f_1(x), ..., f_K(x)) = -\sum_{k=0}^{K} 1(y = k) \ln p_k(x)$, where $p_k(x) = \dfrac{\exp f_k(x)}{\sum_{i=1}^{K} \exp f_i(x)}$ is the estimation of the probability of $y = k$.

As a result, you get the following model:

$$f(x) = f_0 + \nu \cdot \sum_{i=1}^{M} T_i(x),$$

where $f_0$ is the initial guess (the best constant model) and $\nu$ is a regularization parameter from the interval $(0, 1]$, further called *shrinkage*.

## Predicting with the GBT Model

To get the GBT model prediction, you need to compute the sum of responses of all the trees in the ensemble. For regression problems, it is the answer. For classification problems, the result is $\arg\max_{i=1..K}(f_i(x))$.

## CvGBTreesParams

**struct `CvGBTreesParams` : public `CvDTreeParams`**

GBT training parameters.

The structure contains parameters for each single decision tree in the ensemble, as well as the whole model characteristics. The structure is derived from `CvDTreeParams` but not all of the decision tree parameters are supported: cross-validation, pruning, and class priorities are not used.

## CvGBTreesParams::CvGBTreesParams

**C++:** `CvGBTreesParams::`**`CvGBTreesParams`**`()`

**C++:** `CvGBTreesParams::`**`CvGBTreesParams`**`(int **loss_function_type**, int **weak_count**, float **shrinkage**, float **subsample_portion**, int **max_depth**, bool **use_surrogates**)`

> **Parameters**

> > **loss_function_type** – Type of the loss function used for training (see *Training the GBT model*). It must be one of the following types: `CvGBTrees::SQUARED_LOSS`, `CvGBTrees::ABSOLUTE_LOSS`, `CvGBTrees::HUBER_LOSS`, `CvGBTrees::DEVIANCE_LOSS`. The first three types are used for regression problems, and the last one for classification.

> > **weak_count** – Count of boosting algorithm iterations. `weak_count*K` is the total count of trees in the GBT model, where K is the output classes count (equal to one in case of a regression).

> > **shrinkage** – Regularization parameter (see *Training the GBT model*).

> > **subsample_portion** – Portion of the whole training set used for each algorithm iteration. Subset is generated randomly. For more information see http://www.salfordsystems.com/doc/StochasticBoostingSS.pdf.

> > **max_depth** – Maximal depth of each decision tree in the ensemble (see `CvDTree`).

> > **use_surrogates** – If `true`, surrogate splits are built (see `CvDTree`).

By default the following constructor is used:

```
CvGBTreesParams(CvGBTrees::SQUARED_LOSS, 200, 0.8f, 0.01f, 3, false)
    : CvDTreeParams( 3, 10, 0, false, 10, 0, false, false, 0 )
```

## CvGBTrees

**class `CvGBTrees` : public `CvStatModel`**

The class implements the Gradient boosted tree model as described in the beginning of this section.

## CvGBTrees::CvGBTrees

Default and training constructors.

**C++:** `CvGBTrees::``CvGBTrees``()`

**C++:** `CvGBTrees::``CvGBTrees``(const Mat&` **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **var-Type**=Mat(), const Mat& **missingDataMask**=Mat(), CvGBTreesParams **params**=CvGBTreesParams() )

**C++:** `CvGBTrees::``CvGBTrees``(const CvMat*` **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **var-Type**=0, const CvMat* **missingDataMask**=0, CvGBTreesParams **params**=CvGBTreesParams() )

**Python:** `cv2.``GBTrees``(`$\big[$trainData, tflag, responses$\big[$, varIdx$\big[$, sampleIdx$\big[$, varType$\big[$, missingDataMask$\big[$, params$\big]\big]\big]\big]\big]\big]$) → <GBTrees object>

The constructors follow conventions of `CvStatModel::CvStatModel()`. See `CvStatModel::train()` for parameters descriptions.

## CvGBTrees::train

Trains a Gradient boosted tree model.

**C++:** bool `CvGBTrees::``train``(const Mat&` **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **var-Type**=Mat(), const Mat& **missingDataMask**=Mat(), CvGBTreesParams **params**=CvGBTreesParams(), bool **update**=false )

**C++:** bool `CvGBTrees::``train``(const CvMat*` **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **var-Type**=0, const CvMat* **missingDataMask**=0, CvGBTreesParams **params**=CvGBTreesParams(), bool **update**=false )

**C++:** bool `CvGBTrees::``train``(CvMLData*` **data**, CvGBTreesParams **params**=CvGBTreesParams(), bool **update**=false )

**Python:** `cv2.GBTrees.``train``(`trainData, tflag, responses$\big[$, varIdx$\big[$, sampleIdx$\big[$, varType$\big[$, missingData-Mask$\big[$, params$\big[$, update$\big]\big]\big]\big]\big]\big]$) → retval

The first train method follows the common template (see `CvStatModel::train()`). Both tflag values (`CV_ROW_SAMPLE`, `CV_COL_SAMPLE`) are supported. `trainData` must be of the `CV_32F` type. `responses` must be a matrix of type `CV_32S` or `CV_32F`. In both cases it is converted into the `CV_32F` matrix inside the training procedure. `varIdx` and `sampleIdx` must be a list of indices (`CV_32S`) or a mask (`CV_8U` or `CV_8S`). `update` is a dummy parameter.

The second form of `CvGBTrees::train()` function uses `CvMLData` as a data set container. `update` is still a dummy parameter.

All parameters specific to the GBT model are passed into the training function as a `CvGBTreesParams` structure.

## CvGBTrees::predict

Predicts a response for an input sample.

**C++:** float `CvGBTrees::``predict``(const Mat&` **sample**, const Mat& **missing**=Mat(), const Range& **slice**=Range::all(), int **k**=-1) const

**C++:** float CvGBTrees::**predict**(const CvMat* **sample**, const CvMat* **missing**=0, CvMat* **weakResponses**=0, CvSlice **slice**=CV_WHOLE_SEQ, int **k**=-1 ) const

**Python:** cv2.GBTrees.**predict**(sample[, missing[, slice[, k ] ] ]) → retval

> **Parameters**

>> **sample** – Input feature vector that has the same format as every training set element. If not all the variables were actually used during training, sample contains forged values at the appropriate places.

>> **missing** – Missing values mask, which is a dimensional matrix of the same size as sample having the CV_8U type. 1 corresponds to the missing value in the same position in the sample vector. If there are no missing values in the feature vector, an empty matrix can be passed instead of the missing mask.

>> **weakResponses** – Matrix used to obtain predictions of all the trees. The matrix has K rows, where K is the count of output classes (1 for the regression case). The matrix has as many columns as the slice length.

>> **slice** – Parameter defining the part of the ensemble used for prediction. If slice = Range::all(), all trees are used. Use this parameter to get predictions of the GBT models with different ensemble sizes learning only one model.

>> **k** – Number of tree ensembles built in case of the classification problem (see *Training the GBT model*). Use this parameter to change the output to sum of the trees' predictions in the k-th ensemble only. To get the total GBT model prediction, k value must be -1. For regression problems, k is also equal to -1.

The method predicts the response corresponding to the given sample (see *Predicting with the GBT Model*). The result is either the class label or the estimated function value. The CvGBTrees::predict() method enables using the parallel version of the GBT model prediction if the OpenCV is built with the TBB library. In this case, predictions of single trees are computed in a parallel fashion.

## CvGBTrees::clear

Clears the model.

**C++:** void CvGBTrees::**clear**()

**Python:** cv2.GBTrees.**clear**() → None

The function deletes the data set information and all the weak models and sets all internal variables to the initial state. The function is called in CvGBTrees::train() and in the destructor.

## CvGBTrees::calc_error

Calculates a training or testing error.

**C++:** float CvGBTrees::**calc_error**(CvMLData* **_data**, int **type**, std::vector<float>* **resp**=0 )

> **Parameters**

>> **_data** – Data set.

>> **type** – Parameter defining the error that should be computed: train (CV_TRAIN_ERROR) or test (CV_TEST_ERROR).

>> **resp** – If non-zero, a vector of predictions on the corresponding data set is returned.

If the `CvMLData` data is used to store the data set, `CvGBTrees::calc_error()` can be used to get a training/testing error easily and (optionally) all predictions on the training/testing set. If the Intel* TBB* library is used, the error is computed in a parallel way, namely, predictions for different samples are computed at the same time. In case of a regression problem, a mean squared error is returned. For classifications, the result is a misclassification error in percent.

# 9.8 Random Trees

Random trees have been introduced by Leo Breiman and Adele Cutler: http://www.stat.berkeley.edu/users/breiman/RandomForests/ . The algorithm can deal with both classification and regression problems. Random trees is a collection (ensemble) of tree predictors that is called *forest* further in this section (the term has been also introduced by L. Breiman). The classification works as follows: the random trees classifier takes the input feature vector, classifies it with every tree in the forest, and outputs the class label that received the majority of "votes". In case of a regression, the classifier response is the average of the responses over all the trees in the forest.

All the trees are trained with the same parameters but on different training sets. These sets are generated from the original training set using the bootstrap procedure: for each training set, you randomly select the same number of vectors as in the original set ( `=N` ). The vectors are chosen with replacement. That is, some vectors will occur more than once and some will be absent. At each node of each trained tree, not all the variables are used to find the best split, but a random subset of them. With each node a new subset is generated. However, its size is fixed for all the nodes and all the trees. It is a training parameter set to $\sqrt{number\_of\_variables}$ by default. None of the built trees are pruned.

In random trees there is no need for any accuracy estimation procedures, such as cross-validation or bootstrap, or a separate test set to get an estimate of the training error. The error is estimated internally during the training. When the training set for the current tree is drawn by sampling with replacement, some vectors are left out (so-called *oob (out-of-bag) data* ). The size of oob data is about `N/3` . The classification error is estimated by using this oob-data as follows:

1. Get a prediction for each vector, which is oob relative to the i-th tree, using the very i-th tree.

2. After all the trees have been trained, for each vector that has ever been oob, find the class-*winner* for it (the class that has got the majority of votes in the trees where the vector was oob) and compare it to the ground-truth response.

3. Compute the classification error estimate as a ratio of the number of misclassified oob vectors to all the vectors in the original data. In case of regression, the oob-error is computed as the squared error for oob vectors difference divided by the total number of vectors.

For the random trees usage example, please, see letter_recog.cpp sample in OpenCV distribution.

**References:**

- *Machine Learning*, Wald I, July 2002. http://stat-www.berkeley.edu/users/breiman/wald2002-1.pdf

- *Looking Inside the Black Box*, Wald II, July 2002. http://stat-www.berkeley.edu/users/breiman/wald2002-2.pdf

- *Software for the Masses*, Wald III, July 2002. http://stat-www.berkeley.edu/users/breiman/wald2002-3.pdf

- And other articles from the web site http://www.stat.berkeley.edu/users/breiman/RandomForests/cc_home.htm

## CvRTParams

**struct CvRTParams** : **public CvDTreeParams**
    Training parameters of random trees.

The set of training parameters for the forest is a superset of the training parameters for a single tree. However, random trees do not need all the functionality/features of decision trees. Most noticeably, the trees are not pruned, so the cross-validation parameters are not used.

## CvRTParams::CvRTParams:

The constructors.

**C++:** CvRTParams::**CvRTParams**()

**C++:** CvRTParams::**CvRTParams**(int **max_depth**, int **min_sample_count**, float **regression_accuracy**, bool **use_surrogates**, int **max_categories**, const float* **priors**, bool **calc_var_importance**, int **nactive_vars**, int **max_num_of_trees_in_the_forest**, float **forest_accuracy**, int **termcrit_type**)

> **Parameters**

>> **max_depth** – the depth of the tree. A low value will likely underfit and conversely a high value will likely overfit. The optimal value can be obtained using cross validation or other suitable methods.

>> **min_sample_count** – minimum samples required at a leaf node for it to be split. A reasonable value is a small percentage of the total data e.g. 1%.

>> **max_categories** – Cluster possible values of a categorical variable into $K \leq$ `max_categories` clusters to find a suboptimal split. If a discrete variable, on which the training procedure tries to make a split, takes more than `max_categories` values, the precise best subset estimation may take a very long time because the algorithm is exponential. Instead, many decision trees engines (including ML) try to find sub-optimal split in this case by clustering all the samples into `max_categories` clusters that is some categories are merged together. The clustering is applied only in n>2-class classification problems for categorical variables with `N > max_categories` possible values. In case of regression and 2-class classification the optimal split can be found efficiently without employing clustering, thus the parameter is not used in these cases.

>> **calc_var_importance** – If true then variable importance will be calculated and then it can be retrieved by `CvRTrees::get_var_importance()`.

>> **nactive_vars** – The size of the randomly selected subset of features at each tree node and that are used to find the best split(s). If you set it to 0 then the size will be set to the square root of the total number of features.

>> **max_num_of_trees_in_the_forest** – The maximum number of trees in the forest (surprise, surprise). Typically the more trees you have the better the accuracy. However, the improvement in accuracy generally diminishes and asymptotes pass a certain number of trees. Also to keep in mind, the number of tree increases the prediction time linearly.

>> **forest_accuracy** – Sufficient accuracy (OOB error).

>> **termcrit_type** – The type of the termination criteria:

>>> – **CV_TERMCRIT_ITER** Terminate learning by the `max_num_of_trees_in_the_forest`;

>>> – **CV_TERMCRIT_EPS** Terminate learning by the `forest_accuracy`;

>>> – **CV_TERMCRIT_ITER | CV_TERMCRIT_EPS** Use both termination criteria.

For meaning of other parameters see `CvDTreeParams::CvDTreeParams()`.

The default constructor sets all parameters to default values which are different from default values of `CvDTreeParams`:

```
CvRTParams::CvRTParams() : CvDTreeParams( 5, 10, 0, false, 10, 0, false, false, 0 ),
    calc_var_importance(false), nactive_vars(0)
{
    term_crit = cvTermCriteria( CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 50, 0.1 );
}
```

# CvRTrees

**class `CvRTrees` : public `CvStatModel`**
> The class implements the random forest predictor as described in the beginning of this section.

# CvRTrees::train

Trains the Random Trees model.

**C++:** bool CvRTrees::**train**(const Mat& **trainData**, int **tflag**, const Mat& **responses**, const Mat& **varIdx**=Mat(), const Mat& **sampleIdx**=Mat(), const Mat& **varType**=Mat(), const Mat& **missingDataMask**=Mat(), CvRTParams **params**=CvRTParams() )

**C++:** bool CvRTrees::**train**(const CvMat* **trainData**, int **tflag**, const CvMat* **responses**, const CvMat* **varIdx**=0, const CvMat* **sampleIdx**=0, const CvMat* **varType**=0, const CvMat* **missingDataMask**=0, CvRTParams **params**=CvRTParams() )

**C++:** bool CvRTrees::**train**(CvMLData* **data**, CvRTParams **params**=CvRTParams() )

**Python:** cv2.RTrees.**train**(trainData, tflag, responses[, varIdx[, sampleIdx[, varType[, missingDataMask[, params ] ] ] ] ]) → retval

The method `CvRTrees::train()` is very similar to the method `CvDTree::train()` and follows the generic method CvStatModel::train() conventions. All the parameters specific to the algorithm training are passed as a `CvRTParams` instance. The estimate of the training error (`oob-error`) is stored in the protected class member `oob_error`.

The function is parallelized with the TBB library.

# CvRTrees::predict

Predicts the output for an input sample.

**C++:** float CvRTrees::**predict**(const Mat& **sample**, const Mat& **missing**=Mat() ) const

**C++:** float CvRTrees::**predict**(const CvMat* **sample**, const CvMat* **missing**=0 ) const

**Python:** cv2.RTrees.**predict**(sample[, missing ]) → retval

> **Parameters**
>> **sample** – Sample for classification.
>> **missing** – Optional missing measurement mask of the sample.

The input parameters of the prediction method are the same as in `CvDTree::predict()` but the return value type is different. This method returns the cumulative result from all the trees in the forest (the class that receives the majority of voices, or the mean of the regression function estimates).

## CvRTrees::predict_prob

Returns a fuzzy-predicted class label.

**C++:** float CvRTrees::**predict_prob**(const cv::Mat& **sample**, const cv::Mat& **missing**=cv::Mat() ) const

**C++:** float CvRTrees::**predict_prob**(const CvMat* **sample**, const CvMat* **missing**=0 ) const

**Python:** cv2.RTrees.**predict_prob**(sample[, missing ]) → retval

> **Parameters**
>
> > **sample** – Sample for classification.
> >
> > **missing** – Optional missing measurement mask of the sample.

The function works for binary classification problems only. It returns the number between 0 and 1. This number represents probability or confidence of the sample belonging to the second class. It is calculated as the proportion of decision trees that classified the sample to the second class.

## CvRTrees::getVarImportance

Returns the variable importance array.

**C++:** Mat CvRTrees::**getVarImportance**()

**C++:** const CvMat* CvRTrees::**get_var_importance**()

**Python:** cv2.RTrees.**getVarImportance**() → retval

The method returns the variable importance vector, computed at the training stage when CvRTParams::calc_var_importance is set to true. If this flag was set to false, the NULL pointer is returned. This differs from the decision trees where variable importance can be computed anytime after the training.

## CvRTrees::get_proximity

Retrieves the proximity measure between two training samples.

**C++:** float CvRTrees::**get_proximity**(const CvMat* **sample1**, const CvMat* **sample2**, const CvMat* **missing1**=0, const CvMat* **missing2**=0 ) const

> **Parameters**
>
> > **sample1** – The first sample.
> >
> > **sample2** – The second sample.
> >
> > **missing1** – Optional missing measurement mask of the first sample.
> >
> > **missing2** – Optional missing measurement mask of the second sample.

The method returns proximity measure between any two samples. This is a ratio of those trees in the ensemble, in which the samples fall into the same leaf node, to the total number of the trees.

## CvRTrees::calc_error

Returns error of the random forest.

**C++:** float CvRTrees::**calc_error**(CvMLData* **data**, int **type**, std::vector<float>* **resp**=0 )

The method is identical to CvDTree::calc_error() but uses the random forest as predictor.

---

## CvRTrees::get_train_error

Returns the train error.

**C++:** float CvRTrees::**get_train_error**()

The method works for classification problems only. It returns the proportion of incorrectly classified train samples.

## CvRTrees::get_rng

Returns the state of the used random number generator.

**C++:** CvRNG* CvRTrees::**get_rng**()

## CvRTrees::get_tree_count

Returns the number of trees in the constructed random forest.

**C++:** int CvRTrees::**get_tree_count**() const

## CvRTrees::get_tree

Returns the specific decision tree in the constructed random forest.

**C++:** CvForestTree* CvRTrees::**get_tree**(int **i**) const

> **Parameters**
>
> > **i** – Index of the decision tree.

# 9.9 Extremely randomized trees

Extremely randomized trees have been introduced by Pierre Geurts, Damien Ernst and Louis Wehenkel in the article "Extremely randomized trees", 2006 [http://citeseerx.ist.psu.edu/viewdoc/download?doi=10.1.1.65.7485&rep=rep1&type=pdf]. The algorithm of growing Extremely randomized trees is similar to *Random Trees* (Random Forest), but there are two differences:

1. Extremely randomized trees don't apply the bagging procedure to construct a set of the training samples for each tree. The same input training set is used to train all trees.

2. Extremely randomized trees pick a node split very extremely (both a variable index and variable splitting value are chosen randomly), whereas Random Forest finds the best split (optimal one by variable index and variable splitting value) among random subset of variables.

## CvERTrees

**class** `CvERTrees` : **public** `CvRTrees`

> The class implements the Extremely randomized trees algorithm. `CvERTrees` is inherited from `CvRTrees` and has the same interface, so see description of `CvRTrees` class to get details. To set the training parameters of Extremely randomized trees the same class `CvRTParams` is used.

## 9.10 Expectation Maximization

The Expectation Maximization(EM) algorithm estimates the parameters of the multivariate probability density function in the form of a Gaussian mixture distribution with a specified number of mixtures.

Consider the set of the N feature vectors $\{ x_1, x_2, ..., x_N \}$ from a d-dimensional Euclidean space drawn from a Gaussian mixture:

$$p(x; a_k, S_k, \pi_k) = \sum_{k=1}^{m} \pi_k p_k(x), \quad \pi_k \geq 0, \quad \sum_{k=1}^{m} \pi_k = 1,$$

$$p_k(x) = \varphi(x; a_k, S_k) = \frac{1}{(2\pi)^{d/2} \mid S_k \mid^{1/2}} exp \left\{ -\frac{1}{2}(x - a_k)^T S_k^{-1}(x - a_k) \right\},$$

where $m$ is the number of mixtures, $p_k$ is the normal distribution density with the mean $a_k$ and covariance matrix $S_k$, $\pi_k$ is the weight of the k-th mixture. Given the number of mixtures $M$ and the samples $x_i$, $i = 1..N$ the algorithm finds the maximum-likelihood estimates (MLE) of all the mixture parameters, that is, $a_k$, $S_k$ and $\pi_k$ :

$$L(x, \theta) = logp(x, \theta) = \sum_{i=1}^{N} log \left( \sum_{k=1}^{m} \pi_k p_k(x) \right) \rightarrow \max_{\theta \in \Theta},$$

$$\Theta = \left\{ (a_k, S_k, \pi_k) : a_k \in \mathbb{R}^d, S_k = S_k^T > 0, S_k \in \mathbb{R}^{d \times d}, \pi_k \geq 0, \sum_{k=1}^{m} \pi_k = 1 \right\}.$$

The EM algorithm is an iterative procedure. Each iteration includes two steps. At the first step (Expectation step or E-step), you find a probability $p_{i,k}$ (denoted $\alpha_{i,k}$ in the formula below) of sample $i$ to belong to mixture $k$ using the currently available mixture parameter estimates:

$$\alpha_{ki} = \frac{\pi_k \varphi(x; a_k, S_k)}{\sum_{j=1}^{m} \pi_j \varphi(x; a_j, S_j)}.$$

At the second step (Maximization step or M-step), the mixture parameter estimates are refined using the computed probabilities:

$$\pi_k = \frac{1}{N} \sum_{i=1}^{N} \alpha_{ki}, \quad a_k = \frac{\sum_{i=1}^{N} \alpha_{ki} x_i}{\sum_{i=1}^{N} \alpha_{ki}}, \quad S_k = \frac{\sum_{i=1}^{N} \alpha_{ki}(x_i - a_k)(x_i - a_k)^T}{\sum_{i=1}^{N} \alpha_{ki}}$$

Alternatively, the algorithm may start with the M-step when the initial values for $p_{i,k}$ can be provided. Another alternative when $p_{i,k}$ are unknown is to use a simpler clustering algorithm to pre-cluster the input samples and thus obtain initial $p_{i,k}$ . Often (including machine learning) the kmeans() algorithm is used for that purpose.

One of the main problems of the EM algorithm is a large number of parameters to estimate. The majority of the parameters reside in covariance matrices, which are $d \times d$ elements each where d is the feature space dimensionality. However, in many practical problems, the covariance matrices are close to diagonal or even to $\mu_k * I$ , where I is an identity matrix and $\mu_k$ is a mixture-dependent "scale" parameter. So, a robust computation scheme could start with harder constraints on the covariance matrices and then use the estimated parameters as an input for a less constrained optimization problem (often a diagonal covariance matrix is already a good enough approximation).

**References:**

- Bilmes98 J. A. Bilmes. *A Gentle Tutorial of the EM Algorithm and its Application to Parameter Estimation for Gaussian Mixture and Hidden Markov Models*. Technical Report TR-97-021, International Computer Science Institute and Computer Science Division, University of California at Berkeley, April 1998.

# EM

**class EM** : **public Algorithm**

The class implements the EM algorithm as described in the beginning of this section. It is inherited from `Algorithm`.

## EM::EM

The constructor of the class

**C++:** EM::**EM**(int nclusters=EM::DEFAULT_NCLUSTERS, int covMat-
Type=EM::COV_MAT_DIAGONAL, const TermCriteria& term-
Crit=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS,
EM::DEFAULT_MAX_ITERS, FLT_EPSILON) )

**Python:** cv2.**EM**([nclusters[, covMatType[, termCrit]]]) → <EM object>

> **Parameters**

>> **nclusters** – The number of mixture components in the Gaussian mixture model. Default
>> value of the parameter is `EM::DEFAULT_NCLUSTERS=5`. Some of EM implementation could
>> determine the optimal number of mixtures within a specified value range, but that is not the
>> case in ML yet.

>> **covMatType** – Constraint on covariance matrices which defines type of matrices. Possible
>> values are:

>> – **EM::COV_MAT_SPHERICAL** A scaled identity matrix $\mu_k * I$. There is the only
>> parameter $\mu_k$ to be estimated for each matrix. The option may be used in special
>> cases, when the constraint is relevant, or as a first step in the optimization (for ex-
>> ample in case when the data is preprocessed with PCA). The results of such prelim-
>> inary estimation may be passed again to the optimization procedure, this time with
>> `covMatType=EM::COV_MAT_DIAGONAL`.

>> – **EM::COV_MAT_DIAGONAL** A diagonal matrix with positive diagonal elements. The
>> number of free parameters is `d` for each matrix. This is most commonly used option
>> yielding good estimation results.

>> – **EM::COV_MAT_GENERIC** A symmetric positively defined matrix. The number of
>> free parameters in each matrix is about $d^2/2$. It is not recommended to use this option,
>> unless there is pretty accurate initial estimation of the parameters and/or a huge number
>> of training samples.

>> **termCrit** – The termination criteria of the EM algorithm. The EM algorithm can be ter-
>> minated by the number of iterations `termCrit.maxCount` (number of M-steps) or when
>> relative change of likelihood logarithm is less than `termCrit.epsilon`. Default maximum
>> number of iterations is `EM::DEFAULT_MAX_ITERS=100`.

## EM::train

Estimates the Gaussian mixture parameters from a samples set.

**C++:** bool EM::**train**(InputArray **samples**, OutputArray **logLikelihoods**=noArray(), OutputArray **la-
bels**=noArray(), OutputArray **probs**=noArray())

**C++:** bool EM::**trainE**(InputArray **samples**, InputArray **means0**, InputArray **covs0**=noArray(), InputAr-
ray **weights0**=noArray(), OutputArray **logLikelihoods**=noArray(), OutputArray **la-
bels**=noArray(), OutputArray **probs**=noArray())

**C++:** bool EM::**trainM**(InputArray **samples**, InputArray **probs0**, OutputArray **logLikelihoods**=noArray(), OutputArray **labels**=noArray(), OutputArray **probs**=noArray())

**Python:** cv2.EM.**train**(samples[, logLikelihoods[, labels[, probs ] ] ]) → retval, logLikelihoods, labels, probs

**Python:** cv2.EM.**trainE**(samples, means0[, covs0[, weights0[, logLikelihoods[, labels[, probs ] ] ] ] ]) → retval, logLikelihoods, labels, probs

**Python:** cv2.EM.**trainM**(samples, probs0[, logLikelihoods[, labels[, probs ] ] ]) → retval, logLikelihoods, labels, probs

> Parameters

>> **samples** – Samples from which the Gaussian mixture model will be estimated. It should be a one-channel matrix, each row of which is a sample. If the matrix does not have CV_64F type it will be converted to the inner matrix of such type for the further computing.

>> **means0** – Initial means $a_k$ of mixture components. It is a one-channel matrix of nclusters × dims size. If the matrix does not have CV_64F type it will be converted to the inner matrix of such type for the further computing.

>> **covs0** – The vector of initial covariance matrices $S_k$ of mixture components. Each of covariance matrices is a one-channel matrix of dims × dims size. If the matrices do not have CV_64F type they will be converted to the inner matrices of such type for the further computing.

>> **weights0** – Initial weights $\pi_k$ of mixture components. It should be a one-channel floating-point matrix with 1 × nclusters or nclusters × 1 size.

>> **probs0** – Initial probabilities $p_{i,k}$ of sample $i$ to belong to mixture component $k$. It is a one-channel floating-point matrix of nsamples × nclusters size.

>> **logLikelihoods** – The optional output matrix that contains a likelihood logarithm value for each sample. It has nsamples × 1 size and CV_64FC1 type.

>> **labels** – The optional output "class label" for each sample: $labels_i = arg\ max_k(p_{i,k}), i = 1..N$ (indices of the most probable mixture component for each sample). It has nsamples × 1 size and CV_32SC1 type.

>> **probs** – The optional output matrix that contains posterior probabilities of each Gaussian mixture component given the each sample. It has nsamples × nclusters size and CV_64FC1 type.

Three versions of training method differ in the initialization of Gaussian mixture model parameters and start step:

- **train** - Starts with Expectation step. Initial values of the model parameters will be estimated by the k-means algorithm.

- **trainE** - Starts with Expectation step. You need to provide initial means $a_k$ of mixture components. Optionally you can pass initial weights $\pi_k$ and covariance matrices $S_k$ of mixture components.

- **trainM** - Starts with Maximization step. You need to provide initial probabilities $p_{i,k}$ to use this option.

The methods return true if the Gaussian mixture model was trained successfully, otherwise it returns false.

Unlike many of the ML models, EM is an unsupervised learning algorithm and it does not take responses (class labels or function values) as input. Instead, it computes the *Maximum Likelihood Estimate* of the Gaussian mixture parameters from an input sample set, stores all the parameters inside the structure: $p_{i,k}$ in probs, $a_k$ in means, $S_k$ in covs[k], $\pi_k$ in weights, and optionally computes the output "class label" for each sample: $labels_i = arg\ max_k(p_{i,k}), i = 1..N$ (indices of the most probable mixture component for each sample).

The trained model can be used further for prediction, just like any other classifier. The trained model is similar to the `CvNormalBayesClassifier`.

## EM::predict

Returns a likelihood logarithm value and an index of the most probable mixture component for the given sample.

**C++:** Vec2d EM::**predict**(InputArray **sample**, OutputArray **probs**=noArray()) const

**Python:** cv2.EM.**predict**(sample[, probs]) → retval, probs

> **Parameters**
>
> > **sample** – A sample for classification. It should be a one-channel matrix of $1 \times \text{dims}$ or $\text{dims} \times 1$ size.
> >
> > **probs** – Optional output matrix that contains posterior probabilities of each component given the sample. It has $1 \times \text{nclusters}$ size and `CV_64FC1` type.

The method returns a two-element `double` vector. Zero element is a likelihood logarithm value for the sample. First element is an index of the most probable mixture component for the given sample.

## CvEM::isTrained

Returns `true` if the Gaussian mixture model was trained.

**C++:** bool EM::**isTrained**() const

**Python:** cv2.EM.**isTrained**() → retval

## EM::read, EM::write

See `Algorithm::read()` and `Algorithm::write()`.
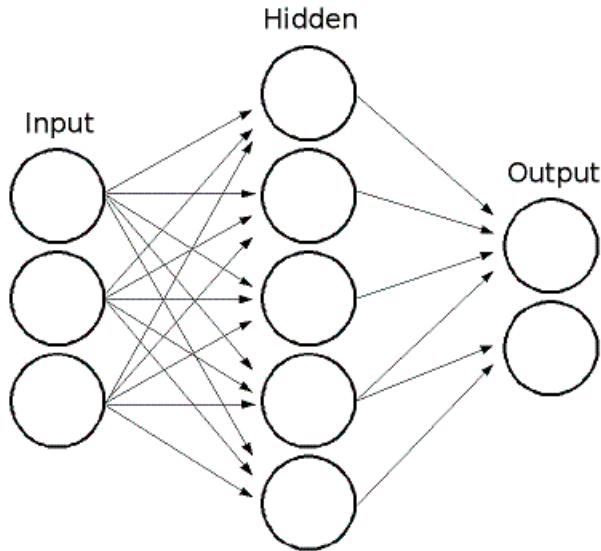
## EM::get, EM::set

See `Algorithm::get()` and `Algorithm::set()`. The following parameters are available:
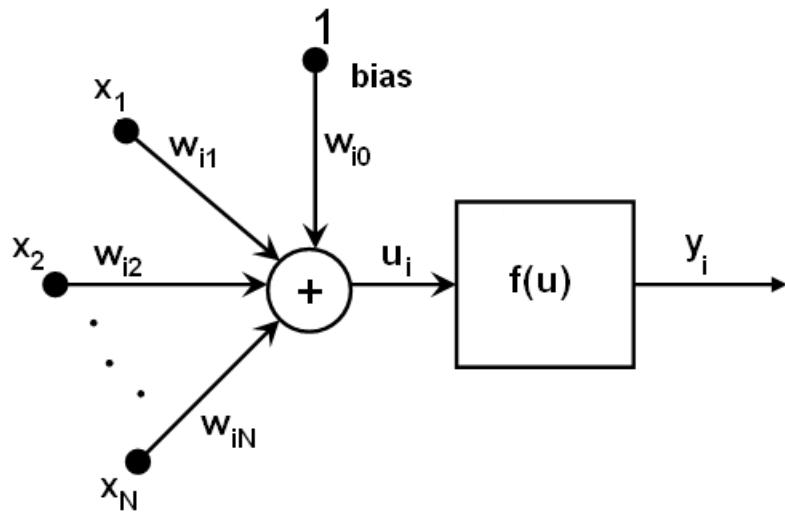
- `"nclusters"`
- `"covMatType"`
- `"maxIters"`
- `"epsilon"`
- `"weights"` *(read-only)*
- `"means"` *(read-only)*
- `"covs"` *(read-only)*

## 9.11 Neural Networks

ML implements feed-forward artificial neural networks or, more particularly, multi-layer perceptrons (MLP), the most commonly used type of neural networks. MLP consists of the input layer, output layer, and one or more hidden layers. Each layer of MLP includes one or more neurons directionally linked with the neurons from the previous and the next layer. The example below represents a 3-layer perceptron with three inputs, two outputs, and the hidden layer including five neurons:



All the neurons in MLP are similar. Each of them has several input links (it takes the output values from several neurons in the previous layer as input) and several output links (it passes the response to several neurons in the next layer). The values retrieved from the previous layer are summed up with certain weights, individual for each neuron, plus the bias term. The sum is transformed using the activation function $f$ that may be also different for different neurons.
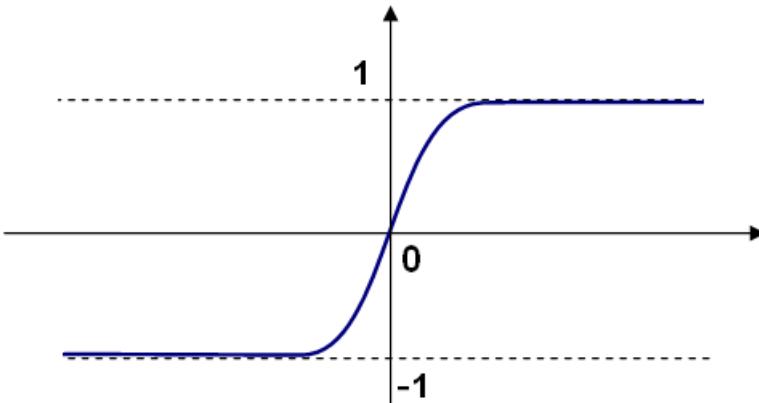


In other words, given the outputs $x_j$ of the layer $n$, the outputs $y_i$ of the layer $n + 1$ are computed as:

$$u_i = \sum_j (w_{i,j}^{n+1} * x_j) + w_{i,bias}^{n+1}$$

$$y_i = f(u_i)$$

Different activation functions may be used. ML implements three standard functions:

- Identity function ( `CvANN_MLP::IDENTITY` ): $f(x) = x$

- Symmetrical sigmoid ( `CvANN_MLP::SIGMOID_SYM` ): $f(x) = \beta * (1 - e^{-\alpha x})/(1 + e^{-\alpha x})$ ), which is the default choice for MLP. The standard sigmoid with $\beta = 1, \alpha = 1$ is shown below:



- Gaussian function ( `CvANN_MLP::GAUSSIAN` ): $f(x) = \beta e^{-\alpha x * x}$ , which is not completely supported at the moment.

In ML, all the neurons have the same activation functions, with the same free parameters ( $\alpha, \beta$ ) that are specified by user and are not altered by the training algorithms.

So, the whole trained network works as follows:

1. Take the feature vector as input. The vector size is equal to the size of the input layer.

2. Pass values as input to the first hidden layer.

3. Compute outputs of the hidden layer using the weights and the activation functions.

4. Pass outputs further downstream until you compute the output layer.

So, to compute the network, you need to know all the weights $w_{i,j}^{n+1)}$ . The weights are computed by the training algorithm. The algorithm takes a training set, multiple input vectors with the corresponding output vectors, and iteratively adjusts the weights to enable the network to give the desired response to the provided input vectors.

The larger the network size (the number of hidden layers and their sizes) is, the more the potential network flexibility is. The error on the training set could be made arbitrarily small. But at the same time the learned network also "learns" the noise present in the training set, so the error on the test set usually starts increasing after the network size reaches a limit. Besides, the larger networks are trained much longer than the smaller ones, so it is reasonable to pre-process the data, using `PCA::operator()` or similar technique, and train a smaller network on only essential features.

Another MLP feature is an inability to handle categorical data as is. However, there is a workaround. If a certain feature in the input or output (in case of $n$ -class classifier for $n > 2$ ) layer is categorical and can take $M > 2$ different values, it makes sense to represent it as a binary tuple of $M$ elements, where the $i$ -th element is 1 if and only if the feature is equal to the $i$ -th value out of $M$ possible. It increases the size of the input/output layer but speeds up the training algorithm convergence and at the same time enables "fuzzy" values of such variables, that is, a tuple of probabilities instead of a fixed value.

ML implements two algorithms for training MLP's. The first algorithm is a classical random sequential back-propagation algorithm. The second (default) one is a batch RPROP algorithm.

## CvANN_MLP_TrainParams

**struct** `CvANN_MLP_TrainParams`

> Parameters of the MLP training algorithm. You can initialize the structure by a constructor or the individual parameters can be adjusted after the structure is created.

> The back-propagation algorithm parameters:

> **double** `bp_dw_scale`

>> Strength of the weight gradient term. The recommended value is about 0.1.

> **double** `bp_moment_scale`

>> Strength of the momentum term (the difference between weights on the 2 previous iterations). This parameter provides some inertia to smooth the random fluctuations of the weights. It can vary from 0 (the feature is disabled) to 1 and beyond. The value 0.1 or so is good enough

> The RPROP algorithm parameters (see [RPROP93] for details):

> **double** `rp_dw0`

>> Initial value $\Delta_0$ of update-values $\Delta_{ij}$.

> **double** `rp_dw_plus`

>> Increase factor $\eta^+$. It must be >1.

> **double** `rp_dw_minus`

>> Decrease factor $\eta^-$. It must be <1.

> **double** `rp_dw_min`

>> Update-values lower limit $\Delta_{min}$. It must be positive.

> **double** `rp_dw_max`

>> Update-values upper limit $\Delta_{max}$. It must be >1.

## CvANN_MLP_TrainParams::CvANN_MLP_TrainParams

The constructors.

**C++:** `CvANN_MLP_TrainParams::`**`CvANN_MLP_TrainParams`**`()`

**C++:** `CvANN_MLP_TrainParams::`**`CvANN_MLP_TrainParams`**`(`CvTermCriteria **term_crit**, int **train_method**, double **param1**, double **param2**=0 )

> **Parameters**

>> **term_crit** – Termination criteria of the training algorithm. You can specify the maximum number of iterations (`max_iter`) and/or how much the error could change between the iterations to make the algorithm continue (`epsilon`).

>> **train_method** – Training method of the MLP. Possible values are:

>> – **CvANN_MLP_TrainParams::BACKPROP** The back-propagation algorithm.

>> – **CvANN_MLP_TrainParams::RPROP** The RPROP algorithm.

>> **param1** – Parameter of the training method. It is `rp_dw0` for RPROP and `bp_dw_scale` for BACKPROP.

>> **param2** – Parameter of the training method. It is `rp_dw_min` for RPROP and `bp_moment_scale` for BACKPROP.

By default the RPROP algorithm is used:

```
CvANN_MLP_TrainParams::CvANN_MLP_TrainParams()
{
    term_crit = cvTermCriteria( CV_TERMCRIT_ITER + CV_TERMCRIT_EPS, 1000, 0.01 );
    train_method = RPROP;
    bp_dw_scale = bp_moment_scale = 0.1;
    rp_dw0 = 0.1; rp_dw_plus = 1.2; rp_dw_minus = 0.5;
    rp_dw_min = FLT_EPSILON; rp_dw_max = 50.;
}
```

## CvANN_MLP

**class `CvANN_MLP` : public `CvStatModel`**

MLP model.

Unlike many other models in ML that are constructed and trained at once, in the MLP model these steps are separated. First, a network with the specified topology is created using the non-default constructor or the method `CvANN_MLP::create()`. All the weights are set to zeros. Then, the network is trained using a set of input and output vectors. The training procedure can be repeated more than once, that is, the weights can be adjusted based on the new training data.

## CvANN_MLP::CvANN_MLP

The constructors.

**C++:** `CvANN_MLP::CvANN_MLP()`

**C++:** `CvANN_MLP::CvANN_MLP`(const CvMat* **layerSizes**, int **activateFunc**=CvANN_MLP::SIGMOID_SYM, double **fparam1**=0, double **fparam2**=0 )

**Python:** `cv2.ANN_MLP`([layerSizes[, activateFunc[, fparam1[, fparam2 ] ] ] ]) → <ANN_MLP object>

The advanced constructor allows to create MLP with the specified topology. See `CvANN_MLP::create()` for details.

## CvANN_MLP::create

Constructs MLP with the specified topology.

**C++:** void `CvANN_MLP::create`(const Mat& **layerSizes**, int **activateFunc**=CvANN_MLP::SIGMOID_SYM, double **fparam1**=0, double **fparam2**=0 )

**C++:** void `CvANN_MLP::create`(const CvMat* **layerSizes**, int **activateFunc**=CvANN_MLP::SIGMOID_SYM, double **fparam1**=0, double **fparam2**=0 )

**Python:** `cv2.ANN_MLP.create`(layerSizes[, activateFunc[, fparam1[, fparam2 ] ] ]) → None

> **Parameters**
>
>> **layerSizes** – Integer vector specifying the number of neurons in each layer including the input and output layers.
>>
>> **activateFunc** – Parameter specifying the activation function for each neuron: one of `CvANN_MLP::IDENTITY`, `CvANN_MLP::SIGMOID_SYM`, and `CvANN_MLP::GAUSSIAN`.
>>
>> **fparam1** – Free parameter of the activation function, $\alpha$. See the formulas in the introduction section.

        **fparam2** – Free parameter of the activation function, β. See the formulas in the introduction section.

The method creates an MLP network with the specified topology and assigns the same activation function to all the neurons.

## CvANN_MLP::train

Trains/updates MLP.

**C++:** int CvANN_MLP::**train**(const Mat& **inputs**, const Mat& **outputs**, const Mat& **sampleWeights**, const Mat& **sampleIdx**=Mat(), CvANN_MLP_TrainParams **params**=CvANN_MLP_TrainParams(), int **flags**=0 )

**C++:** int CvANN_MLP::**train**(const CvMat* **inputs**, const CvMat* **outputs**, const CvMat* **sampleWeights**, const CvMat* **sampleIdx**=0, CvANN_MLP_TrainParams **params**=CvANN_MLP_TrainParams(), int **flags**=0 )

**Python:** cv2.ANN_MLP.**train**(inputs, outputs, sampleWeights[, sampleIdx[, params[, flags ] ] ]) → retval

    **Parameters**

        **inputs** – Floating-point matrix of input vectors, one vector per row.

        **outputs** – Floating-point matrix of the corresponding output vectors, one vector per row.

        **sampleWeights** – (RPROP only) Optional floating-point vector of weights for each sample. Some samples may be more important than others for training. You may want to raise the weight of certain classes to find the right balance between hit-rate and false-alarm rate, and so on.

        **sampleIdx** – Optional integer vector indicating the samples (rows of `inputs` and `outputs`) that are taken into account.

        **params** – Training parameters. See the `CvANN_MLP_TrainParams` description.

        **flags** – Various parameters to control the training algorithm. A combination of the following parameters is possible:

        – **UPDATE_WEIGHTS** Algorithm updates the network weights, rather than computes them from scratch. In the latter case the weights are initialized using the Nguyen-Widrow algorithm.

        – **NO_INPUT_SCALE** Algorithm does not normalize the input vectors. If this flag is not set, the training algorithm normalizes each input feature independently, shifting its mean value to 0 and making the standard deviation equal to 1. If the network is assumed to be updated frequently, the new training data could be much different from original one. In this case, you should take care of proper normalization.

        – **NO_OUTPUT_SCALE** Algorithm does not normalize the output vectors. If the flag is not set, the training algorithm normalizes each output feature independently, by transforming it to the certain range depending on the used activation function.

This method applies the specified training algorithm to computing/adjusting the network weights. It returns the number of done iterations.

The RPROP training algorithm is parallelized with the TBB library.

## CvANN_MLP::predict

Predicts responses for input samples.

**C++:** float CvANN_MLP::**predict**(const Mat& **inputs**, Mat& **outputs**) const

**C++:** float CvANN_MLP::**predict**(const CvMat* **inputs**, CvMat* **outputs**) const

**Python:** cv2.ANN_MLP.**predict**(inputs[, outputs]) → retval, outputs

>      **Parameters**

>>          **inputs** – Input samples.

>>          **outputs** – Predicted responses for corresponding samples.

The method returns a dummy value which should be ignored.

## CvANN_MLP::get_layer_count

Returns the number of layers in the MLP.

**C++:** int CvANN_MLP::**get_layer_count**()

## CvANN_MLP::get_layer_sizes

Returns numbers of neurons in each layer of the MLP.

**C++:** const CvMat* CvANN_MLP::**get_layer_sizes**()

The method returns the integer vector specifying the number of neurons in each layer including the input and output layers of the MLP.

## CvANN_MLP::get_weights

Returns neurons weights of the particular layer.

**C++:** double* CvANN_MLP::**get_weights**(int **layer**)

>      **Parameters**

>>          **layer** – Index of the particular layer.

# 9.12 MLData

For the machine learning algorithms, the data set is often stored in a file of the `.csv`-like format. The file contains a table of predictor and response values where each row of the table corresponds to a sample. Missing values are supported. The UC Irvine Machine Learning Repository (http://archive.ics.uci.edu/ml/) provides many data sets stored in such a format to the machine learning community. The class `MLData` is implemented to easily load the data for training one of the OpenCV machine learning algorithms. For float values, only the `'.'` separator is supported.

## CvMLData

**class CvMLData**

Class for loading the data from a `.csv` file.

```
class CV_EXPORTS CvMLData
{
public:
    CvMLData();
    virtual ~CvMLData();

    int read_csv(const char* filename);

    const CvMat* get_values() const;
    const CvMat* get_responses();
    const CvMat* get_missing() const;

    void set_response_idx( int idx );
    int get_response_idx() const;


    void set_train_test_split( const CvTrainTestSplit * spl);
    const CvMat* get_train_sample_idx() const;
    const CvMat* get_test_sample_idx() const;
    void mix_train_and_test_idx();

    const CvMat* get_var_idx();
    void change_var_idx( int vi, bool state );

    const CvMat* get_var_types();
    void set_var_types( const char* str );

    int get_var_type( int var_idx ) const;
    void change_var_type( int var_idx, int type);

    void set_delimiter( char ch );
    char get_delimiter() const;

    void set_miss_ch( char ch );
    char get_miss_ch() const;

    const std::map<std::string, int>& get_class_labels_map() const;

protected:
    ...
};
```

## CvMLData::read_csv

Reads the data set from a `.csv`-like `filename` file and stores all read values in a matrix.

**C++:** int CvMLData::**read_csv**(const char* **filename**)

> **Parameters**
>
> > **filename** – The input file name

While reading the data, the method tries to define the type of variables (predictors and responses): ordered or categorical. If a value of the variable is not numerical (except for the label for a missing value), the type of the variable is set to `CV_VAR_CATEGORICAL`. If all existing values of the variable are numerical, the type of the variable is set to `CV_VAR_ORDERED`. So, the default definition of variables types works correctly for all cases except the case of a categorical variable with numerical class labels. In this case, the type `CV_VAR_ORDERED` is set. You should change the type to `CV_VAR_CATEGORICAL` using the method `CvMLData::change_var_type()`. For categorical variables, a common map is built to convert a string class label to the numerical class label. Use `CvMLData::get_class_labels_map()` to obtain this map.

Also, when reading the data, the method constructs the mask of missing values. For example, values are equal to '*?*'.

## CvMLData::get_values

Returns a pointer to the matrix of predictors and response values

**C++:** const CvMat* CvMLData::**get_values**() const

The method returns a pointer to the matrix of predictor and response `values` or `0` if the data has not been loaded from the file yet.

The row count of this matrix equals the sample count. The column count equals predictors + 1 for the response (if exists) count. This means that each row of the matrix contains values of one sample predictor and response. The matrix type is `CV_32FC1`.

## CvMLData::get_responses

Returns a pointer to the matrix of response values

**C++:** const CvMat* CvMLData::**get_responses**()

The method returns a pointer to the matrix of response values or throws an exception if the data has not been loaded from the file yet.

This is a single-column matrix of the type `CV_32FC1`. Its row count is equal to the sample count, one column and .

## CvMLData::get_missing

Returns a pointer to the mask matrix of missing values

**C++:** const CvMat* CvMLData::**get_missing**() const

The method returns a pointer to the mask matrix of missing values or throws an exception if the data has not been loaded from the file yet.

This matrix has the same size as the `values` matrix (see `CvMLData::get_values()`) and the type `CV_8UC1`.

## CvMLData::set_response_idx

Specifies index of response column in the data matrix

**C++:** void CvMLData::**set_response_idx**(int **idx**)

The method sets the index of a response column in the `values` matrix (see `CvMLData::get_values()`) or throws an exception if the data has not been loaded from the file yet.

The old response columns become predictors. If `idx < 0`, there is no response.

## CvMLData::get_response_idx

Returns index of the response column in the loaded data matrix

**C++:** int CvMLData::**get_response_idx**() const

The method returns the index of a response column in the `values` matrix (see `CvMLData::get_values()`) or throws an exception if the data has not been loaded from the file yet.

If `idx < 0`, there is no response.

## CvMLData::set_train_test_split

Divides the read data set into two disjoint training and test subsets.

**C++:** void CvMLData::**set_train_test_split**(const CvTrainTestSplit* **spl**)

This method sets parameters for such a split using `spl` (see `CvTrainTestSplit`) or throws an exception if the data has not been loaded from the file yet.

## CvMLData::get_train_sample_idx

Returns the matrix of sample indices for a training subset

**C++:** const CvMat* CvMLData::**get_train_sample_idx**() const

The method returns the matrix of sample indices for a training subset. This is a single-row matrix of the type `CV_32SC1`. If data split is not set, the method returns 0. If the data has not been loaded from the file yet, an exception is thrown.

## CvMLData::get_test_sample_idx

Returns the matrix of sample indices for a testing subset

**C++:** const CvMat* CvMLData::**get_test_sample_idx**() const

## CvMLData::mix_train_and_test_idx

Mixes the indices of training and test samples

**C++:** void CvMLData::**mix_train_and_test_idx**()

The method shuffles the indices of training and test samples preserving sizes of training and test subsets if the data split is set by `CvMLData::get_values()`. If the data has not been loaded from the file yet, an exception is thrown.

## CvMLData::get_var_idx

Returns the indices of the active variables in the data matrix

**C++:** const CvMat* CvMLData::**get_var_idx**()

The method returns the indices of variables (columns) used in the `values` matrix (see `CvMLData::get_values()`).

It returns 0 if the used subset is not set. It throws an exception if the data has not been loaded from the file yet. Returned matrix is a single-row matrix of the type `CV_32SC1`. Its column count is equal to the size of the used variable subset.

## CvMLData::change_var_idx

Enables or disables particular variable in the loaded data

**C++:** void CvMLData::**change_var_idx**(int **vi**, bool **state**)

By default, after reading the data set all variables in the values matrix (see `CvMLData::get_values()`) are used. But you may want to use only a subset of variables and include/exclude (depending on state value) a variable with the vi index from the used subset. If the data has not been loaded from the file yet, an exception is thrown.

## CvMLData::get_var_types

Returns a matrix of the variable types.

**C++:** const CvMat* CvMLData::**get_var_types**()

The function returns a single-row matrix of the type `CV_8UC1`, where each element is set to either `CV_VAR_ORDERED` or `CV_VAR_CATEGORICAL`. The number of columns is equal to the number of variables. If data has not been loaded from file yet an exception is thrown.

## CvMLData::set_var_types

Sets the variables types in the loaded data.

**C++:** void CvMLData::**set_var_types**(const char* **str**)

In the string, a variable type is followed by a list of variables indices. For example: `"ord[0-17],cat[18]"`, `"ord[0,2,4,10-12], cat[1,3,5-9,13,14]"`, `"cat"` (all variables are categorical), `"ord"` (all variables are ordered).

## CvMLData::get_var_type

Returns type of the specified variable

**C++:** int CvMLData::**get_var_type**(int **var_idx**) const

The method returns the type of a variable by the index var_idx ( `CV_VAR_ORDERED` or `CV_VAR_CATEGORICAL`).

## CvMLData::change_var_type

Changes type of the specified variable

**C++:** void CvMLData::**change_var_type**(int **var_idx**, int **type**)

The method changes type of variable with index var_idx from existing type to type ( `CV_VAR_ORDERED` or `CV_VAR_CATEGORICAL`).

## CvMLData::set_delimiter

Sets the delimiter in the file used to separate input numbers

**C++:** void CvMLData::**set_delimiter**(char **ch**)

The method sets the delimiter for variables in a file. For example: `','` (default), `';'`, `' '` (space), or other characters. The floating-point separator `'.'` is not allowed.

## CvMLData::get_delimiter

Returns the currently used delimiter character.

**C++:** char CvMLData::**get_delimiter**() const

## CvMLData::set_miss_ch

Sets the character used to specify missing values

**C++:** void CvMLData::**set_miss_ch**(char **ch**)

The method sets the character used to specify missing values. For example: '?' (default), '-'. The floating-point separator '.' is not allowed.

## CvMLData::get_miss_ch

Returns the currently used missing value character.

**C++:** char CvMLData::**get_miss_ch**() const

## CvMLData::get_class_labels_map

Returns a map that converts strings to labels.

**C++:** const std::map<std::string, int>& CvMLData::**get_class_labels_map**() const

The method returns a map that converts string class labels to the numerical class labels. It can be used to get an original class label as in a file.

## CvTrainTestSplit

**struct `CvTrainTestSplit`**

Structure setting the split of a data set read by CvMLData.

```
struct CvTrainTestSplit
{
    CvTrainTestSplit();
    CvTrainTestSplit( int train_sample_count, bool mix = true);
    CvTrainTestSplit( float train_sample_portion, bool mix = true);

    union
    {
        int count;
        float portion;
    } train_sample_part;
    int train_sample_part_mode;

    bool mix;
};
```

There are two ways to construct a split:

- Set the training sample count (subset size) `train_sample_count`. Other existing samples are located in a test subset.

---

• Set a training sample portion in `[0,..1]`. The flag `mix` is used to mix training and test samples indices when the split is set. Otherwise, the data set is split in the storing order: the first part of samples of a given size is a training subset, the second part is a test subset.

# FLANN. CLUSTERING AND SEARCH IN MULTI-DIMENSIONAL SPACES

## 10.1 Fast Approximate Nearest Neighbor Search

This section documents OpenCV's interface to the FLANN library. FLANN (Fast Library for Approximate Nearest Neighbors) is a library that contains a collection of algorithms optimized for fast nearest neighbor search in large datasets and for high dimensional features. More information about FLANN can be found in [Muja2009] .

### flann::Index_

**class** `flann::`**`Index_`**

The FLANN nearest neighbor index class. This class is templated with the type of elements for which the index is built.

### flann::Index_<T>::Index_

Constructs a nearest neighbor search index for a given dataset.

**C++:** `flann::Index_<T>::`**`Index_`**`(const Mat&` **features**`, const IndexParams&` **params**`)`

> **Parameters**
>
>> **features** – Matrix of containing the features(points) to index. The size of the matrix is `num_features x feature_dimensionality` and the data type of the elements in the matrix must coincide with the type of the index.
>>
>> **params** – Structure containing the index parameters. The type of index that will be constructed depends on the type of this parameter. See the description.

The method constructs a fast search structure from a set of features using the specified algorithm with specified parameters, as defined by `params`. `params` is a reference to one of the following class `IndexParams` descendants:

- **LinearIndexParams** When passing an object of this type, the index will perform a linear, brute-force search.

  ```
  struct LinearIndexParams : public IndexParams
  {
  };
  ```

- **KDTreeIndexParams** When passing an object of this type the index constructed will consist of a set of randomized kd-trees which will be searched in parallel.

```
struct KDTreeIndexParams : public IndexParams
{
    KDTreeIndexParams( int trees = 4 );
};
```

  – **trees** The number of parallel kd-trees to use. Good values are in the range [1..16]

• **KMeansIndexParams** When passing an object of this type the index constructed will be a hierarchical k-means tree.

```
struct KMeansIndexParams : public IndexParams
{
    KMeansIndexParams(
        int branching = 32,
        int iterations = 11,
        flann_centers_init_t centers_init = CENTERS_RANDOM,
        float cb_index = 0.2 );
};
```

  – **branching** The branching factor to use for the hierarchical k-means tree

  – **iterations** The maximum number of iterations to use in the k-means clustering stage when building the k-means tree. A value of -1 used here means that the k-means clustering should be iterated until convergence

  – **centers_init** The algorithm to use for selecting the initial centers when performing a k-means clustering step. The possible values are CENTERS_RANDOM (picks the initial cluster centers randomly), CENTERS_GONZALES (picks the initial centers using Gonzales' algorithm) and CENTERS_KMEANSPP (picks the initial centers using the algorithm suggested in arthur_kmeanspp_2007 )

  – **cb_index** This parameter (cluster boundary index) influences the way exploration is performed in the hierarchical kmeans tree. When cb_index is zero the next kmeans domain to be explored is chosen to be the one with the closest center. A value greater then zero also takes into account the size of the domain.

• **CompositeIndexParams** When using a parameters object of this type the index created combines the randomized kd-trees and the hierarchical k-means tree.

```
struct CompositeIndexParams : public IndexParams
{
    CompositeIndexParams(
        int trees = 4,
        int branching = 32,
        int iterations = 11,
        flann_centers_init_t centers_init = CENTERS_RANDOM,
        float cb_index = 0.2 );
};
```

• **LshIndexParams** When using a parameters object of this type the index created uses multi-probe LSH (by Multi-Probe LSH: Efficient Indexing for High-Dimensional Similarity Search by Qin Lv, William Josephson, Zhe Wang, Moses Charikar, Kai Li., Proceedings of the 33rd International Conference on Very Large Data Bases (VLDB). Vienna, Austria. September 2007)

```
struct LshIndexParams : public IndexParams
{
    LshIndexParams(
        unsigned int table_number,
        unsigned int key_size,
        unsigned int multi_probe_level );
};
```

  – **table_number** the number of hash tables to use (between 10 and 30 usually).

- **key_size** the size of the hash key in bits (between 10 and 20 usually).

- **multi_probe_level** the number of bits to shift to check for neighboring buckets (0 is regular LSH, 2 is recommended).

- **AutotunedIndexParams** When passing an object of this type the index created is automatically tuned to offer the best performance, by choosing the optimal index type (randomized kd-trees, hierarchical kmeans, linear) and parameters for the dataset provided.

```
struct AutotunedIndexParams : public IndexParams
{
    AutotunedIndexParams(
        float target_precision = 0.9,
        float build_weight = 0.01,
        float memory_weight = 0,
        float sample_fraction = 0.1 );
};
```

- **target_precision** Is a number between 0 and 1 specifying the percentage of the approximate nearest-neighbor searches that return the exact nearest-neighbor. Using a higher value for this parameter gives more accurate results, but the search takes longer. The optimum value usually depends on the application.

- **build_weight** Specifies the importance of the index build time raported to the nearest-neighbor search time. In some applications it's acceptable for the index build step to take a long time if the subsequent searches in the index can be performed very fast. In other applications it's required that the index be build as fast as possible even if that leads to slightly longer search times.

- **memory_weight** Is used to specify the tradeoff between time (index build time and search time) and memory used by the index. A value less than 1 gives more importance to the time spent and a value greater than 1 gives more importance to the memory usage.

- **sample_fraction** Is a number between 0 and 1 indicating what fraction of the dataset to use in the automatic parameter configuration algorithm. Running the algorithm on the full dataset gives the most accurate results, but for very large datasets can take longer than desired. In such case using just a fraction of the data helps speeding up this algorithm while still giving good approximations of the optimum parameters.

- **SavedIndexParams** This object type is used for loading a previously saved index from the disk.

```
struct SavedIndexParams : public IndexParams
{
    SavedIndexParams( std::string filename );
};
```

- **filename** The filename in which the index was saved.

## flann::Index_<T>::knnSearch

Performs a K-nearest neighbor search for a given query point using the index.

**C++:** void flann::Index_<T>::**knnSearch**(const vector<T>& **query**, vector<int>& **indices**, vector<float>& **dists**, int **knn**, const SearchParams& **params**)

**C++:** void flann::Index_<T>::**knnSearch**(const Mat& **queries**, Mat& **indices**, Mat& **dists**, int **knn**, const SearchParams& **params**)

   **Parameters**

   **query** – The query point

   **indices** – Vector that will contain the indices of the K-nearest neighbors found. It must have at least knn size.

> **dists** – Vector that will contain the distances to the K-nearest neighbors found. It must have at least knn size.
>
> **knn** – Number of nearest neighbors to search for.
>
> **params** – Search parameters

```
struct SearchParams {
        SearchParams(int checks = 32);
};
```

> – **checks** The number of times the tree(s) in the index should be recursively traversed. A higher value for this parameter would give better search precision, but also take more time. If automatic configuration was used when the index was created, the number of checks required to achieve the specified precision was also computed, in which case this parameter is ignored.

## flann::Index_<T>::radiusSearch

Performs a radius nearest neighbor search for a given query point.

**C++:** int flann::Index_<T>::**radiusSearch**(const vector<T>& **query**, vector<int>& **indices**, vector<float>& **dists**, float **radius**, const SearchParams& **params**)

**C++:** int flann::Index_<T>::**radiusSearch**(const Mat& **query**, Mat& **indices**, Mat& **dists**, float **radius**, const SearchParams& **params**)

> **Parameters**
>
>> **query** – The query point
>>
>> **indices** – Vector that will contain the indices of the points found within the search radius in decreasing order of the distance to the query point. If the number of neighbors in the search radius is bigger than the size of this vector, the ones that don't fit in the vector are ignored.
>>
>> **dists** – Vector that will contain the distances to the points found within the search radius
>>
>> **radius** – The search radius
>>
>> **params** – Search parameters

## flann::Index_<T>::save

Saves the index to a file.

**C++:** void flann::Index_<T>::**save**(std::string **filename**)

> **Parameters**
>
>> **filename** – The file to save the index to

## flann::Index_<T>::getIndexParameters

Returns the index parameters.

**C++:** const IndexParams* flann::Index_<T>::**getIndexParameters**()

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The method is useful in the case of auto-tuned indices, when the parameters are chosen during the index construction. Then, the method can be used to retrieve the actual parameter values.

## 10.2 Clustering

### flann::hierarchicalClustering<Distance>

Clusters features using hierarchical k-means algorithm.

**C++:** template<typename Distance> int flann::**hierarchicalClustering**(const Mat& **features**, Mat& **centers**, const cvflann::KMeansIndexParams& **params**, Distance **d**=Distance())

> **Parameters**
>
> > **features** – The points to be clustered. The matrix must have elements of type `Distance::ElementType`.
> >
> > **centers** – The centers of the clusters obtained. The matrix must have type `Distance::ResultType`. The number of rows in this matrix represents the number of clusters desired, however, because of the way the cut in the hierarchical tree is chosen, the number of clusters computed will be the highest number of the form `(branching-1)*k+1` that's lower than the number of clusters desired, where `branching` is the tree's branching factor (see description of the KMeansIndexParams).
> >
> > **params** – Parameters used in the construction of the hierarchical k-means tree.
> >
> > **d** – Distance to be used for clustering.

The method clusters the given feature vectors by constructing a hierarchical k-means tree and choosing a cut in the tree that minimizes the cluster's variance. It returns the number of clusters found.

# GPU. GPU-ACCELERATED COMPUTER VISION

## 11.1 GPU Module Introduction

### General Information

The OpenCV GPU module is a set of classes and functions to utilize GPU computational capabilities. It is implemented using NVIDIA* CUDA* Runtime API and supports only NVIDIA GPUs. The OpenCV GPU module includes utility functions, low-level vision primitives, and high-level algorithms. The utility functions and low-level primitives provide a powerful infrastructure for developing fast vision algorithms taking advantage of GPU whereas the high-level functionality includes some state-of-the-art algorithms (such as stereo correspondence, face and people detectors, and others) ready to be used by the application developers.

The GPU module is designed as a host-level API. This means that if you have pre-compiled OpenCV GPU binaries, you are not required to have the CUDA Toolkit installed or write any extra code to make use of the GPU.

The OpenCV GPU module is designed for ease of use and does not require any knowledge of CUDA. Though, such a knowledge will certainly be useful to handle non-trivial cases or achieve the highest performance. It is helpful to understand the cost of various operations, what the GPU does, what the preferred data formats are, and so on. The GPU module is an effective instrument for quick implementation of GPU-accelerated computer vision algorithms. However, if your algorithm involves many simple operations, then, for the best possible performance, you may still need to write your own kernels to avoid extra write and read operations on the intermediate results.

To enable CUDA support, configure OpenCV using `CMake` with `WITH_CUDA=ON` . When the flag is set and if CUDA is installed, the full-featured OpenCV GPU module is built. Otherwise, the module is still built but at runtime all functions from the module throw `Exception` with `CV_GpuNotSupported` error code, except for `gpu::getCudaEnabledDeviceCount()`. The latter function returns zero GPU count in this case. Building OpenCV without CUDA support does not perform device code compilation, so it does not require the CUDA Toolkit installed. Therefore, using the `gpu::getCudaEnabledDeviceCount()` function, you can implement a high-level algorithm that will detect GPU presence at runtime and choose an appropriate implementation (CPU or GPU) accordingly.

### Compilation for Different NVIDIA* Platforms

NVIDIA* compiler enables generating binary code (cubin and fatbin) and intermediate code (PTX). Binary code often implies a specific GPU architecture and generation, so the compatibility with other GPUs is not guaranteed. PTX is targeted for a virtual platform that is defined entirely by the set of capabilities or features. Depending on the selected virtual platform, some of the instructions are emulated or disabled, even if the real hardware supports all the features.

At the first call, the PTX code is compiled to binary code for the particular GPU using a JIT compiler. When the target GPU has a compute capability (CC) lower than the PTX code, JIT fails. By default, the OpenCV GPU module includes:

- Binaries for compute capabilities 1.3 and 2.0 (controlled by `CUDA_ARCH_BIN` in `CMake`)

- PTX code for compute capabilities 1.1 and 1.3 (controlled by `CUDA_ARCH_PTX` in `CMake`)

This means that for devices with CC 1.3 and 2.0 binary images are ready to run. For all newer platforms, the PTX code for 1.3 is JIT'ed to a binary image. For devices with CC 1.1 and 1.2, the PTX for 1.1 is JIT'ed. For devices with CC 1.0, no code is available and the functions throw `Exception`. For platforms where JIT compilation is performed first, the run is slow.

On a GPU with CC 1.0, you can still compile the GPU module and most of the functions will run flawlessly. To achieve this, add "1.0" to the list of binaries, for example, `CUDA_ARCH_BIN="1.0 1.3 2.0"`. The functions that cannot be run on CC 1.0 GPUs throw an exception.

You can always determine at runtime whether the OpenCV GPU-built binaries (or PTX code) are compatible with your GPU. The function `gpu::DeviceInfo::isCompatible()` returns the compatibility status (true/false).

### Utilizing Multiple GPUs

In the current version, each of the OpenCV GPU algorithms can use only a single GPU. So, to utilize multiple GPUs, you have to manually distribute the work between GPUs. Switching active devie can be done using `gpu::setDevice()` function. For more details please read Cuda C Programing Guide.

While developing algorithms for multiple GPUs, note a data passing overhead. For primitive functions and small images, it can be significant, which may eliminate all the advantages of having multiple GPUs. But for high-level algorithms, consider using multi-GPU acceleration. For example, the Stereo Block Matching algorithm has been successfully parallelized using the following algorithm:

1. Split each image of the stereo pair into two horizontal overlapping stripes.

2. Process each pair of stripes (from the left and right images) on a separate Fermi* GPU.

3. Merge the results into a single disparity map.

With this algorithm, a dual GPU gave a 180 % performance increase comparing to the single Fermi GPU. For a source code example, see http://code.opencv.org/projects/opencv/repository/revisions/master/entry/samples/gpu/.

## 11.2 Initalization and Information

### gpu::getCudaEnabledDeviceCount

Returns the number of installed CUDA-enabled devices.

**C++:** int gpu::**getCudaEnabledDeviceCount**()

Use this function before any other GPU functions calls. If OpenCV is compiled without GPU support, this function returns 0.

### gpu::setDevice

Sets a device and initializes it for the current thread.

**C++:** void gpu::**setDevice**(int **device**)

> **Parameters**
>
> > **device** – System index of a GPU device starting with 0.

If the call of this function is omitted, a default device is initialized at the fist GPU usage.

## gpu::getDevice

Returns the current device index set by `gpu::setDevice()` or initialized by default.

**C++:** int gpu::**getDevice**()

## gpu::resetDevice

Explicitly destroys and cleans up all resources associated with the current device in the current process.

**C++:** void gpu::**resetDevice**()

Any subsequent API call to this device will reinitialize the device.

## gpu::FeatureSet

Class providing GPU computing features.

```
enum FeatureSet
{
    FEATURE_SET_COMPUTE_10,
    FEATURE_SET_COMPUTE_11,
    FEATURE_SET_COMPUTE_12,
    FEATURE_SET_COMPUTE_13,
    FEATURE_SET_COMPUTE_20,
    FEATURE_SET_COMPUTE_21,
    GLOBAL_ATOMICS,
    SHARED_ATOMICS,
    NATIVE_DOUBLE
};
```

## gpu::TargetArchs

**class** gpu::**TargetArchs**

Class providing a set of static methods to check what NVIDIA* card architecture the GPU module was built for.

The following method checks whether the module was built with the support of the given feature:

> **C++:** **static** bool gpu::TargetArchs::**builtWith**(FeatureSet **feature_set**)
>
> > **Parameters**
> >
> > > **feature_set** – Features to be checked. See `gpu::FeatureSet`.

There is a set of methods to check whether the module contains intermediate (PTX) or binary GPU code for the given architecture(s):

> **C++:** **static** bool gpu::TargetArchs::**has**(int **major**, int **minor**)

> **C++:** **static** bool gpu::TargetArchs::**hasPtx**(int **major**, int **minor**)

**C++:** **static** bool gpu::TargetArchs::**hasBin**(int **major**, int **minor**)

**C++:** **static** bool gpu::TargetArchs::**hasEqualOrLessPtx**(int **major**, int **minor**)

**C++:** **static** bool gpu::TargetArchs::**hasEqualOrGreater**(int **major**, int **minor**)

**C++:** **static** bool gpu::TargetArchs::**hasEqualOrGreaterPtx**(int **major**, int **minor**)

**C++:** **static** bool gpu::TargetArchs::**hasEqualOrGreaterBin**(int **major**, int **minor**)

> **Parameters**
>
>> **major** – Major compute capability version.
>>
>> **minor** – Minor compute capability version.

According to the CUDA C Programming Guide Version 3.2: "PTX code produced for some specific compute capability can always be compiled to binary code of greater or equal compute capability".

## gpu::DeviceInfo

**class** gpu::**DeviceInfo**

Class providing functionality for querying the specified GPU properties.

```cpp
class CV_EXPORTS DeviceInfo
{
public:
    DeviceInfo();
    DeviceInfo(int device_id);

    string name() const;

    int majorVersion() const;
    int minorVersion() const;

    int multiProcessorCount() const;

    size_t freeMemory() const;
    size_t totalMemory() const;

    bool supports(FeatureSet feature) const;
    bool isCompatible() const;

    int deviceID() const;
};
```

## gpu::DeviceInfo::DeviceInfo

The constructors.

**C++:** gpu::DeviceInfo::**DeviceInfo**()

**C++:** gpu::DeviceInfo::**DeviceInfo**(int **device_id**)

> **Parameters**
>
>> **device_id** – System index of the GPU device starting with 0.

Constructs the DeviceInfo object for the specified device. If device_id parameter is missed, it constructs an object for the current device.

## gpu::DeviceInfo::name

Returns the device name.

**C++:** string gpu::DeviceInfo::**name**() const

## gpu::DeviceInfo::majorVersion

Returns the major compute capability version.

**C++:** int gpu::DeviceInfo::**majorVersion**()

## gpu::DeviceInfo::minorVersion

Returns the minor compute capability version.

**C++:** int gpu::DeviceInfo::**minorVersion**()

## gpu::DeviceInfo::multiProcessorCount

Returns the number of streaming multiprocessors.

**C++:** int gpu::DeviceInfo::**multiProcessorCount**()

## gpu::DeviceInfo::freeMemory

Returns the amount of free memory in bytes.

**C++:** size_t gpu::DeviceInfo::**freeMemory**()

## gpu::DeviceInfo::totalMemory

Returns the amount of total memory in bytes.

**C++:** size_t gpu::DeviceInfo::**totalMemory**()

## gpu::DeviceInfo::supports

Provides information on GPU feature support.

**C++:** bool gpu::DeviceInfo::**supports**(FeatureSet **feature_set**) const

> **Parameters**
>
> > **feature_set** – Features to be checked. See gpu::FeatureSet.

This function returns true if the device has the specified GPU feature. Otherwise, it returns false .

### gpu::DeviceInfo::isCompatible

Checks the GPU module and device compatibility.

**C++:** bool gpu::DeviceInfo::**isCompatible**()

This function returns `true` if the GPU module can be run on the specified device. Otherwise, it returns `false` .

### gpu::DeviceInfo::deviceID

Returns system index of the GPU device starting with 0.

**C++:** int gpu::DeviceInfo::**deviceID**()

# 11.3 Data Structures

### gpu::PtrStepSz

**class** gpu::**PtrStepSz**

Lightweight class encapsulating pitched memory on a GPU and passed to nvcc-compiled code (CUDA kernels). Typically, it is used internally by OpenCV and by users who write device code. You can call its members from both host and device code.

```cpp
template <typename T> struct PtrStepSz
{
    int cols;
    int rows;
    T* data;
    size_t step;

    PtrStepSz() : cols(0), rows(0), data(0), step(0){};
    PtrStepSz(int rows, int cols, T *data, size_t step);

    template <typename U>
    explicit PtrStepSz(const PtrStepSz<U>& d);

    typedef T elem_type;
    enum { elem_size = sizeof(elem_type) };

    __CV_GPU_HOST_DEVICE__ size_t elemSize() const;

    /* returns pointer to the beginning of the given image row */
    __CV_GPU_HOST_DEVICE__ T* ptr(int y = 0);
    __CV_GPU_HOST_DEVICE__ const T* ptr(int y = 0) const;
};

typedef PtrStepSz<unsigned char> PtrStepSzb;
typedef PtrStepSz<float> PtrStepSzf;
typedef PtrStepSz<int> PtrStepSzi;
```

### gpu::PtrStep

**class** gpu::**PtrStep**

Structure similar to `gpu::PtrStepSz` but containing only a pointer and row step. Width and height fields are excluded due to performance reasons. The structure is intended for internal use or for users who write device code.

```cpp
template<typename T> struct PtrStep
{
        T* data;
        size_t step;

        PtrStep();
        PtrStep(const PtrStepSz<T>& mem);

        typedef T elem_type;
        enum { elem_size = sizeof(elem_type) };

        __CV_GPU_HOST_DEVICE__ size_t elemSize() const;
        __CV_GPU_HOST_DEVICE__ T* ptr(int y = 0);
        __CV_GPU_HOST_DEVICE__ const T* ptr(int y = 0) const;
};

typedef PtrStep<unsigned char> PtrStep;
typedef PtrStep<float> PtrStepf;
typedef PtrStep<int> PtrStepi;
```

## gpu::GpuMat

**class** `gpu::`**`GpuMat`**

Base storage class for GPU memory with reference counting. Its interface matches the `Mat` interface with the following limitations:

- no arbitrary dimensions support (only 2D)

- no functions that return references to their data (because references on GPU are not valid for CPU)

- no expression templates technique support

Beware that the latter limitation may lead to overloaded matrix operators that cause memory allocations. The `GpuMat` class is convertible to `gpu::PtrStepSz` and `gpu::PtrStep` so it can be passed directly to the kernel.

---

**Note:** In contrast with `Mat`, in most cases `GpuMat::isContinuous() == false`. This means that rows are aligned to a size depending on the hardware. Single-row `GpuMat` is always a continuous matrix.

---

```cpp
class CV_EXPORTS GpuMat
{
public:
        //! default constructor
        GpuMat();

        GpuMat(int rows, int cols, int type);
        GpuMat(Size size, int type);


        .....

        //! builds GpuMat from Mat. Blocks uploading to device.
        explicit GpuMat (const Mat& m);

        //! returns lightweight PtrStepSz structure for passing
```

```
        //to nvcc-compiled code. Contains size, data ptr and step.
        template <class T> operator PtrStepSz<T>() const;
        template <class T> operator PtrStep<T>() const;

        //! blocks uploading data to GpuMat.
        void upload(const cv::Mat& m);
        void upload(const CudaMem& m, Stream& stream);

        //! downloads data from device to host memory. Blocking calls.
        void download(cv::Mat& m) const;

        //! download async
        void download(CudaMem& m, Stream& stream) const;
};
```

**Note:** You are not recommended to leave static or global `GpuMat` variables allocated, that is, to rely on its destructor. The destruction order of such variables and CUDA context is undefined. GPU memory release function returns error if the CUDA context has been destroyed before.

**See Also:**

`Mat`

## gpu::createContinuous

Creates a continuous matrix in the GPU memory.

**C++:** void gpu::**createContinuous**(int **rows**, int **cols**, int **type**, GpuMat& **m**)

**C++:** GpuMat gpu::**createContinuous**(int **rows**, int **cols**, int **type**)

**C++:** void gpu::**createContinuous**(Size **size**, int **type**, GpuMat& **m**)

**C++:** GpuMat gpu::**createContinuous**(Size **size**, int **type**)

> **Parameters**
>
> > **rows** – Row count.
> >
> > **cols** – Column count.
> >
> > **type** – Type of the matrix.
> >
> > **m** – Destination matrix. This parameter changes only if it has a proper type and area ( rows $\times$ cols ).

Matrix is called continuous if its elements are stored continuously, that is, without gaps at the end of each row.

## gpu::ensureSizeIsEnough

Ensures that the size of a matrix is big enough and the matrix has a proper type.

**C++:** void gpu::**ensureSizeIsEnough**(int **rows**, int **cols**, int **type**, GpuMat& **m**)

**C++:** void gpu::**ensureSizeIsEnough**(Size **size**, int **type**, GpuMat& **m**)

> **Parameters**
>
> > **rows** – Minimum desired number of rows.

> > > **cols** – Minimum desired number of columns.
> > >
> > > **size** – Rows and columns passed as a structure.
> > >
> > > **type** – Desired matrix type.
> > >
> > > **m** – Destination matrix.

The function does not reallocate memory if the matrix has proper attributes already.

## gpu::registerPageLocked

Page-locks the memory of matrix and maps it for the device(s).

**C++:** void gpu::`registerPageLocked`(Mat& **m**)

> > **Parameters**
> >
> > > **m** – Input matrix.

## gpu::unregisterPageLocked

Unmaps the memory of matrix and makes it pageable again.

**C++:** void gpu::`unregisterPageLocked`(Mat& **m**)

> > **Parameters**
> >
> > > **m** – Input matrix.

## gpu::CudaMem

**class** gpu::`CudaMem`

Class with reference counting wrapping special memory type allocation functions from CUDA. Its interface is also `Mat()`-like but with additional memory type parameters.

- **ALLOC_PAGE_LOCKED** sets a page locked memory type used commonly for fast and asynchronous up-loading/downloading data from/to GPU.

- **ALLOC_ZEROCOPY** specifies a zero copy memory allocation that enables mapping the host memory to GPU address space, if supported.

- **ALLOC_WRITE_COMBINED** sets the write combined buffer that is not cached by CPU. Such buffers are used to supply GPU with data when GPU only reads it. The advantage is a better CPU cache utilization.

---

**Note:** Allocation size of such memory types is usually limited. For more details, see *CUDA 2.2 Pinned Memory APIs* document or *CUDA C Programming Guide*.

---

```
class CV_EXPORTS CudaMem
{
public:
        enum  { ALLOC_PAGE_LOCKED = 1, ALLOC_ZEROCOPY = 2,
                ALLOC_WRITE_COMBINED = 4 };

        CudaMem(Size size, int type, int alloc_type = ALLOC_PAGE_LOCKED);

        //! creates from cv::Mat with coping data
```

```
        explicit CudaMem(const Mat& m, int alloc_type = ALLOC_PAGE_LOCKED);

         ......

        void create(Size size, int type, int alloc_type = ALLOC_PAGE_LOCKED);

        //! returns matrix header with disabled ref. counting for CudaMem data.
        Mat createMatHeader() const;
        operator Mat() const;

        //! maps host memory into device address space
        GpuMat createGpuMatHeader() const;
        operator GpuMat() const;

        //if host memory can be mapped to gpu address space;
        static bool canMapHostMemory();

        int alloc_type;
};
```

## gpu::CudaMem::createMatHeader

Creates a header without reference counting to gpu::CudaMem data.

**C++:** Mat gpu::CudaMem::**createMatHeader**() const

## gpu::CudaMem::createGpuMatHeader

Maps CPU memory to GPU address space and creates the gpu::GpuMat header without reference counting for it.

**C++:** GpuMat gpu::CudaMem::**createGpuMatHeader**() const

This can be done only if memory was allocated with the ALLOC_ZEROCOPY flag and if it is supported by the hardware. Laptops often share video and CPU memory, so address spaces can be mapped, which eliminates an extra copy.

## gpu::CudaMem::canMapHostMemory

Returns true if the current hardware supports address space mapping and ALLOC_ZEROCOPY memory allocation.

**C++:** **static** bool gpu::CudaMem::**canMapHostMemory**()

## gpu::Stream

**class** gpu::**Stream**

This class encapsulates a queue of asynchronous calls. Some functions have overloads with the additional gpu::Stream parameter. The overloads do initialization work (allocate output buffers, upload constants, and so on), start the GPU kernel, and return before results are ready. You can check whether all operations are complete via gpu::Stream::queryIfComplete(). You can asynchronously upload/download data from/to page-locked buffers, using the gpu::CudaMem or Mat header that points to a region of gpu::CudaMem.

**Note:** Currently, you may face problems if an operation is enqueued twice with different data. Some functions use the constant GPU memory, and next call may update the memory before the previous one has been finished.

But calling different operations asynchronously is safe because each operation has its own constant buffer. Memory copy/upload/download/set operations to the buffers you hold are also safe.

```cpp
class CV_EXPORTS Stream
{
public:
        Stream();
        ~Stream();

        Stream(const Stream&);
        Stream& operator=(const Stream&);

        bool queryIfComplete();
        void waitForCompletion();

        //! downloads asynchronously.
        // Warning! cv::Mat must point to page locked memory
                 (i.e. to CudaMem data or to its subMat)
        void enqueueDownload(const GpuMat& src, CudaMem& dst);
        void enqueueDownload(const GpuMat& src, Mat& dst);

        //! uploads asynchronously.
        // Warning! cv::Mat must point to page locked memory
                 (i.e. to CudaMem data or to its ROI)
        void enqueueUpload(const CudaMem& src, GpuMat& dst);
        void enqueueUpload(const Mat& src, GpuMat& dst);

        void enqueueCopy(const GpuMat& src, GpuMat& dst);

        void enqueueMemSet(const GpuMat& src, Scalar val);
        void enqueueMemSet(const GpuMat& src, Scalar val, const GpuMat& mask);

        // converts matrix type, ex from float to uchar depending on type
        void enqueueConvert(const GpuMat& src, GpuMat& dst, int type,
                double a = 1, double b = 0);
};
```

## gpu::Stream::queryIfComplete

Returns `true` if the current stream queue is finished. Otherwise, it returns false.

**C++:** bool gpu::Stream::**queryIfComplete**()

## gpu::Stream::waitForCompletion

Blocks the current CPU thread until all operations in the stream are complete.

**C++:** void gpu::Stream::**waitForCompletion**()

## gpu::StreamAccessor

**struct** gpu::**StreamAccessor**

Class that enables getting cudaStream_t from gpu::Stream and is declared in stream_accessor.hpp because it is the only public header that depends on the CUDA Runtime API. Including it brings a dependency to your code.

```
struct StreamAccessor
{
    CV_EXPORTS static cudaStream_t getStream(const Stream& stream);
};
```

# 11.4 Operations on Matrices

## gpu::gemm

Performs generalized matrix multiplication.

**C++:** void gpu::**gemm**(const GpuMat& **src1**, const GpuMat& **src2**, double **alpha**, const GpuMat& **src3**, double **beta**, GpuMat& **dst**, int **flags**=0, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src1** – First multiplied input matrix that should have `CV_32FC1` , `CV_64FC1` , `CV_32FC2` , or `CV_64FC2` type.
> >
> > **src2** – Second multiplied input matrix of the same type as `src1` .
> >
> > **alpha** – Weight of the matrix product.
> >
> > **src3** – Third optional delta matrix added to the matrix product. It should have the same type as `src1` and `src2` .
> >
> > **beta** – Weight of `src3` .
> >
> > **dst** – Destination matrix. It has the proper size and the same type as input matrices.
> >
> > **flags** – Operation flags:
> >
> > – **GEMM_1_T** transpose `src1`
> >
> > – **GEMM_2_T** transpose `src2`
> >
> > – **GEMM_3_T** transpose `src3`
> >
> > **stream** – Stream for the asynchronous version.

The function performs generalized matrix multiplication similar to the `gemm` functions in BLAS level 3. For example, `gemm(src1, src2, alpha, src3, beta, dst, GEMM_1_T + GEMM_3_T)` corresponds to

$$dst = alpha \cdot src1^T \cdot src2 + beta \cdot src3^T$$

---

**Note:** Transposition operation doesn't support `CV_64FC2` input type.

---

**See Also:**

`gemm()`

## gpu::transpose

Transposes a matrix.

**C++:** void gpu::**transpose**(const GpuMat& **src1**, GpuMat& **dst**, Stream& **stream**=Stream::Null() )

---

**Parameters**

> **src1** – Source matrix. 1-, 4-, 8-byte element sizes are supported for now (CV_8UC1, CV_8UC4, CV_16UC2, CV_32FC1, etc).
>
> **dst** – Destination matrix.
>
> **stream** – Stream for the asynchronous version.

**See Also:**

transpose()

## gpu::flip

Flips a 2D matrix around vertical, horizontal, or both axes.

**C++:** void gpu::**flip**(const GpuMat& **a**, GpuMat& **b**, int **flipCode**, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **a** – Source matrix. Supports 1, 3 and 4 channels images with CV_8U, CV_16U, CV_32S or CV_32F depth.
> >
> > **b** – Destination matrix.
> >
> > **flipCode** – Flip mode for the source:
> >
> > – 0 Flips around x-axis.
> >
> > – >0 Flips around y-axis.
> >
> > – <0 Flips around both axes.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

flip()

## gpu::LUT

Transforms the source matrix into the destination matrix using the given look-up table: dst(I) = lut(src(I))

**C++:** void gpu::**LUT**(const GpuMat& **src**, const Mat& **lut**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source matrix. CV_8UC1 and CV_8UC3 matrices are supported for now.
> >
> > **lut** – Look-up table of 256 elements. It is a continuous CV_8U matrix.
> >
> > **dst** – Destination matrix with the same depth as lut and the same number of channels as src.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

LUT()

## gpu::merge

Makes a multi-channel matrix out of several single-channel matrices.

**C++:** void gpu::**merge**(const GpuMat* **src**, size_t **n**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**merge**(const vector<GpuMat>& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Array/vector of source matrices.
> >
> > **n** – Number of source matrices.
> >
> > **dst** – Destination matrix.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

merge()


## gpu::split

Copies each plane of a multi-channel matrix into an array.

**C++:** void gpu::**split**(const GpuMat& **src**, GpuMat* **dst**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**split**(const GpuMat& **src**, vector<GpuMat>& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source matrix.
> >
> > **dst** – Destination array/vector of single-channel matrices.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

split()


## gpu::magnitude

Computes magnitudes of complex matrix elements.

**C++:** void gpu::**magnitude**(const GpuMat& **xy**, GpuMat& **magnitude**, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**magnitude**(const GpuMat& **x**, const GpuMat& **y**, GpuMat& **magnitude**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **xy** – Source complex matrix in the interleaved format ( CV_32FC2 ).
> >
> > **x** – Source matrix containing real components ( CV_32FC1 ).
> >
> > **y** – Source matrix containing imaginary components ( CV_32FC1 ).
> >
> > **magnitude** – Destination matrix of float magnitudes ( CV_32FC1 ).
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

magnitude()

## gpu::magnitudeSqr

Computes squared magnitudes of complex matrix elements.

**C++:** void gpu::**magnitudeSqr**(const GpuMat& **xy**, GpuMat& **magnitude**, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**magnitudeSqr**(const GpuMat& **x**, const GpuMat& **y**, GpuMat& **magnitude**, Stream& **stream**=Stream::Null())

> **Parameters**

>> **xy** – Source complex matrix in the interleaved format ( CV_32FC2 ).

>> **x** – Source matrix containing real components ( CV_32FC1 ).

>> **y** – Source matrix containing imaginary components ( CV_32FC1 ).

>> **magnitude** – Destination matrix of float magnitude squares ( CV_32FC1 ).

>> **stream** – Stream for the asynchronous version.

## gpu::phase

Computes polar angles of complex matrix elements.

**C++:** void gpu::**phase**(const GpuMat& **x**, const GpuMat& **y**, GpuMat& **angle**, bool **angleInDegrees**=false, Stream& **stream**=Stream::Null())

> **Parameters**

>> **x** – Source matrix containing real components ( CV_32FC1 ).

>> **y** – Source matrix containing imaginary components ( CV_32FC1 ).

>> **angle** – Destination matrix of angles ( CV_32FC1 ).

>> **angleInDegrees** – Flag for angles that must be evaluated in degrees.

>> **stream** – Stream for the asynchronous version.

**See Also:**

phase()

## gpu::cartToPolar

Converts Cartesian coordinates into polar.

**C++:** void gpu::**cartToPolar**(const GpuMat& **x**, const GpuMat& **y**, GpuMat& **magnitude**, GpuMat& **angle**, bool **angleInDegrees**=false, Stream& **stream**=Stream::Null())

> **Parameters**

>> **x** – Source matrix containing real components ( CV_32FC1 ).

>> **y** – Source matrix containing imaginary components ( CV_32FC1 ).

>> **magnitude** – Destination matrix of float magnitudes ( CV_32FC1 ).

>> **angle** – Destination matrix of angles ( CV_32FC1 ).

>> **angleInDegrees** – Flag for angles that must be evaluated in degrees.

>> **stream** – Stream for the asynchronous version.

**See Also:**

cartToPolar()

## gpu::polarToCart

Converts polar coordinates into Cartesian.

**C++:** void gpu::**polarToCart**(const GpuMat& **magnitude**, const GpuMat& **angle**, GpuMat& **x**, GpuMat& **y**, bool **angleInDegrees**=false, Stream& **stream**=Stream::Null())

>       **Parameters**

>               **magnitude** – Source matrix containing magnitudes ( CV_32FC1 ).

>               **angle** – Source matrix containing angles ( CV_32FC1 ).

>               **x** – Destination matrix of real components ( CV_32FC1 ).

>               **y** – Destination matrix of imaginary components ( CV_32FC1 ).

>               **angleInDegrees** – Flag that indicates angles in degrees.

>               **stream** – Stream for the asynchronous version.

**See Also:**

polarToCart()

# 11.5 Per-element Operations

## gpu::add

Computes a matrix-matrix or matrix-scalar sum.

**C++:** void gpu::**add**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, const GpuMat& **mask**=GpuMat(), int **dtype**=-1, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**add**(const GpuMat& **a**, const Scalar& **sc**, GpuMat& **c**, const GpuMat& **mask**=GpuMat(), int **dtype**=-1, Stream& **stream**=Stream::Null() )

>       **Parameters**

>               **a** – First source matrix.

>               **b** – Second source matrix to be added to a . Matrix should have the same size and type as a .

>               **sc** – A scalar to be added to a .

>               **c** – Destination matrix that has the same size and number of channels as the input array(s). The depth is defined by dtype or a depth.

>               **mask** – Optional operation mask, 8-bit single channel array, that specifies elements of the destination array to be changed.

>               **dtype** – Optional depth of the output array.

>               **stream** – Stream for the asynchronous version.

**See Also:**

add()

## gpu::subtract

Computes a matrix-matrix or matrix-scalar difference.

**C++:** void gpu::**subtract**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, const GpuMat& **mask**=GpuMat(), int **dtype**=-1, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**subtract**(const GpuMat& **a**, const Scalar& **sc**, GpuMat& **c**, const GpuMat& **mask**=GpuMat(), int **dtype**=-1, Stream& **stream**=Stream::Null() )

> **Parameters**

>> **a** – First source matrix.

>> **b** – Second source matrix to be added to `a` . Matrix should have the same size and type as `a` .

>> **sc** – A scalar to be added to `a` .

>> **c** – Destination matrix that has the same size and number of channels as the input array(s). The depth is defined by `dtype` or `a` depth.

>> **mask** – Optional operation mask, 8-bit single channel array, that specifies elements of the destination array to be changed.

>> **dtype** – Optional depth of the output array.

>> **stream** – Stream for the asynchronous version.

**See Also:**

subtract()

## gpu::multiply

Computes a matrix-matrix or matrix-scalar per-element product.

**C++:** void gpu::**multiply**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, double **scale**=1, int **dtype**=-1, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**multiply**(const GpuMat& **a**, const Scalar& **sc**, GpuMat& **c**, double **scale**=1, int **dtype**=-1, Stream& **stream**=Stream::Null() )

> **Parameters**

>> **a** – First source matrix.

>> **b** – Second source matrix to be multiplied by `a` elements.

>> **sc** – A scalar to be multiplied by `a` elements.

>> **c** – Destination matrix that has the same size and number of channels as the input array(s). The depth is defined by `dtype` or `a` depth.

>> **scale** – Optional scale factor.

>> **dtype** – Optional depth of the output array.

>> **stream** – Stream for the asynchronous version.

**See Also:**

multiply()

## gpu::divide

Computes a matrix-matrix or matrix-scalar division.

**C++:** void gpu::**divide**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, double **scale**=1, int **dtype**=-1, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**divide**(const GpuMat& **a**, const Scalar& **sc**, GpuMat& **c**, double **scale**=1, int **dtype**=-1, Stream& **stream**=Stream::Null())

**C++:** void gpu::**divide**(double **scale**, const GpuMat& **b**, GpuMat& **c**, int **dtype**=-1, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **a** – First source matrix or a scalar.
> >
> > **b** – Second source matrix. The a elements are divided by it.
> >
> > **sc** – A scalar to be divided by the elements of a matrix.
> >
> > **c** – Destination matrix that has the same size and number of channels as the input array(s). The depth is defined by dtype or a depth.
> >
> > **scale** – Optional scale factor.
> >
> > **dtype** – Optional depth of the output array.
> >
> > **stream** – Stream for the asynchronous version.

This function, in contrast to divide(), uses a round-down rounding mode.

**See Also:**

divide()

## gpu::addWeighted

Computes the weighted sum of two arrays.

**C++:** void gpu::**addWeighted**(const GpuMat& **src1**, double **alpha**, const GpuMat& **src2**, double **beta**, double **gamma**, GpuMat& **dst**, int **dtype**=-1, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src1** – First source array.
> >
> > **alpha** – Weight for the first array elements.
> >
> > **src2** – Second source array of the same size and channel number as src1 .
> >
> > **beta** – Weight for the second array elements.
> >
> > **dst** – Destination array that has the same size and number of channels as the input arrays.
> >
> > **gamma** – Scalar added to each sum.
> >
> > **dtype** – Optional depth of the destination array. When both input arrays have the same depth, dtype can be set to -1, which will be equivalent to src1.depth().
> >
> > **stream** – Stream for the asynchronous version.

The function addWeighted calculates the weighted sum of two arrays as follows:

$$\texttt{dst}(I) = \texttt{saturate}(\texttt{src1}(I) * \texttt{alpha} + \texttt{src2}(I) * \texttt{beta} + \texttt{gamma})$$

where I is a multi-dimensional index of array elements. In case of multi-channel arrays, each channel is processed independently.

**See Also:**

addWeighted()

## gpu::abs

Computes an absolute value of each matrix element.

**C++:** void gpu::**abs**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source matrix. Supports CV_16S and CV_32F depth.
> >
> > **dst** – Destination matrix with the same size and type as src .
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

abs()

## gpu::sqr

Computes a square value of each matrix element.

**C++:** void gpu::**sqr**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source matrix. Supports CV_8U , CV_16U , CV_16S and CV_32F depth.
> >
> > **dst** – Destination matrix with the same size and type as src .
> >
> > **stream** – Stream for the asynchronous version.

## gpu::sqrt

Computes a square root of each matrix element.

**C++:** void gpu::**sqrt**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source matrix. Supports CV_8U , CV_16U , CV_16S and CV_32F depth.
> >
> > **dst** – Destination matrix with the same size and type as src .
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

sqrt()

## gpu::exp

Computes an exponent of each matrix element.

**C++:** void gpu::**exp**(const GpuMat& **a**, GpuMat& **b**, Stream& **stream**=Stream::Null() )

> **Parameters**
>
>> **a** – Source matrix. Supports `CV_8U` , `CV_16U` , `CV_16S` and `CV_32F` depth.
>>
>> **b** – Destination matrix with the same size and type as `a` .
>>
>> **stream** – Stream for the asynchronous version.

See Also:

exp()

## gpu::log

Computes a natural logarithm of absolute value of each matrix element.

**C++:** void gpu::**log**(const GpuMat& **a**, GpuMat& **b**, Stream& **stream**=Stream::Null() )

> **Parameters**
>
>> **a** – Source matrix. Supports `CV_8U` , `CV_16U` , `CV_16S` and `CV_32F` depth.
>>
>> **b** – Destination matrix with the same size and type as `a` .
>>
>> **stream** – Stream for the asynchronous version.

See Also:

log()

## gpu::pow

Raises every matrix element to a power.

**C++:** void gpu::**pow**(const GpuMat& **src**, double **power**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
>> **src** – Source matrix. Supports all type, except `CV_64F` depth.
>>
>> **power** – Exponent of power.
>>
>> **dst** – Destination matrix with the same size and type as `src` .
>>
>> **stream** – Stream for the asynchronous version.

The function pow raises every element of the input matrix to p :

$$\mathtt{dst}(I) = \begin{cases} \mathtt{src}(I)^p & \text{if p is integer} \\ |\mathtt{src}(I)|^p & \text{otherwise} \end{cases}$$

See Also:

pow()

## gpu::absdiff

Computes per-element absolute difference of two matrices (or of a matrix and scalar).

**C++:** void gpu::**absdiff**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, Stream& **stream**=Stream::Null() )

**C++:** void gpu::**absdiff**(const GpuMat& **a**, const Scalar& **s**, GpuMat& **c**, Stream& **stream**=Stream::Null() )

> Parameters
>
> > **a** – First source matrix.
> >
> > **b** – Second source matrix to be added to `a` .
> >
> > **s** – A scalar to be added to `a` .
> >
> > **c** – Destination matrix with the same size and type as `a` .
> >
> > **stream** – Stream for the asynchronous version.

See Also:

absdiff()


## gpu::compare

Compares elements of two matrices.

**C++:** void gpu::**compare**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, int **cmpop**, Stream& **stream**=Stream::Null() )

> Parameters
>
> > **a** – First source matrix.
> >
> > **b** – Second source matrix with the same size and type as `a` .
> >
> > **sc** – A scalar to be compared with `a` .
> >
> > **c** – Destination matrix with the same size as `a` and the `CV_8UC1` type.
> >
> > **cmpop** – Flag specifying the relation between the elements to be checked:
> >
> > – **CMP_EQ:** `a(.) == b(.)`
> >
> > – **CMP_GT:** `a(.) < b(.)`
> >
> > – **CMP_GE:** `a(.) <= b(.)`
> >
> > – **CMP_LT:** `a(.) < b(.)`
> >
> > – **CMP_LE:** `a(.) <= b(.)`
> >
> > – **CMP_NE:** `a(.) != b(.)`
> >
> > **stream** – Stream for the asynchronous version.

See Also:

compare()

## gpu::bitwise_not

Performs a per-element bitwise inversion.

**C++:** void gpu::**bitwise_not**(const GpuMat& **src**, GpuMat& **dst**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

> **Parameters**
>
>> **src** – Source matrix.
>>
>> **dst** – Destination matrix with the same size and type as `src` .
>>
>> **mask** – Optional operation mask. 8-bit single channel image.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::bitwise_or

Performs a per-element bitwise disjunction of two matrices or of matrix and scalar.

**C++:** void gpu::**bitwise_or**(const GpuMat& **src1**, const GpuMat& **src2**, GpuMat& **dst**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::**bitwise_or**(const GpuMat& **src1**, const Scalar& **sc**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
>> **src1** – First source matrix.
>>
>> **src2** – Second source matrix with the same size and type as `src1` .
>>
>> **dst** – Destination matrix with the same size and type as `src1` .
>>
>> **mask** – Optional operation mask. 8-bit single channel image.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::bitwise_and

Performs a per-element bitwise conjunction of two matrices or of matrix and scalar.

**C++:** void gpu::**bitwise_and**(const GpuMat& **src1**, const GpuMat& **src2**, GpuMat& **dst**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::**bitwise_and**(const GpuMat& **src1**, const Scalar& **sc**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> **Parameters**
>
>> **src1** – First source matrix.
>>
>> **src2** – Second source matrix with the same size and type as `src1` .
>>
>> **dst** – Destination matrix with the same size and type as `src1` .
>>
>> **mask** – Optional operation mask. 8-bit single channel image.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::bitwise_xor

Performs a per-element bitwise `exclusive or` operation of two matrices of matrix and scalar.

**C++:** void gpu::**bitwise_xor**(const GpuMat& **src1**, const GpuMat& **src2**, GpuMat& **dst**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::**bitwise_xor**(const GpuMat& **src1**, const Scalar& **sc**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

> Parameters

>> **src1** – First source matrix.

>> **src2** – Second source matrix with the same size and type as `src1` .

>> **dst** – Destination matrix with the same size and type as `src1` .

>> **mask** – Optional operation mask. 8-bit single channel image.

>> **stream** – Stream for the asynchronous version.

## gpu::rshift

Performs pixel by pixel right shift of an image by a constant value.

**C++:** void gpu::**rshift**(const GpuMat& **src**, Scalar_<int> **sc**, GpuMat& **dst**, Stream& **stream**=Stream::Null() )

> Parameters

>> **src** – Source matrix. Supports 1, 3 and 4 channels images with integers elements.

>> **sc** – Constant values, one per channel.

>> **dst** – Destination matrix with the same size and type as `src` .

>> **stream** – Stream for the asynchronous version.

## gpu::lshift

Performs pixel by pixel right left of an image by a constant value.

**C++:** void gpu::**lshift**(const GpuMat& **src**, Scalar_<int> **sc**, GpuMat& **dst**, Stream& **stream**=Stream::Null() )

> Parameters

>> **src** – Source matrix. Supports 1, 3 and 4 channels images with `CV_8U` , `CV_16U` or `CV_32S` depth.

>> **sc** – Constant values, one per channel.

>> **dst** – Destination matrix with the same size and type as `src` .

>> **stream** – Stream for the asynchronous version.

## gpu::min

Computes the per-element minimum of two matrices (or a matrix and a scalar).

**C++:** void gpu::**min**(const GpuMat& **src1**, const GpuMat& **src2**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

---

**C++:** void gpu::**min**(const GpuMat& **src1**, double **src2**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

>> **Parameters**

>>> **src1** – First source matrix.

>>> **src2** – Second source matrix or a scalar to compare `src1` elements with.

>>> **dst** – Destination matrix with the same size and type as `src1` .

>>> **stream** – Stream for the asynchronous version.

**See Also:**

min()

## gpu::max

Computes the per-element maximum of two matrices (or a matrix and a scalar).

**C++:** void gpu::**max**(const GpuMat& **src1**, const GpuMat& **src2**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**max**(const GpuMat& **src1**, double **src2**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

>> **Parameters**

>>> **src1** – First source matrix.

>>> **src2** – Second source matrix or a scalar to compare `src1` elements with.

>>> **dst** – Destination matrix with the same size and type as `src1` .

>>> **stream** – Stream for the asynchronous version.

**See Also:**

max()

# 11.6 Image Processing

## gpu::meanShiftFiltering

Performs mean-shift filtering for each point of the source image.

**C++:** void gpu::**meanShiftFiltering**(const GpuMat& **src**, GpuMat& **dst**, int **sp**, int **sr**, TermCriteria **criteria**=TermCriteria(TermCriteria::MAX_ITER + TermCriteria::EPS, 5, 1), Stream& **stream**=Stream::Null() )

>> **Parameters**

>>> **src** – Source image. Only `CV_8UC4` images are supported for now.

>>> **dst** – Destination image containing the color of mapped points. It has the same size and type as `src` .

>>> **sp** – Spatial window radius.

>>> **sr** – Color window radius.

>>> **criteria** – Termination criteria. See `TermCriteria`.

It maps each point of the source image into another point. As a result, you have a new color and new position of each point.

## gpu::meanShiftProc

Performs a mean-shift procedure and stores information about processed points (their colors and positions) in two images.

**C++:** void gpu::**meanShiftProc**(const GpuMat& **src**, GpuMat& **dstr**, GpuMat& **dstsp**, int **sp**, int **sr**, TermCriteria **criteria**=TermCriteria(TermCriteria::MAX_ITER + TermCriteria::EPS, 5, 1), Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC4` images are supported for now.
> >
> > **dstr** – Destination image containing the color of mapped points. The size and type is the same as `src` .
> >
> > **dstsp** – Destination image containing the position of mapped points. The size is the same as `src` size. The type is `CV_16SC2` .
> >
> > **sp** – Spatial window radius.
> >
> > **sr** – Color window radius.
> >
> > **criteria** – Termination criteria. See `TermCriteria`.

**See Also:**

`gpu::meanShiftFiltering()`

## gpu::meanShiftSegmentation

Performs a mean-shift segmentation of the source image and eliminates small segments.

**C++:** void gpu::**meanShiftSegmentation**(const GpuMat& **src**, Mat& **dst**, int **sp**, int **sr**, int **minsize**, TermCriteria **criteria**=TermCriteria(TermCriteria::MAX_ITER + TermCriteria::EPS, 5, 1))

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC4` images are supported for now.
> >
> > **dst** – Segmented image with the same size and type as `src` .
> >
> > **sp** – Spatial window radius.
> >
> > **sr** – Color window radius.
> >
> > **minsize** – Minimum segment size. Smaller segments are merged.
> >
> > **criteria** – Termination criteria. See `TermCriteria`.

## gpu::integral

Computes an integral image.

**C++:** void gpu::**integral**(const GpuMat& **src**, GpuMat& **sum**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC1` images are supported for now.
> >
> > **sum** – Integral image containing 32-bit unsigned integer values packed into `CV_32SC1` .
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

`integral()`

## gpu::sqrIntegral

Computes a squared integral image.

**C++:** void gpu::**sqrIntegral**(const GpuMat& **src**, GpuMat& **sqsum**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC1` images are supported for now.
> >
> > **sqsum** – Squared integral image containing 64-bit unsigned integer values packed into `CV_64FC1`.
> >
> > **stream** – Stream for the asynchronous version.

## gpu::columnSum

Computes a vertical (column) sum.

**C++:** void gpu::**columnSum**(const GpuMat& **src**, GpuMat& **sum**)

> **Parameters**
>
> > **src** – Source image. Only `CV_32FC1` images are supported for now.
> >
> > **sum** – Destination image of the `CV_32FC1` type.

## gpu::cornerHarris

Computes the Harris cornerness criteria at each image pixel.

**C++:** void gpu::**cornerHarris**(const GpuMat& **src**, GpuMat& **dst**, int **blockSize**, int **ksize**, double **k**, int **borderType**=BORDER_REFLECT101)

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC1` and `CV_32FC1` images are supported for now.
> >
> > **dst** – Destination image containing cornerness values. It has the same size as `src` and `CV_32FC1` type.
> >
> > **blockSize** – Neighborhood size.
> >
> > **ksize** – Aperture parameter for the Sobel operator.
> >
> > **k** – Harris detector free parameter.
> >
> > **borderType** – Pixel extrapolation method. Only `BORDER_REFLECT101` and `BORDER_REPLICATE` are supported for now.

**See Also:**

`cornerHarris()`

## gpu::cornerMinEigenVal

Computes the minimum eigen value of a 2x2 derivative covariation matrix at each pixel (the cornerness criteria).

**C++:** void gpu::**cornerMinEigenVal**(const GpuMat& **src**, GpuMat& **dst**, int **blockSize**, int **ksize**, int **borderType**=BORDER_REFLECT101)

**C++:** void gpu::**cornerMinEigenVal**(const GpuMat& **src**, GpuMat& **dst**, GpuMat& **Dx**, GpuMat& **Dy**, int **blockSize**, int **ksize**, int **borderType**=BORDER_REFLECT101)

**C++:** void gpu::**cornerMinEigenVal**(const GpuMat& **src**, GpuMat& **dst**, GpuMat& **Dx**, GpuMat& **Dy**, GpuMat& **buf**, int **blockSize**, int **ksize**, int **borderType**=BORDER_REFLECT101, Stream& **stream**=Stream::Null())

> Parameters
>> **src** – Source image. Only `CV_8UC1` and `CV_32FC1` images are supported for now.
>>
>> **dst** – Destination image containing cornerness values. The size is the same. The type is `CV_32FC1` .
>>
>> **blockSize** – Neighborhood size.
>>
>> **ksize** – Aperture parameter for the Sobel operator.
>>
>> **borderType** – Pixel extrapolation method. Only `BORDER_REFLECT101` and `BORDER_REPLICATE` are supported for now.

**See Also:**

cornerMinEigenVal()

## gpu::mulSpectrums

Performs a per-element multiplication of two Fourier spectrums.

**C++:** void gpu::**mulSpectrums**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, int **flags**, bool **conjB**=false, Stream& **stream**=Stream::Null() )

> Parameters
>> **a** – First spectrum.
>>
>> **b** – Second spectrum with the same size and type as a .
>>
>> **c** – Destination spectrum.
>>
>> **flags** – Mock parameter used for CPU/GPU interfaces similarity.
>>
>> **conjB** – Optional flag to specify if the second spectrum needs to be conjugated before the multiplication.
>
> Only full (not packed) `CV_32FC2` complex spectrums in the interleaved format are supported for now.

**See Also:**

mulSpectrums()

## gpu::mulAndScaleSpectrums

Performs a per-element multiplication of two Fourier spectrums and scales the result.

**C++:** void gpu::**mulAndScaleSpectrums**(const GpuMat& **a**, const GpuMat& **b**, GpuMat& **c**, int **flags**, float **scale**, bool **conjB**=false, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **a** – First spectrum.
> >
> > **b** – Second spectrum with the same size and type as `a` .
> >
> > **c** – Destination spectrum.
> >
> > **flags** – Mock parameter used for CPU/GPU interfaces similarity.
> >
> > **scale** – Scale constant.
> >
> > **conjB** – Optional flag to specify if the second spectrum needs to be conjugated before the multiplication.
>
> Only full (not packed) `CV_32FC2` complex spectrums in the interleaved format are supported for now.

**See Also:**

`mulSpectrums()`

## gpu::dft

Performs a forward or inverse discrete Fourier transform (1D or 2D) of the floating point matrix.

**C++:** void gpu::**dft**(const GpuMat& **src**, GpuMat& **dst**, Size **dft_size**, int **flags**=0, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source matrix (real or complex).
> >
> > **dst** – Destination matrix (real or complex).
> >
> > **dft_size** – Size of a discrete Fourier transform.
> >
> > **flags** – Optional flags:
> >
> > – **DFT_ROWS** transforms each individual row of the source matrix.
> >
> > – **DFT_SCALE** scales the result: divide it by the number of elements in the transform (obtained from `dft_size` ).
> >
> > – **DFT_INVERSE** inverts DFT. Use for complex-complex cases (real-complex and complex-real cases are always forward and inverse, respectively).
> >
> > – **DFT_REAL_OUTPUT** specifies the output as real. The source matrix is the result of real-complex transform, so the destination matrix must be real.

Use to handle real matrices ( `CV32FC1` ) and complex matrices in the interleaved format ( `CV32FC2` ).

The source matrix should be continuous, otherwise reallocation and data copying is performed. The function chooses an operation mode depending on the flags, size, and channel count of the source matrix:

- If the source matrix is complex and the output is not specified as real, the destination matrix is complex and has the `dft_size` size and `CV_32FC2` type. The destination matrix contains a full result of the DFT (forward or inverse).

- If the source matrix is complex and the output is specified as real, the function assumes that its input is the result of the forward transform (see the next item). The destination matrix has the `dft_size` size and `CV_32FC1` type. It contains the result of the inverse DFT.

- If the source matrix is real (its type is `CV_32FC1` ), forward DFT is performed. The result of the DFT is packed into complex ( `CV_32FC2` ) matrix. So, the width of the destination matrix is `dft_size.width / 2 + 1` . But if the source is a single column, the height is reduced instead of the width.

**See Also:**

dft()

## gpu::ConvolveBuf

**struct** gpu::**ConvolveBuf**

Class providing a memory buffer for gpu::convolve() function, plus it allows to adjust some specific parameters.

```
struct CV_EXPORTS ConvolveBuf
{
    Size result_size;
    Size block_size;
    Size user_block_size;
    Size dft_size;
    int spect_len;

    GpuMat image_spect, templ_spect, result_spect;
    GpuMat image_block, templ_block, result_data;

    void create(Size image_size, Size templ_size);
    static Size estimateBlockSize(Size result_size, Size templ_size);
};
```

You can use field *user_block_size* to set specific block size for gpu::convolve() function. If you leave its default value *Size(0,0)* then automatic estimation of block size will be used (which is optimized for speed). By varying *user_block_size* you can reduce memory requirements at the cost of speed.

## gpu::ConvolveBuf::create

**C++:** gpu::ConvolveBuf::**create**(Size **image_size**, Size **templ_size**)

Constructs a buffer for gpu::convolve() function with respective arguments.

## gpu::convolve

Computes a convolution (or cross-correlation) of two images.

**C++:** void gpu::**convolve**(const GpuMat& **image**, const GpuMat& **templ**, GpuMat& **result**, bool **ccorr**=false)

**C++:** void gpu::**convolve**(const GpuMat& **image**, const GpuMat& **templ**, GpuMat& **result**, bool **ccorr**, ConvolveBuf& **buf**, Stream& **stream**=Stream::Null() )

>   **Parameters**

>>      **image** – Source image. Only CV_32FC1 images are supported for now.

>>      **templ** – Template image. The size is not greater than the image size. The type is the same as image .

>>      **result** – Result image. If image is *W x H* and templ is *w x h*, then result must be *W-w+1 x H-h+1*.

>>      **ccorr** – Flags to evaluate cross-correlation instead of convolution.

>>      **buf** – Optional buffer to avoid extra memory allocations and to adjust some specific parameters. See gpu::ConvolveBuf.

---

> **stream** – Stream for the asynchronous version.

**See Also:**

`gpu::filter2D()`

## gpu::MatchTemplateBuf

**struct** gpu::**MatchTemplateBuf**

Class providing memory buffers for `gpu::matchTemplate()` function, plus it allows to adjust some specific parameters.

```
struct CV_EXPORTS MatchTemplateBuf
{
    Size user_block_size;
    GpuMat imagef, templf;
    std::vector<GpuMat> images;
    std::vector<GpuMat> image_sums;
    std::vector<GpuMat> image_sqsums;
};
```

You can use field *user_block_size* to set specific block size for `gpu::matchTemplate()` function. If you leave its default value *Size(0,0)* then automatic estimation of block size will be used (which is optimized for speed). By varying *user_block_size* you can reduce memory requirements at the cost of speed.

## gpu::matchTemplate

Computes a proximity map for a raster template and an image where the template is searched for.

**C++:** void gpu::**matchTemplate**(const GpuMat& **image**, const GpuMat& **templ**, GpuMat& **result**, int **method**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**matchTemplate**(const GpuMat& **image**, const GpuMat& **templ**, GpuMat& **result**, int **method**, MatchTemplateBuf& **buf**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **image** – Source image. `CV_32F` and `CV_8U` depth images (1..4 channels) are supported for now.
> >
> > **templ** – Template image with the size and type the same as `image` .
> >
> > **result** – Map containing comparison results ( `CV_32FC1` ). If `image` is *W x H* and `templ` is *w x h*, then `result` must be *W-w+1 x H-h+1*.
> >
> > **method** – Specifies the way to compare the template with the image.
> >
> > **buf** – Optional buffer to avoid extra memory allocations and to adjust some specific parameters. See `gpu::MatchTemplateBuf`.
> >
> > **stream** – Stream for the asynchronous version.
>
> The following methods are supported for the `CV_8U` depth images for now:
>
> > - `CV_TM_SQDIFF`
> > - `CV_TM_SQDIFF_NORMED`
> > - `CV_TM_CCORR`
> > - `CV_TM_CCORR_NORMED`

- `CV_TM_CCOEFF`

- `CV_TM_CCOEFF_NORMED`

The following methods are supported for the `CV_32F` images for now:

- `CV_TM_SQDIFF`

- `CV_TM_CCORR`

**See Also:**

`matchTemplate()`

## gpu::remap

Applies a generic geometrical transformation to an image.

**C++:** void gpu::**remap**(const GpuMat& **src**, GpuMat& **dst**, const GpuMat& **xmap**, const GpuMat& **ymap**, int **interpolation**, int **borderMode**=BORDER_CONSTANT, Scalar **border-Value**=Scalar(), Stream& **stream**=Stream::Null() )

> **Parameters**

>> **src** – Source image.

>> **dst** – Destination image with the size the same as xmap and the type the same as `src` .

>> **xmap** – X values. Only `CV_32FC1` type is supported.

>> **ymap** – Y values. Only `CV_32FC1` type is supported.

>> **interpolation** – Interpolation method (see `resize()` ). INTER_NEAREST , INTER_LINEAR and INTER_CUBIC are supported for now.

>> **borderMode** – Pixel extrapolation method (see `borderInterpolate()` ). BORDER_REFLECT101 , BORDER_REPLICATE , BORDER_CONSTANT , BORDER_REFLECT and BORDER_WRAP are supported for now.

>> **borderValue** – Value used in case of a constant border. By default, it is 0.

>> **stream** – Stream for the asynchronous version.

The function transforms the source image using the specified map:

$$\texttt{dst}(x, y) = \texttt{src}(xmap(x, y), ymap(x, y))$$

Values of pixels with non-integer coordinates are computed using the bilinear interpolation.

**See Also:**

`remap()`

## gpu::cvtColor

Converts an image from one color space to another.

**C++:** void gpu::**cvtColor**(const GpuMat& **src**, GpuMat& **dst**, int **code**, int **dcn**=0, Stream& **stream**=Stream::Null())

> **Parameters**

>> **src** – Source image with `CV_8U` , `CV_16U` , or `CV_32F` depth and 1, 3, or 4 channels.

>> **dst** – Destination image with the same size and depth as `src` .

> **code** – Color space conversion code. For details, see `cvtColor()` . Conversion to/from Luv and Bayer color spaces is not supported.
>
> **dcn** – Number of channels in the destination image. If the parameter is 0, the number of the channels is derived automatically from `src` and the `code` .
>
> **stream** – Stream for the asynchronous version.

3-channel color spaces (like `HSV`, `XYZ`, and so on) can be stored in a 4-channel image for better performance.

**See Also:**

`cvtColor()`

## gpu::swapChannels

Exchanges the color channels of an image in-place.

**C++:** void gpu::**swapChannels**(GpuMat& **image**, const int **dstOrder**[4], Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **image** – Source image. Supports only `CV_8UC4` type.
> >
> > **dstOrder** – Integer array describing how channel values are permutated. The n-th entry of the array contains the number of the channel that is stored in the n-th channel of the output image. E.g. Given an RGBA image, aDstOrder = [3,2,1,0] converts this to ABGR channel order.
> >
> > **stream** – Stream for the asynchronous version.

The methods support arbitrary permutations of the original channels, including replication.

## gpu::threshold

Applies a fixed-level threshold to each array element.

**C++:** double gpu::**threshold**(const GpuMat& **src**, GpuMat& **dst**, double **thresh**, double **maxval**, int **type**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source array (single-channel).
> >
> > **dst** – Destination array with the same size and type as `src` .
> >
> > **thresh** – Threshold value.
> >
> > **maxval** – Maximum value to use with `THRESH_BINARY` and `THRESH_BINARY_INV` threshold types.
> >
> > **type** – Threshold type. For details, see `threshold()` . The `THRESH_OTSU` threshold type is not supported.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

`threshold()`

## gpu::resize

Resizes an image.

**C++:** void gpu::**resize**(const GpuMat& **src**, GpuMat& **dst**, Size **dsize**, double **fx**=0, double **fy**=0, int **interpolation**=INTER_LINEAR, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image.
> >
> > **dst** – Destination image with the same type as `src`. The size is `dsize` (when it is non-zero) or the size is computed from `src.size()`, `fx`, and `fy`.
> >
> > **dsize** – Destination image size. If it is zero, it is computed as:
> >
> > > ```
> > > dsize = Size(round(fx*src.cols), round(fy*src.rows))
> > > ```
> >
> > Either `dsize` or both `fx` and `fy` must be non-zero.
> >
> > **fx** – Scale factor along the horizontal axis. If it is zero, it is computed as:
> >
> > > ```
> > > (double)dsize.width/src.cols
> > > ```
> >
> > **fy** – Scale factor along the vertical axis. If it is zero, it is computed as:
> >
> > > ```
> > > (double)dsize.height/src.rows
> > > ```
> >
> > **interpolation** – Interpolation method. `INTER_NEAREST`, `INTER_LINEAR` and `INTER_CUBIC` are supported for now.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

resize()

## gpu::warpAffine

Applies an affine transformation to an image.

**C++:** void gpu::**warpAffine**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **M**, Size **dsize**, int **flags**=INTER_LINEAR, int **borderMode**=BORDER_CONSTANT, Scalar **borderValue**=Scalar(), Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. `CV_8U`, `CV_16U`, `CV_32S`, or `CV_32F` depth and 1, 3, or 4 channels are supported.
> >
> > **dst** – Destination image with the same type as `src`. The size is `dsize`.
> >
> > **M** – *2x3* transformation matrix.
> >
> > **dsize** – Size of the destination image.
> >
> > **flags** – Combination of interpolation methods (see resize()) and the optional flag WARP_INVERSE_MAP specifying that M is an inverse transformation ( `dst=>src` ). Only `INTER_NEAREST`, `INTER_LINEAR`, and `INTER_CUBIC` interpolation methods are supported.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

warpAffine()

## gpu::buildWarpAffineMaps

Builds transformation maps for affine transformation.

**C++:** void gpu::**buildWarpAffineMaps**(const Mat& **M**, bool **inverse**, Size **dsize**, GpuMat& **xmap**, GpuMat& **ymap**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **M** – *2x3* transformation matrix.
> >
> > **inverse** – Flag specifying that M is an inverse transformation ( dst=>src ).
> >
> > **dsize** – Size of the destination image.
> >
> > **xmap** – X values with CV_32FC1 type.
> >
> > **ymap** – Y values with CV_32FC1 type.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

gpu::warpAffine() , gpu::remap()

## gpu::warpPerspective

Applies a perspective transformation to an image.

**C++:** void gpu::**warpPerspective**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **M**, Size **dsize**, int **flags**=INTER_LINEAR, int **borderMode**=BORDER_CONSTANT, Scalar **borderValue**=Scalar(), Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. CV_8U , CV_16U , CV_32S , or CV_32F depth and 1, 3, or 4 channels are supported.
> >
> > **dst** – Destination image with the same type as src . The size is dsize .
> >
> > **M** – *3x3* transformation matrix.
> >
> > **dsize** – Size of the destination image.
> >
> > **flags** – Combination of interpolation methods (see resize() ) and the optional flag WARP_INVERSE_MAP specifying that M is the inverse transformation ( dst => src ). Only INTER_NEAREST , INTER_LINEAR , and INTER_CUBIC interpolation methods are supported.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

warpPerspective()

## gpu::buildWarpPerspectiveMaps

Builds transformation maps for perspective transformation.

**C++:** void gpu::**buildWarpAffineMaps**(const Mat& **M**, bool **inverse**, Size **dsize**, GpuMat& **xmap**, GpuMat& **ymap**, Stream& **stream**=Stream::Null())

> Parameters

>> **M** – *3x3* transformation matrix.

>> **inverse** – Flag specifying that `M` is an inverse transformation ( `dst=>src` ).

>> **dsize** – Size of the destination image.

>> **xmap** – X values with `CV_32FC1` type.

>> **ymap** – Y values with `CV_32FC1` type.

>> **stream** – Stream for the asynchronous version.

See Also:

gpu::warpPerspective() , gpu::remap()

## gpu::rotate

Rotates an image around the origin (0,0) and then shifts it.

**C++:** void gpu::**rotate**(const GpuMat& **src**, GpuMat& **dst**, Size **dsize**, double **angle**, double **xShift**=0, double **yShift**=0, int **interpolation**=INTER_LINEAR, Stream& **stream**=Stream::Null())

> Parameters

>> **src** – Source image. Supports 1, 3 or 4 channels images with `CV_8U` , `CV_16U` or `CV_32F` depth.

>> **dst** – Destination image with the same type as `src` . The size is `dsize` .

>> **dsize** – Size of the destination image.

>> **angle** – Angle of rotation in degrees.

>> **xShift** – Shift along the horizontal axis.

>> **yShift** – Shift along the vertical axis.

>> **interpolation** – Interpolation method. Only `INTER_NEAREST` , `INTER_LINEAR` , and `INTER_CUBIC` are supported.

>> **stream** – Stream for the asynchronous version.

See Also:

gpu::warpAffine()

## gpu::copyMakeBorder

Forms a border around an image.

**C++:** void gpu::**copyMakeBorder**(const GpuMat& **src**, GpuMat& **dst**, int **top**, int **bottom**, int **left**, int **right**, int **borderType**, const Scalar& **value**=Scalar(), Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. `CV_8UC1` , `CV_8UC4` , `CV_32SC1` , and `CV_32FC1` types are supported.
> >
> > **dst** – Destination image with the same type as `src`. The size is `Size(src.cols+left+right, src.rows+top+bottom)`.
> >
> > **top** –
> >
> > **bottom** –
> >
> > **left** –
> >
> > **right** – Number of pixels in each direction from the source image rectangle to extrapolate. For example: `top=1, bottom=1, left=1, right=1` mean that 1 pixel-wide border needs to be built.
> >
> > **borderType** – Border type. See `borderInterpolate()` for details. `BORDER_REFLECT101` , `BORDER_REPLICATE` , `BORDER_CONSTANT` , `BORDER_REFLECT` and `BORDER_WRAP` are supported for now.
> >
> > **value** – Border value.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

copyMakeBorder()

## gpu::rectStdDev

Computes a standard deviation of integral images.

**C++:** void gpu::**rectStdDev**(const GpuMat& **src**, const GpuMat& **sqr**, GpuMat& **dst**, const Rect& **rect**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. Only the `CV_32SC1` type is supported.
> >
> > **sqr** – Squared source image. Only the `CV_32FC1` type is supported.
> >
> > **dst** – Destination image with the same type and size as `src` .
> >
> > **rect** – Rectangular window.
> >
> > **stream** – Stream for the asynchronous version.

## gpu::evenLevels

Computes levels with even distribution.

**C++:** void gpu::**evenLevels**(GpuMat& **levels**, int **nLevels**, int **lowerLevel**, int **upperLevel**)

> **Parameters**
>
> > **levels** – Destination array. `levels` has 1 row, `nLevels` columns, and the `CV_32SC1` type.
> >
> > **nLevels** – Number of computed levels. `nLevels` must be at least 2.
> >
> > **lowerLevel** – Lower boundary value of the lowest level.
> >
> > **upperLevel** – Upper boundary value of the greatest level.

## gpu::histEven

Calculates a histogram with evenly distributed bins.

**C++:** void gpu::**histEven**(const GpuMat& **src**, GpuMat& **hist**, int **histSize**, int **lowerLevel**, int **upperLevel**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**histEven**(const GpuMat& **src**, GpuMat& **hist**, GpuMat& **buf**, int **histSize**, int **lowerLevel**, int **upperLevel**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**histEven**(const GpuMat& **src**, GpuMat **hist**[4], int **histSize**[4], int **lowerLevel**[4], int **upperLevel**[4], Stream& **stream**=Stream::Null() )

**C++:** void gpu::**histEven**(const GpuMat& **src**, GpuMat **hist**[4], GpuMat& **buf**, int **histSize**[4], int **lowerLevel**[4], int **upperLevel**[4], Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. `CV_8U`, `CV_16U`, or `CV_16S` depth and 1 or 4 channels are supported. For a four-channel image, all channels are processed separately.
> >
> > **hist** – Destination histogram with one row, `histSize` columns, and the `CV_32S` type.
> >
> > **histSize** – Size of the histogram.
> >
> > **lowerLevel** – Lower boundary of lowest-level bin.
> >
> > **upperLevel** – Upper boundary of highest-level bin.
> >
> > **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).
> >
> > **stream** – Stream for the asynchronous version.

## gpu::histRange

Calculates a histogram with bins determined by the `levels` array.

**C++:** void gpu::**histRange**(const GpuMat& **src**, GpuMat& **hist**, const GpuMat& **levels**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**histRange**(const GpuMat& **src**, GpuMat& **hist**, const GpuMat& **levels**, GpuMat& **buf**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. `CV_8U` , `CV_16U` , or `CV_16S` depth and 1 or 4 channels are supported. For a four-channel image, all channels are processed separately.
> >
> > **hist** – Destination histogram with one row, (`levels.cols-1`) columns, and the `CV_32SC1` type.
> >
> > **levels** – Number of levels in the histogram.
> >
> > **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).
> >
> > **stream** – Stream for the asynchronous version.

## gpu::calcHist

Calculates histogram for one channel 8-bit image.

**C++:** void gpu::**calcHist**(const GpuMat& **src**, GpuMat& **hist**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**calcHist**(const GpuMat& **src**, GpuMat& **hist**, GpuMat& **buf**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image.
> >
> > **hist** – Destination histogram with one row, 256 columns, and the `CV_32SC1` type.
> >
> > **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).
> >
> > **stream** – Stream for the asynchronous version.

## gpu::equalizeHist

Equalizes the histogram of a grayscale image.

**C++:** void gpu::**equalizeHist**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**equalizeHist**(const GpuMat& **src**, GpuMat& **dst**, GpuMat& **hist**, Stream& **stream**=Stream::Null())

**C++:** void gpu::**equalizeHist**(const GpuMat& **src**, GpuMat& **dst**, GpuMat& **hist**, GpuMat& **buf**, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Source image.
> >
> > **dst** – Destination image.
> >
> > **hist** – Destination histogram with one row, 256 columns, and the `CV_32SC1` type.
> >
> > **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

`equalizeHist()`

## gpu::buildWarpPlaneMaps

Builds plane warping maps.

**C++:** void gpu::**buildWarpPlaneMaps**(Size **src_size**, Rect **dst_roi**, const Mat& **K**, const Mat& **R**, const Mat& **T**, float **scale**, GpuMat& **map_x**, GpuMat& **map_y**, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **stream** – Stream for the asynchronous version.

## gpu::buildWarpCylindricalMaps

Builds cylindrical warping maps.

**C++:** void gpu::**buildWarpCylindricalMaps**(Size **src_size**, Rect **dst_roi**, const Mat& **K**, const Mat& **R**, float **scale**, GpuMat& **map_x**, GpuMat& **map_y**, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **stream** – Stream for the asynchronous version.

## gpu::buildWarpSphericalMaps

Builds spherical warping maps.

**C++:** void gpu::**buildWarpSphericalMaps**(Size **src_size**, Rect **dst_roi**, const Mat& **K**, const Mat& **R**, float **scale**, GpuMat& **map_x**, GpuMat& **map_y**, Stream& **stream**=Stream::Null() )

>    **Parameters**

>        **stream** – Stream for the asynchronous version.

## gpu::pyrDown

Smoothes an image and downsamples it.

**C++:** void gpu::**pyrDown**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

>    **Parameters**

>        **src** – Source image.

>        **dst** – Destination image. Will have `Size((src.cols+1)/2, (src.rows+1)/2)` size and the same type as `src`.

>        **stream** – Stream for the asynchronous version.

**See Also:**

`pyrDown()`

## gpu::pyrUp

Upsamples an image and then smoothes it.

**C++:** void gpu::**pyrUp**(const GpuMat& **src**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

>    **Parameters**

>        **src** – Source image.

>        **dst** – Destination image. Will have `Size(src.cols*2, src.rows*2)` size and the same type as `src`.

>        **stream** – Stream for the asynchronous version.

**See Also:**

`pyrUp()`

## gpu::blendLinear

Performs linear blending of two images.

**C++:** void gpu::**blendLinear**(const GpuMat& **img1**, const GpuMat& **img2**, const GpuMat& **weights1**, const GpuMat& **weights2**, GpuMat& **result**, Stream& **stream**=Stream::Null())

>    **Parameters**

>        **img1** – First image. Supports only `CV_8U` and `CV_32F` depth.

> **img2** – Second image. Must have the same size and the same type as `img1` .
>
> **weights1** – Weights for first image. Must have tha same size as `img1` . Supports only `CV_32F` type.
>
> **weights2** – Weights for second image. Must have tha same size as `img2` . Supports only `CV_32F` type.
>
> **result** – Destination image.
>
> **stream** – Stream for the asynchronous version.

## gpu::bilateralFilter

Performs bilateral filtering of passed image

**C++:** void gpu::**bilateralFilter**(const GpuMat& **src**, GpuMat& **dst**, int **kernel_size**, float **sigma_color**, float **sigma_spatial**, int **borderMode**=BORDER_DEFAULT, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. Supports only (channles != 2 && depth() != CV_8S && depth() != CV_32S && depth() != CV_64F).
> >
> > **dst** – Destination imagwe.
> >
> > **kernel_size** – Kernel window size.
> >
> > **sigma_color** – Filter sigma in the color space.
> >
> > **sigma_spatial** – Filter sigma in the coordinate space.
> >
> > **borderMode** – Border type. See `borderInterpolate()` for details. `BORDER_REFLECT101` , `BORDER_REPLICATE` , `BORDER_CONSTANT` , `BORDER_REFLECT` and `BORDER_WRAP` are supported for now.
> >
> > **stream** – Stream for the asynchronous version.

**See Also:**

`bilateralFilter()`,

## gpu::nonLocalMeans

Performs pure non local means denoising without any simplification, and thus it is not fast.

**C++:** void gpu::**nonLocalMeans**(const GpuMat& **src**, GpuMat& **dst**, float **h**, int **search_window**=21, int **block_size**=7, int **borderMode**=BORDER_DEFAULT, Stream& **s**=Stream::Null())

> **Parameters**
>
> > **src** – Source image. Supports only CV_8UC1, CV_8UC2 and CV_8UC3.
> >
> > **dst** – Destination image.
> >
> > **h** – Filter sigma regulating filter strength for color.
> >
> > **search_window** – Size of search window.
> >
> > **block_size** – Size of block used for computing weights.

> **borderMode** – Border type. See `borderInterpolate()` for details. `BORDER_REFLECT101`
> , `BORDER_REPLICATE` , `BORDER_CONSTANT` , `BORDER_REFLECT` and `BORDER_WRAP` are supported for now.
>
> **stream** – Stream for the asynchronous version.

**See Also:**

`fastNlMeansDenoising()`

## gpu::FastNonLocalMeansDenoising

**class** gpu::**FastNonLocalMeansDenoising**

> class FastNonLocalMeansDenoising { public:
>
> > //! Simple method, recommended for grayscale images (though it supports multichannel images) void simpleMethod(const GpuMat& src, GpuMat& dst, float h, int search_window = 21, int block_size = 7, Stream& s = Stream::Null())
> >
> > //! Processes luminance and color components separatelly void labMethod(const GpuMat& src, GpuMat& dst, float h_luminance, float h_color, int search_window = 21, int block_size = 7, Stream& s = Stream::Null())
>
> };

The class implements fast approximate Non Local Means Denoising algorithm.

## gpu::FastNonLocalMeansDenoising::simpleMethod()

Perform image denoising using Non-local Means Denoising algorithm http://www.ipol.im/pub/algo/bcm_non_local_means_denoising with several computational optimizations. Noise expected to be a gaussian white noise

**C++:** void gpu::FastNonLocalMeansDenoising::**simpleMethod**(const GpuMat& **src**, GpuMat& **dst**, float **h**, int **search_window**=21, int **block_size**=7, Stream& **s**=Stream::Null())

> **Parameters**
>
> > **src** – Input 8-bit 1-channel, 2-channel or 3-channel image.
> >
> > **dst** – Output image with the same size and type as `src` .
> >
> > **h** – Parameter regulating filter strength. Big h value perfectly removes noise but also removes image details, smaller h value preserves details but also preserves some noise
> >
> > **search_window** – Size in pixels of the window that is used to compute weighted average for given pixel. Should be odd. Affect performance linearly: greater search_window - greater denoising time. Recommended value 21 pixels
> >
> > **block_size** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels
> >
> > **stream** – Stream for the asynchronous invocations.

This function expected to be applied to grayscale images. For colored images look at FastNonLocalMeansDenoising::labMethod.

**See Also:**

`fastNlMeansDenoising()`

## gpu::FastNonLocalMeansDenoising::labMethod()

Modification of `FastNonLocalMeansDenoising::simpleMethod` for color images

**C++:** void gpu::FastNonLocalMeansDenoising::**labMethod**(const GpuMat& **src**, GpuMat& **dst**, float **h_luminance**, float **h_color**, int **search_window**=21, int **block_size**=7, Stream& **s**=Stream::Null())

> Parameters

>> **src** – Input 8-bit 3-channel image.

>> **dst** – Output image with the same size and type as `src` .

>> **h_luminance** – Parameter regulating filter strength. Big h value perfectly removes noise but also removes image details, smaller h value preserves details but also preserves some noise

>> **float** – The same as h but for color components. For most images value equals 10 will be enought to remove colored noise and do not distort colors

>> **search_window** – Size in pixels of the window that is used to compute weighted average for given pixel. Should be odd. Affect performance linearly: greater search_window - greater denoising time. Recommended value 21 pixels

>> **block_size** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels

>> **stream** – Stream for the asynchronous invocations.

The function converts image to CIELAB colorspace and then separately denoise L and AB components with given h parameters using `FastNonLocalMeansDenoising::simpleMethod` function.

**See Also:**

`fastNlMeansDenoisingColored()`

## gpu::alphaComp

Composites two images using alpha opacity values contained in each image.

**C++:** void gpu::**alphaComp**(const GpuMat& **img1**, const GpuMat& **img2**, GpuMat& **dst**, int **alpha_op**, Stream& **stream**=Stream::Null())

> Parameters

>> **img1** – First image. Supports `CV_8UC4` , `CV_16UC4` , `CV_32SC4` and `CV_32FC4` types.

>> **img2** – Second image. Must have the same size and the same type as `img1` .

>> **dst** – Destination image.

>> **alpha_op** – Flag specifying the alpha-blending operation:

>> – **ALPHA_OVER**

>> – **ALPHA_IN**

>> – **ALPHA_OUT**

>> – **ALPHA_ATOP**

>> – **ALPHA_XOR**

>> – **ALPHA_PLUS**

- **ALPHA_OVER_PREMUL**

- **ALPHA_IN_PREMUL**

- **ALPHA_OUT_PREMUL**

- **ALPHA_ATOP_PREMUL**

- **ALPHA_XOR_PREMUL**

- **ALPHA_PLUS_PREMUL**

- **ALPHA_PREMUL**

**stream** – Stream for the asynchronous version.

## gpu::Canny

Finds edges in an image using the [Canny86] algorithm.

**C++:** void gpu::**Canny**(const GpuMat& **image**, GpuMat& **edges**, double **low_thresh**, double **high_thresh**, int **apperture_size**=3, bool **L2gradient**=false)

**C++:** void gpu::**Canny**(const GpuMat& **image**, CannyBuf& **buf**, GpuMat& **edges**, double **low_thresh**, double **high_thresh**, int **apperture_size**=3, bool **L2gradient**=false)

**C++:** void gpu::**Canny**(const GpuMat& **dx**, const GpuMat& **dy**, GpuMat& **edges**, double **low_thresh**, double **high_thresh**, bool **L2gradient**=false)

**C++:** void gpu::**Canny**(const GpuMat& **dx**, const GpuMat& **dy**, CannyBuf& **buf**, GpuMat& **edges**, double **low_thresh**, double **high_thresh**, bool **L2gradient**=false)

Parameters

**image** – Single-channel 8-bit input image.

**dx** – First derivative of image in the vertical direction. Support only `CV_32S` type.

**dy** – First derivative of image in the horizontal direction. Support only `CV_32S` type.

**edges** – Output edge map. It has the same size and type as `image` .

**low_thresh** – First threshold for the hysteresis procedure.

**high_thresh** – Second threshold for the hysteresis procedure.

**apperture_size** – Aperture size for the `Sobel()` operator.

**L2gradient** – Flag indicating whether a more accurate $L_2$ norm $= \sqrt{(dI/dx)^2 + (dI/dy)^2}$ should be used to compute the image gradient magnitude ( `L2gradient=true` ), or a faster default $L_1$ norm $= |dI/dx| + |dI/dy|$ is enough ( `L2gradient=false` ).

**buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).

**See Also:**

`Canny()`

## gpu::HoughLines

Finds lines in a binary image using the classical Hough transform.

**C++:** void gpu::**HoughLines**(const GpuMat& **src**, GpuMat& **lines**, float **rho**, float **theta**, int **threshold**, bool **doSort**=false, int **maxLines**=4096)

**C++:** void gpu::**HoughLines**(const GpuMat& **src**, GpuMat& **lines**, HoughLinesBuf& **buf**, float **rho**, float **theta**, int **threshold**, bool **doSort**=false, int **maxLines**=4096)

> **Parameters**

>> **src** – 8-bit, single-channel binary source image.

>> **lines** – Output vector of lines. Each line is represented by a two-element vector $(\rho, \theta)$ . $\rho$ is the distance from the coordinate origin $(0, 0)$ (top-left corner of the image). $\theta$ is the line rotation angle in radians ( $0 \sim$ vertical line, $\pi/2 \sim$ horizontal line ).

>> **rho** – Distance resolution of the accumulator in pixels.

>> **theta** – Angle resolution of the accumulator in radians.

>> **threshold** – Accumulator threshold parameter. Only those lines are returned that get enough votes ( $>$ `threshold` ).

>> **doSort** – Performs lines sort by votes.

>> **maxLines** – Maximum number of output lines.

>> **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).

**See Also:**

`HoughLines()`

## gpu::HoughLinesDownload

Downloads results from `gpu::HoughLines()` to host memory.

**C++:** void gpu::**HoughLinesDownload**(const GpuMat& **d_lines**, OutputArray **h_lines**, OutputArray **h_votes**=noArray())

> **Parameters**

>> **d_lines** – Result of `gpu::HoughLines()` .

>> **h_lines** – Output host array.

>> **h_votes** – Optional output array for line's votes.

**See Also:**

`gpu::HoughLines()`

## gpu::HoughCircles

Finds circles in a grayscale image using the Hough transform.

**C++:** void gpu::**HoughCircles**(const GpuMat& **src**, GpuMat& **circles**, int **method**, float **dp**, float **minDist**, int **cannyThreshold**, int **votesThreshold**, int **minRadius**, int **maxRadius**, int **maxCircles**=4096)

**C++:** void gpu::**HoughCircles**(const GpuMat& **src**, GpuMat& **circles**, HoughCirclesBuf& **buf**, int **method**, float **dp**, float **minDist**, int **cannyThreshold**, int **votesThreshold**, int **min-Radius**, int **maxRadius**, int **maxCircles**=4096)

> **Parameters**

>> **src** – 8-bit, single-channel grayscale input image.

>> **circles** – Output vector of found circles. Each vector is encoded as a 3-element floating-point vector $(x, y, \texttt{radius})$ .

> **method** – Detection method to use. Currently, the only implemented method is `CV_HOUGH_GRADIENT`, which is basically *21HT*, described in [Yuen90].
>
> **dp** – Inverse ratio of the accumulator resolution to the image resolution. For example, if `dp=1`, the accumulator has the same resolution as the input image. If `dp=2`, the accumulator has half as big width and height.
>
> **minDist** – Minimum distance between the centers of the detected circles. If the parameter is too small, multiple neighbor circles may be falsely detected in addition to a true one. If it is too large, some circles may be missed.
>
> **cannyThreshold** – The higher threshold of the two passed to the `gpu::Canny()` edge detector (the lower one is twice smaller).
>
> **votesThreshold** – The accumulator threshold for the circle centers at the detection stage. The smaller it is, the more false circles may be detected.
>
> **minRadius** – Minimum circle radius.
>
> **maxRadius** – Maximum circle radius.
>
> **maxCircles** – Maximum number of output circles.
>
> **buf** – Optional buffer to avoid extra memory allocations (for many calls with the same sizes).

**See Also:**

`HoughCircles()`

## gpu::HoughCirclesDownload

Downloads results from `gpu::HoughCircles()` to host memory.

**C++:** void gpu::**HoughCirclesDownload**(const GpuMat& **d_circles**, OutputArray **h_circles**)

> **Parameters**
>
> > **d_circles** – Result of `gpu::HoughCircles()`.
> >
> > **h_circles** – Output host array.

**See Also:**

`gpu::HoughCircles()`

# 11.7 Matrix Reductions

## gpu::meanStdDev

Computes a mean value and a standard deviation of matrix elements.

**C++:** void gpu::**meanStdDev**(const GpuMat& **mtx**, Scalar& **mean**, Scalar& **stddev**)

**C++:** void gpu::**meanStdDev**(const GpuMat& **mtx**, Scalar& **mean**, Scalar& **stddev**, GpuMat& **buf**)

> **Parameters**
>
> > **mtx** – Source matrix. `CV_8UC1` matrices are supported for now.
> >
> > **mean** – Mean value.
> >
> > **stddev** – Standard deviation value.

> **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

**See Also:**

meanStdDev()

## gpu::norm

Returns the norm of a matrix (or difference of two matrices).

**C++:** double gpu::**norm**(const GpuMat& **src1**, int **normType**=NORM_L2)

**C++:** double gpu::**norm**(const GpuMat& **src1**, int **normType**, GpuMat& **buf**)

**C++:** double gpu::**norm**(const GpuMat& **src1**, const GpuMat& **src2**, int **normType**=NORM_L2)

> **Parameters**
>
> > **src1** – Source matrix. Any matrices except 64F are supported.
> >
> > **src2** – Second source matrix (if any) with the same size and type as src1.
> >
> > **normType** – Norm type. NORM_L1 , NORM_L2 , and NORM_INF are supported for now.
> >
> > **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

**See Also:**

norm()

## gpu::sum

Returns the sum of matrix elements.

**C++:** Scalar gpu::**sum**(const GpuMat& **src**)

**C++:** Scalar gpu::**sum**(const GpuMat& **src**, GpuMat& **buf**)

> **Parameters**
>
> > **src** – Source image of any depth except for CV_64F .
> >
> > **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

**See Also:**

sum()

## gpu::absSum

Returns the sum of absolute values for matrix elements.

**C++:** Scalar gpu::**absSum**(const GpuMat& **src**)

**C++:** Scalar gpu::**absSum**(const GpuMat& **src**, GpuMat& **buf**)

> **Parameters**
>
> > **src** – Source image of any depth except for CV_64F .
> >
> > **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

## gpu::sqrSum

Returns the squared sum of matrix elements.

**C++:** Scalar gpu::**sqrSum**(const GpuMat& **src**)

**C++:** Scalar gpu::**sqrSum**(const GpuMat& **src**, GpuMat& **buf**)

> **Parameters**

>> **src** – Source image of any depth except for `CV_64F` .

>> **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

## gpu::minMax

Finds global minimum and maximum matrix elements and returns their values.

**C++:** void gpu::**minMax**(const GpuMat& **src**, double* **minVal**, double* **maxVal**=0, const GpuMat& **mask**=GpuMat())

**C++:** void gpu::**minMax**(const GpuMat& **src**, double* **minVal**, double* **maxVal**, const GpuMat& **mask**, GpuMat& **buf**)

> **Parameters**

>> **src** – Single-channel source image.

>> **minVal** – Pointer to the returned minimum value. Use `NULL` if not required.

>> **maxVal** – Pointer to the returned maximum value. Use `NULL` if not required.

>> **mask** – Optional mask to select a sub-matrix.

>> **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

The function does not work with `CV_64F` images on GPUs with the compute capability < 1.3.

**See Also:**

minMaxLoc()

## gpu::minMaxLoc

Finds global minimum and maximum matrix elements and returns their values with locations.

**C++:** void gpu::**minMaxLoc**(const GpuMat& **src**, double* **minVal**, double* **maxVal**=0, Point* **minLoc**=0, Point* **maxLoc**=0, const GpuMat& **mask**=GpuMat())

**C++:** void gpu::**minMaxLoc**(const GpuMat& **src**, double* **minVal**, double* **maxVal**, Point* **minLoc**, Point* **maxLoc**, const GpuMat& **mask**, GpuMat& **valbuf**, GpuMat& **locbuf**)

> **Parameters**

>> **src** – Single-channel source image.

>> **minVal** – Pointer to the returned minimum value. Use `NULL` if not required.

>> **maxVal** – Pointer to the returned maximum value. Use `NULL` if not required.

>> **minLoc** – Pointer to the returned minimum location. Use `NULL` if not required.

>> **maxLoc** – Pointer to the returned maximum location. Use `NULL` if not required.

>> **mask** – Optional mask to select a sub-matrix.

> **valbuf** – Optional values buffer to avoid extra memory allocations. It is resized automatically.

> **locbuf** – Optional locations buffer to avoid extra memory allocations. It is resized automatically.

The function does not work with `CV_64F` images on GPU with the compute capability < 1.3.

**See Also:**

`minMaxLoc()`

## gpu::countNonZero

Counts non-zero matrix elements.

**C++:** int gpu::**countNonZero**(const GpuMat& **src**)

**C++:** int gpu::**countNonZero**(const GpuMat& **src**, GpuMat& **buf**)

> **Parameters**

>> **src** – Single-channel source image.

>> **buf** – Optional buffer to avoid extra memory allocations. It is resized automatically.

The function does not work with `CV_64F` images on GPUs with the compute capability < 1.3.

**See Also:**

`countNonZero()`

## gpu::reduce

Reduces a matrix to a vector.

**C++:** void gpu::**reduce**(const GpuMat& **mtx**, GpuMat& **vec**, int **dim**, int **reduceOp**, int **dtype**=-1, Stream& **stream**=Stream::Null())

> **Parameters**

>> **mtx** – Source 2D matrix.

>> **vec** – Destination vector. Its size and type is defined by `dim` and `dtype` parameters.

>> **dim** – Dimension index along which the matrix is reduced. 0 means that the matrix is reduced to a single row. 1 means that the matrix is reduced to a single column.

>> **reduceOp** – Reduction operation that could be one of the following:

>> – **CV_REDUCE_SUM** The output is the sum of all rows/columns of the matrix.

>> – **CV_REDUCE_AVG** The output is the mean vector of all rows/columns of the matrix.

>> – **CV_REDUCE_MAX** The output is the maximum (column/row-wise) of all rows/columns of the matrix.

>> – **CV_REDUCE_MIN** The output is the minimum (column/row-wise) of all rows/columns of the matrix.

>> **dtype** – When it is negative, the destination vector will have the same type as the source matrix. Otherwise, its type will be `CV_MAKE_TYPE(CV_MAT_DEPTH(dtype), mtx.channels())`.

The function reduce reduces the matrix to a vector by treating the matrix rows/columns as a set of 1D vectors and performing the specified operation on the vectors until a single row/column is obtained. For example, the function can be used to compute horizontal and vertical projections of a raster image. In case of CV_REDUCE_SUM and CV_REDUCE_AVG, the output may have a larger element bit-depth to preserve accuracy. And multi-channel arrays are also supported in these two reduction modes.

**See Also:**

reduce()

# 11.8 Object Detection

## gpu::HOGDescriptor

**struct** gpu::**HOGDescriptor**

The class implements Histogram of Oriented Gradients ([Dalal2005]) object detector.

```
struct CV_EXPORTS HOGDescriptor
{
    enum { DEFAULT_WIN_SIGMA = -1 };
    enum { DEFAULT_NLEVELS = 64 };
    enum { DESCR_FORMAT_ROW_BY_ROW, DESCR_FORMAT_COL_BY_COL };

    HOGDescriptor(Size win_size=Size(64, 128), Size block_size=Size(16, 16),
                  Size block_stride=Size(8, 8), Size cell_size=Size(8, 8),
                  int nbins=9, double win_sigma=DEFAULT_WIN_SIGMA,
                  double threshold_L2hys=0.2, bool gamma_correction=true,
                  int nlevels=DEFAULT_NLEVELS);

    size_t getDescriptorSize() const;
    size_t getBlockHistogramSize() const;

    void setSVMDetector(const vector<float>& detector);

    static vector<float> getDefaultPeopleDetector();
    static vector<float> getPeopleDetector48x96();
    static vector<float> getPeopleDetector64x128();

    void detect(const GpuMat& img, vector<Point>& found_locations,
                double hit_threshold=0, Size win_stride=Size(),
                Size padding=Size());

    void detectMultiScale(const GpuMat& img, vector<Rect>& found_locations,
                          double hit_threshold=0, Size win_stride=Size(),
                          Size padding=Size(), double scale0=1.05,
                          int group_threshold=2);

    void getDescriptors(const GpuMat& img, Size win_stride,
                        GpuMat& descriptors,
                        int descr_format=DESCR_FORMAT_COL_BY_COL);

    Size win_size;
    Size block_size;
    Size block_stride;
    Size cell_size;
    int nbins;
```

```
    double win_sigma;
    double threshold_L2hys;
    bool gamma_correction;
    int nlevels;

private:
    // Hidden
}
```

Interfaces of all methods are kept similar to the `CPU HOG` descriptor and detector analogues as much as possible.

## gpu::HOGDescriptor::HOGDescriptor

Creates the `HOG` descriptor and detector.

**C++:** `gpu::HOGDescriptor::`**`HOGDescriptor`**(Size **win_size**=Size(64, 128), Size **block_size**=Size(16, 16), Size **block_stride**=Size(8, 8), Size **cell_size**=Size(8, 8), int **nbins**=9, double **win_sigma**=DEFAULT_WIN_SIGMA, double **threshold_L2hys**=0.2, bool **gamma_correction**=true, int **nlevels**=DEFAULT_NLEVELS)

> Parameters
>
> > **win_size** – Detection window size. Align to block size and block stride.
> >
> > **block_size** – Block size in pixels. Align to cell size. Only (16,16) is supported for now.
> >
> > **block_stride** – Block stride. It must be a multiple of cell size.
> >
> > **cell_size** – Cell size. Only (8, 8) is supported for now.
> >
> > **nbins** – Number of bins. Only 9 bins per cell are supported for now.
> >
> > **win_sigma** – Gaussian smoothing window parameter.
> >
> > **threshold_L2hys** – L2-Hys normalization method shrinkage.
> >
> > **gamma_correction** – Flag to specify whether the gamma correction preprocessing is required or not.
> >
> > **nlevels** – Maximum number of detection window increases.

## gpu::HOGDescriptor::getDescriptorSize

Returns the number of coefficients required for the classification.

**C++:** size_t gpu::HOGDescriptor::**getDescriptorSize**() const

## gpu::HOGDescriptor::getBlockHistogramSize

Returns the block histogram size.

**C++:** size_t gpu::HOGDescriptor::**getBlockHistogramSize**() const

## gpu::HOGDescriptor::setSVMDetector

Sets coefficients for the linear SVM classifier.

**C++:** void gpu::HOGDescriptor::**setSVMDetector**(const vector<float>& **detector**)

## gpu::HOGDescriptor::getDefaultPeopleDetector

Returns coefficients of the classifier trained for people detection (for default window size).

**C++:** **static** vector<float> gpu::HOGDescriptor::**getDefaultPeopleDetector**()

## gpu::HOGDescriptor::getPeopleDetector48x96

Returns coefficients of the classifier trained for people detection (for 48x96 windows).

**C++:** **static** vector<float> gpu::HOGDescriptor::**getPeopleDetector48x96**()

## gpu::HOGDescriptor::getPeopleDetector64x128

Returns coefficients of the classifier trained for people detection (for 64x128 windows).

**C++:** **static** vector<float> gpu::HOGDescriptor::**getPeopleDetector64x128**()

## gpu::HOGDescriptor::detect

Performs object detection without a multi-scale window.

**C++:** void gpu::HOGDescriptor::**detect**(const GpuMat& **img**, vector<Point>& **found_locations**, double **hit_threshold**=0, Size **win_stride**=Size(), Size **padding**=Size())

>    **Parameters**

>    >    **img** – Source image. `CV_8UC1` and `CV_8UC4` types are supported for now.

>    >    **found_locations** – Left-top corner points of detected objects boundaries.

>    >    **hit_threshold** – Threshold for the distance between features and SVM classifying plane. Usually it is 0 and should be specfied in the detector coefficients (as the last free coefficient). But if the free coefficient is omitted (which is allowed), you can specify it manually here.

>    >    **win_stride** – Window stride. It must be a multiple of block stride.

>    >    **padding** – Mock parameter to keep the CPU interface compatibility. It must be (0,0).

## gpu::HOGDescriptor::detectMultiScale

Performs object detection with a multi-scale window.

**C++:** void gpu::HOGDescriptor::**detectMultiScale**(const GpuMat& **img**, vector<Rect>& **found_locations**, double **hit_threshold**=0, Size **win_stride**=Size(), Size **padding**=Size(), double **scale0**=1.05, int **group_threshold**=2)

>    **Parameters**

>    >    **img** – Source image. See `gpu::HOGDescriptor::detect()` for type limitations.

> **found_locations** – Detected objects boundaries.
>
> **hit_threshold** – Threshold for the distance between features and SVM classifying plane. See `gpu::HOGDescriptor::detect()` for details.
>
> **win_stride** – Window stride. It must be a multiple of block stride.
>
> **padding** – Mock parameter to keep the CPU interface compatibility. It must be (0,0).
>
> **scale0** – Coefficient of the detection window increase.
>
> **group_threshold** – Coefficient to regulate the similarity threshold. When detected, some objects can be covered by many rectangles. 0 means not to perform grouping. See `groupRectangles()`.

## gpu::HOGDescriptor::getDescriptors

Returns block descriptors computed for the whole image.

**C++:** void gpu::HOGDescriptor::**getDescriptors**(const GpuMat& **img**, Size **win_stride**, GpuMat& **descriptors**, int **descr_format**=DESCR_FORMAT_COL_BY_COL)

> **Parameters**
>
> > **img** – Source image. See `gpu::HOGDescriptor::detect()` for type limitations.
> >
> > **win_stride** – Window stride. It must be a multiple of block stride.
> >
> > **descriptors** – 2D array of descriptors.
> >
> > **descr_format** – Descriptor storage format:
> >
> > – **DESCR_FORMAT_ROW_BY_ROW** - Row-major order.
> >
> > – **DESCR_FORMAT_COL_BY_COL** - Column-major order.

The function is mainly used to learn the classifier.

## gpu::CascadeClassifier_GPU

**class** gpu::**CascadeClassifier_GPU**

Cascade classifier class used for object detection. Supports HAAR and LBP cascades.

```
class CV_EXPORTS CascadeClassifier_GPU
{
public:
        CascadeClassifier_GPU();
        CascadeClassifier_GPU(const string& filename);
        ~CascadeClassifier_GPU();

        bool empty() const;
        bool load(const string& filename);
        void release();

        /* Returns number of detected objects */
        int detectMultiScale( const GpuMat& image, GpuMat& objectsBuf, double scaleFactor=1.2, int minNeighbors=4, Siz
        int detectMultiScale( const GpuMat& image, GpuMat& objectsBuf, Size maxObjectSize, Size minSize = Size(), doub

        /* Finds only the largest object. Special mode if training is required.*/
```

```
        bool findLargestObject;

        /* Draws rectangles in input image */
        bool visualizeInPlace;

        Size getClassifierSize() const;
};
```

## gpu::CascadeClassifier_GPU::CascadeClassifier_GPU

Loads the classifier from a file. Cascade type is detected automatically by constructor parameter.

**C++:** `gpu::CascadeClassifier_GPU::`**`CascadeClassifier_GPU`**`(const string&` **filename**`)`

> **Parameters**
>
> > **filename** – Name of the file from which the classifier is loaded. Only the old `haar` classifier (trained by the `haar` training application) and NVIDIA's `nvbin` are supported for HAAR and only new type of OpenCV XML cascade supported for LBP.

## gpu::CascadeClassifier_GPU::empty

Checks whether the classifier is loaded or not.

**C++:** `bool gpu::CascadeClassifier_GPU::`**`empty`**`() const`

## gpu::CascadeClassifier_GPU::load

Loads the classifier from a file. The previous content is destroyed.

**C++:** `bool gpu::CascadeClassifier_GPU::`**`load`**`(const string&` **filename**`)`

> **Parameters**
>
> > **filename** – Name of the file from which the classifier is loaded. Only the old `haar` classifier (trained by the `haar` training application) and NVIDIA's `nvbin` are supported for HAAR and only new type of OpenCV XML cascade supported for LBP.

## gpu::CascadeClassifier_GPU::release

Destroys the loaded classifier.

**C++:** `void gpu::CascadeClassifier_GPU::`**`release`**`()`

## gpu::CascadeClassifier_GPU::detectMultiScale

Detects objects of different sizes in the input image.

**C++:** `int gpu::CascadeClassifier_GPU::`**`detectMultiScale`**`(const GpuMat&` **image**`, GpuMat&` **objectsBuf**`, double` **scaleFactor**`=1.1, int` **minNeighbors**`=4, Size` **minSize**`=Size() )`

> **Parameters**
>
> > **image** – Matrix of type `CV_8U` containing an image where objects should be detected.

> **objectsBuf** – Buffer to store detected objects (rectangles). If it is empty, it is allocated with the default size. If not empty, the function searches not more than N objects, where `N = sizeof(objectsBufer's data)/sizeof(cv::Rect)`.
>
> **maxObjectSize** – Maximum possible object size. Objects larger than that are ignored. Used for second signature and supported only for LBP cascades.
>
> **scaleFactor** – Parameter specifying how much the image size is reduced at each image scale.
>
> **minNeighbors** – Parameter specifying how many neighbors each candidate rectangle should have to retain it.
>
> **minSize** – Minimum possible object size. Objects smaller than that are ignored.

The detected objects are returned as a list of rectangles.

The function returns the number of detected objects, so you can retrieve them as in the following example:

```
gpu::CascadeClassifier_GPU cascade_gpu(...);

Mat image_cpu = imread(...)
GpuMat image_gpu(image_cpu);

GpuMat objbuf;
int detections_number = cascade_gpu.detectMultiScale( image_gpu,
          objbuf, 1.2, minNeighbors);

Mat obj_host;
// download only detected number of rectangles
objbuf.colRange(0, detections_number).download(obj_host);

Rect* faces = obj_host.ptr<Rect>();
for(int i = 0; i < detections_num; ++i)
   cv::rectangle(image_cpu, faces[i], Scalar(255));

imshow("Faces", image_cpu);
```

**See Also:**

`CascadeClassifier::detectMultiScale()`

# 11.9 Feature Detection and Description

## gpu::SURF_GPU

**class** gpu::**SURF_GPU**

Class used for extracting Speeded Up Robust Features (SURF) from an image.

```
class SURF_GPU
{
public:
    enum KeypointLayout
    {
        X_ROW = 0,
        Y_ROW,
        LAPLACIAN_ROW,
        OCTAVE_ROW,
```

```
        SIZE_ROW,
        ANGLE_ROW,
        HESSIAN_ROW,
        ROWS_COUNT
    };

    //! the default constructor
    SURF_GPU();
    //! the full constructor taking all the necessary parameters
    explicit SURF_GPU(double _hessianThreshold, int _nOctaves=4,
        int _nOctaveLayers=2, bool _extended=false, float _keypointsRatio=0.01f);

    //! returns the descriptor size in float's (64 or 128)
    int descriptorSize() const;

    //! upload host keypoints to device memory
    void uploadKeypoints(const vector<KeyPoint>& keypoints,
        GpuMat& keypointsGPU);
    //! download keypoints from device to host memory
    void downloadKeypoints(const GpuMat& keypointsGPU,
        vector<KeyPoint>& keypoints);

    //! download descriptors from device to host memory
    void downloadDescriptors(const GpuMat& descriptorsGPU,
        vector<float>& descriptors);

    void operator()(const GpuMat& img, const GpuMat& mask,
        GpuMat& keypoints);

    void operator()(const GpuMat& img, const GpuMat& mask,
        GpuMat& keypoints, GpuMat& descriptors,
        bool useProvidedKeypoints = false,
        bool calcOrientation = true);

    void operator()(const GpuMat& img, const GpuMat& mask,
        std::vector<KeyPoint>& keypoints);

    void operator()(const GpuMat& img, const GpuMat& mask,
        std::vector<KeyPoint>& keypoints, GpuMat& descriptors,
        bool useProvidedKeypoints = false,
        bool calcOrientation = true);

    void operator()(const GpuMat& img, const GpuMat& mask,
        std::vector<KeyPoint>& keypoints,
        std::vector<float>& descriptors,
        bool useProvidedKeypoints = false,
        bool calcOrientation = true);

    void releaseMemory();

    // SURF parameters
    double hessianThreshold;
    int nOctaves;
    int nOctaveLayers;
    bool extended;
    bool upright;

    //! max keypoints = keypointsRatio * img.size().area()
```

```
    float keypointsRatio;

    GpuMat sum, mask1, maskSum, intBuffer;

    GpuMat det, trace;

    GpuMat maxPosBuffer;
};
```

The class SURF_GPU implements Speeded Up Robust Features descriptor. There is a fast multi-scale Hessian keypoint detector that can be used to find the keypoints (which is the default option). But the descriptors can also be computed for the user-specified keypoints. Only 8-bit grayscale images are supported.

The class SURF_GPU can store results in the GPU and CPU memory. It provides functions to convert results between CPU and GPU version ( uploadKeypoints, downloadKeypoints, downloadDescriptors ). The format of CPU results is the same as SURF results. GPU results are stored in GpuMat. The keypoints matrix is $nFeatures \times 7$ matrix with the CV_32FC1 type.

- keypoints.ptr<float>(X_ROW)[i] contains x coordinate of the i-th feature.

- keypoints.ptr<float>(Y_ROW)[i] contains y coordinate of the i-th feature.

- keypoints.ptr<float>(LAPLACIAN_ROW)[i] contains the laplacian sign of the i-th feature.

- keypoints.ptr<float>(OCTAVE_ROW)[i] contains the octave of the i-th feature.

- keypoints.ptr<float>(SIZE_ROW)[i] contains the size of the i-th feature.

- keypoints.ptr<float>(ANGLE_ROW)[i] contain orientation of the i-th feature.

- keypoints.ptr<float>(HESSIAN_ROW)[i] contains the response of the i-th feature.

The descriptors matrix is $nFeatures \times descriptorSize$ matrix with the CV_32FC1 type.

The class SURF_GPU uses some buffers and provides access to it. All buffers can be safely released between function calls.

**See Also:**

SURF

# gpu::FAST_GPU

class gpu::**FAST_GPU**

Class used for corner detection using the FAST algorithm.

```
class FAST_GPU
{
public:
    enum
    {
        LOCATION_ROW = 0,
        RESPONSE_ROW,
        ROWS_COUNT
    };

    // all features have same size
    static const int FEATURE_SIZE = 7;

    explicit FAST_GPU(int threshold, bool nonmaxSupression = true,
```

```
                        double keypointsRatio = 0.05);

    void operator ()(const GpuMat& image, const GpuMat& mask, GpuMat& keypoints);
    void operator ()(const GpuMat& image, const GpuMat& mask,
                     std::vector<KeyPoint>& keypoints);

    void downloadKeypoints(const GpuMat& d_keypoints,
                           std::vector<KeyPoint>& keypoints);

    void convertKeypoints(const Mat& h_keypoints,
                          std::vector<KeyPoint>& keypoints);

    void release();

    bool nonmaxSupression;

    int threshold;

    double keypointsRatio;

    int calcKeyPointsLocation(const GpuMat& image, const GpuMat& mask);

    int getKeyPoints(GpuMat& keypoints);
};
```

The class FAST_GPU implements FAST corner detection algorithm.

**See Also:**

FAST()

## gpu::FAST_GPU::FAST_GPU

Constructor.

**C++:** gpu::FAST_GPU::**FAST_GPU**(int **threshold**, bool **nonmaxSupression**=true, double **keypointsRatio**=0.05)

>       **Parameters**
>
>>           **threshold** – Threshold on difference between intensity of the central pixel and pixels on a circle around this pixel.
>>
>>           **nonmaxSupression** – If it is true, non-maximum suppression is applied to detected corners (keypoints).
>>
>>           **keypointsRatio** – Inner buffer size for keypoints store is determined as (keypointsRatio * image_width * image_height).

## gpu::FAST_GPU::operator ()

Finds the keypoints using FAST detector.

**C++:** void gpu::FAST_GPU::**operator()**(const GpuMat& **image**, const GpuMat& **mask**, GpuMat& **keypoints**)

**C++:** void gpu::FAST_GPU::**operator()**(const GpuMat& **image**, const GpuMat& **mask**, std::vector<KeyPoint>& **keypoints**)

>       **Parameters**

> **image** – Image where keypoints (corners) are detected. Only 8-bit grayscale images are supported.
>
> **mask** – Optional input mask that marks the regions where we should detect features.
>
> **keypoints** – The output vector of keypoints. Can be stored both in CPU and GPU memory. For GPU memory:
>
> – keypoints.ptr<Vec2s>(LOCATION_ROW)[i] will contain location of i'th point
>
> – keypoints.ptr<float>(RESPONSE_ROW)[i] will contain response of i'th point (if non-maximum suppression is applied)

## gpu::FAST_GPU::downloadKeypoints

Download keypoints from GPU to CPU memory.

**C++:** void gpu::FAST_GPU::**downloadKeypoints**(const GpuMat& **d_keypoints**, std::vector<KeyPoint>& **keypoints**)

## gpu::FAST_GPU::convertKeypoints

Converts keypoints from GPU representation to vector of `KeyPoint`.

**C++:** void gpu::FAST_GPU::**convertKeypoints**(const Mat& **h_keypoints**, std::vector<KeyPoint>& **keypoints**)

## gpu::FAST_GPU::release

Releases inner buffer memory.

**C++:** void gpu::FAST_GPU::**release**()

## gpu::FAST_GPU::calcKeyPointsLocation

Find keypoints and compute it's response if `nonmaxSupression` is true.

**C++:** int gpu::FAST_GPU::**calcKeyPointsLocation**(const GpuMat& **image**, const GpuMat& **mask**)

>   **Parameters**
>
>> **image** – Image where keypoints (corners) are detected. Only 8-bit grayscale images are supported.
>>
>> **mask** – Optional input mask that marks the regions where we should detect features.

The function returns count of detected keypoints.

## gpu::FAST_GPU::getKeyPoints

Gets final array of keypoints.

**C++:** int gpu::FAST_GPU::**getKeyPoints**(GpuMat& **keypoints**)

>   **Parameters**
>
>> **keypoints** – The output vector of keypoints.

The function performs non-max suppression if needed and returns final count of keypoints.

## gpu::ORB_GPU

**class** gpu::**ORB_GPU**

Class for extracting ORB features and descriptors from an image.

```cpp
class ORB_GPU
{
public:
    enum
    {
        X_ROW = 0,
        Y_ROW,
        RESPONSE_ROW,
        ANGLE_ROW,
        OCTAVE_ROW,
        SIZE_ROW,
        ROWS_COUNT
    };

    enum
    {
        DEFAULT_FAST_THRESHOLD = 20
    };

    explicit ORB_GPU(int nFeatures = 500, float scaleFactor = 1.2f,
                     int nLevels = 8, int edgeThreshold = 31,
                     int firstLevel = 0, int WTA_K = 2,
                     int scoreType = 0, int patchSize = 31);

    void operator()(const GpuMat& image, const GpuMat& mask,
                    std::vector<KeyPoint>& keypoints);
    void operator()(const GpuMat& image, const GpuMat& mask, GpuMat& keypoints);

    void operator()(const GpuMat& image, const GpuMat& mask,
                    std::vector<KeyPoint>& keypoints, GpuMat& descriptors);
    void operator()(const GpuMat& image, const GpuMat& mask,
                    GpuMat& keypoints, GpuMat& descriptors);

    void downloadKeyPoints(GpuMat& d_keypoints, std::vector<KeyPoint>& keypoints);

    void convertKeyPoints(Mat& d_keypoints, std::vector<KeyPoint>& keypoints);

    int descriptorSize() const;

    void setParams(size_t n_features, const ORB::CommonParams& detector_params);
    void setFastParams(int threshold, bool nonmaxSupression = true);

    void release();

    bool blurForDescriptor;
};
```

The class implements ORB feature detection and description algorithm.

## gpu::ORB_GPU::ORB_GPU

Constructor.

**C++:** `gpu::ORB_GPU::`**`ORB_GPU`**`(`int **nFeatures**=500, float **scaleFactor**=1.2f, int **nLevels**=8, int **edgeThreshold**=31, int **firstLevel**=0, int **WTA_K**=2, int **scoreType**=0, int **patchSize**=31`)`

>   **Parameters**

>>   **nFeatures** – The number of desired features.

>>   **scaleFactor** – Coefficient by which we divide the dimensions from one scale pyramid level to the next.

>>   **nLevels** – The number of levels in the scale pyramid.

>>   **edgeThreshold** – How far from the boundary the points should be.

>>   **firstLevel** – The level at which the image is given. If 1, that means we will also look at the image *scaleFactor* times bigger.

## gpu::ORB_GPU::operator()

Detects keypoints and computes descriptors for them.

**C++:** void `gpu::ORB_GPU::`**`operator()`**`(`const GpuMat& **image**, const GpuMat& **mask**, std::vector<KeyPoint>& **keypoints**`)`

**C++:** void `gpu::ORB_GPU::`**`operator()`**`(`const GpuMat& **image**, const GpuMat& **mask**, GpuMat& **keypoints**`)`

**C++:** void `gpu::ORB_GPU::`**`operator()`**`(`const GpuMat& **image**, const GpuMat& **mask**, std::vector<KeyPoint>& **keypoints**, GpuMat& **descriptors**`)`

**C++:** void `gpu::ORB_GPU::`**`operator()`**`(`const GpuMat& **image**, const GpuMat& **mask**, GpuMat& **keypoints**, GpuMat& **descriptors**`)`

>   **Parameters**

>>   **image** – Input 8-bit grayscale image.

>>   **mask** – Optional input mask that marks the regions where we should detect features.

>>   **keypoints** – The input/output vector of keypoints. Can be stored both in CPU and GPU memory. For GPU memory:

>>>   – `keypoints.ptr<float>(X_ROW)[i]` contains x coordinate of the i'th feature.

>>>   – `keypoints.ptr<float>(Y_ROW)[i]` contains y coordinate of the i'th feature.

>>>   – `keypoints.ptr<float>(RESPONSE_ROW)[i]` contains the response of the i'th feature.

>>>   – `keypoints.ptr<float>(ANGLE_ROW)[i]` contains orientation of the i'th feature.

>>>   – `keypoints.ptr<float>(OCTAVE_ROW)[i]` contains the octave of the i'th feature.

>>>   – `keypoints.ptr<float>(SIZE_ROW)[i]` contains the size of the i'th feature.

>>   **descriptors** – Computed descriptors. if `blurForDescriptor` is true, image will be blurred before descriptors calculation.

## gpu::ORB_GPU::downloadKeyPoints

Download keypoints from GPU to CPU memory.

**C++:** void gpu::ORB_GPU::**downloadKeyPoints**(GpuMat& **d_keypoints**, std::vector<KeyPoint>& **keypoints**)

## gpu::ORB_GPU::convertKeyPoints

Converts keypoints from GPU representation to vector of KeyPoint.

**C++:** void gpu::ORB_GPU::**convertKeyPoints**(Mat& **d_keypoints**, std::vector<KeyPoint>& **keypoints**)

## gpu::ORB_GPU::release

Releases inner buffer memory.

**C++:** void gpu::ORB_GPU::**release**()

## gpu::BruteForceMatcher_GPU_base

class gpu::**BruteForceMatcher_GPU_base**

Brute-force descriptor matcher. For each descriptor in the first set, this matcher finds the closest descriptor in the second set by trying each one. This descriptor matcher supports masking permissible matches between descriptor sets.

```cpp
class BruteForceMatcher_GPU_base
{
public:
    explicit BruteForceMatcher_GPU_base(int norm = cv::NORM_L2);

    // Add descriptors to train descriptor collection.
    void add(const std::vector<GpuMat>& descCollection);

    // Get train descriptors collection.
    const std::vector<GpuMat>& getTrainDescriptors() const;

    // Clear train descriptors collection.
    void clear();

    // Return true if there are no train descriptors in collection.
    bool empty() const;

    // Return true if the matcher supports mask in match methods.
    bool isMaskSupported() const;

    void matchSingle(const GpuMat& query, const GpuMat& train,
        GpuMat& trainIdx, GpuMat& distance,
        const GpuMat& mask = GpuMat(), Stream& stream = Stream::Null());

    static void matchDownload(const GpuMat& trainIdx,
        const GpuMat& distance, std::vector<DMatch>& matches);
    static void matchConvert(const Mat& trainIdx,
        const Mat& distance, std::vector<DMatch>& matches);

    void match(const GpuMat& query, const GpuMat& train,
```

```
        std::vector<DMatch>& matches, const GpuMat& mask = GpuMat());

    void makeGpuCollection(GpuMat& trainCollection, GpuMat& maskCollection,
        const vector<GpuMat>& masks = std::vector<GpuMat>());

    void matchCollection(const GpuMat& query, const GpuMat& trainCollection,
        GpuMat& trainIdx, GpuMat& imgIdx, GpuMat& distance,
        const GpuMat& maskCollection, Stream& stream = Stream::Null());

    static void matchDownload(const GpuMat& trainIdx, GpuMat& imgIdx,
        const GpuMat& distance, std::vector<DMatch>& matches);
    static void matchConvert(const Mat& trainIdx, const Mat& imgIdx,
        const Mat& distance, std::vector<DMatch>& matches);

    void match(const GpuMat& query, std::vector<DMatch>& matches,
        const std::vector<GpuMat>& masks = std::vector<GpuMat>());

    void knnMatchSingle(const GpuMat& query, const GpuMat& train,
        GpuMat& trainIdx, GpuMat& distance, GpuMat& allDist, int k,
        const GpuMat& mask = GpuMat(), Stream& stream = Stream::Null());

    static void knnMatchDownload(const GpuMat& trainIdx, const GpuMat& distance,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);
    static void knnMatchConvert(const Mat& trainIdx, const Mat& distance,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);

    void knnMatch(const GpuMat& query, const GpuMat& train,
        std::vector< std::vector<DMatch> >& matches, int k,
        const GpuMat& mask = GpuMat(), bool compactResult = false);

    void knnMatch2Collection(const GpuMat& query, const GpuMat& trainCollection,
        GpuMat& trainIdx, GpuMat& imgIdx, GpuMat& distance,
        const GpuMat& maskCollection = GpuMat(), Stream& stream = Stream::Null());

    static void knnMatch2Download(const GpuMat& trainIdx, const GpuMat& imgIdx, const GpuMat& distance,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);
    static void knnMatch2Convert(const Mat& trainIdx, const Mat& imgIdx, const Mat& distance,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);

    void knnMatch(const GpuMat& query, std::vector< std::vector<DMatch> >& matches, int k,
        const std::vector<GpuMat>& masks = std::vector<GpuMat>(),
        bool compactResult = false);

    void radiusMatchSingle(const GpuMat& query, const GpuMat& train,
        GpuMat& trainIdx, GpuMat& distance, GpuMat& nMatches, float maxDistance,
        const GpuMat& mask = GpuMat(), Stream& stream = Stream::Null());

    static void radiusMatchDownload(const GpuMat& trainIdx, const GpuMat& distance, const GpuMat& nMatches,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);
    static void radiusMatchConvert(const Mat& trainIdx, const Mat& distance, const Mat& nMatches,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);

    void radiusMatch(const GpuMat& query, const GpuMat& train,
        std::vector< std::vector<DMatch> >& matches, float maxDistance,
        const GpuMat& mask = GpuMat(), bool compactResult = false);

    void radiusMatchCollection(const GpuMat& query, GpuMat& trainIdx, GpuMat& imgIdx, GpuMat& distance, GpuMat& nMatch
        const std::vector<GpuMat>& masks = std::vector<GpuMat>(), Stream& stream = Stream::Null());
```

```
    static void radiusMatchDownload(const GpuMat& trainIdx, const GpuMat& imgIdx, const GpuMat& distance, const GpuMat&
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);
    static void radiusMatchConvert(const Mat& trainIdx, const Mat& imgIdx, const Mat& distance, const Mat& nMatches,
        std::vector< std::vector<DMatch> >& matches, bool compactResult = false);

    void radiusMatch(const GpuMat& query, std::vector< std::vector<DMatch> >& matches, float maxDistance,
        const std::vector<GpuMat>& masks = std::vector<GpuMat>(), bool compactResult = false);

private:
    std::vector<GpuMat> trainDescCollection;
};
```

The class `BruteForceMatcher_GPU_base` has an interface similar to the class `DescriptorMatcher`. It has two groups of `match` methods: for matching descriptors of one image with another image or with an image set. Also, all functions have an alternative to save results either to the GPU memory or to the CPU memory.

**See Also:**

`DescriptorMatcher`, `BFMatcher`

## gpu::BruteForceMatcher_GPU_base::match

Finds the best match for each descriptor from a query set with train descriptors.

**C++:** void gpu::BruteForceMatcher_GPU_base::**match**(const GpuMat& **query**, const GpuMat& **train**, std::vector<DMatch>& **matches**, const GpuMat& **mask**=GpuMat())

**C++:** void gpu::BruteForceMatcher_GPU_base::**matchSingle**(const GpuMat& **query**, const GpuMat& **train**, GpuMat& **trainIdx**, GpuMat& **distance**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::BruteForceMatcher_GPU_base::**match**(const GpuMat& **query**, std::vector<DMatch>& **matches**, const std::vector<GpuMat>& **masks**=std::vector<GpuMat>())

**C++:** void gpu::BruteForceMatcher_GPU_base::**matchCollection**(const GpuMat& **query**, const GpuMat& **trainCollection**, GpuMat& **trainIdx**, GpuMat& **imgIdx**, GpuMat& **distance**, const GpuMat& **masks**=GpuMat(), Stream& **stream**=Stream::Null() )

**See Also:**

`DescriptorMatcher::match()`

## gpu::BruteForceMatcher_GPU_base::makeGpuCollection

Performs a GPU collection of train descriptors and masks in a suitable format for the `gpu::BruteForceMatcher_GPU_base::matchCollection()` function.

**C++:** void gpu::BruteForceMatcher_GPU_base::**makeGpuCollection**(GpuMat& **trainCollection**, GpuMat& **maskCollection**, const vector<GpuMat>& **masks**=std::vector<GpuMat>())

---

## gpu::BruteForceMatcher_GPU_base::matchDownload

Downloads matrices obtained via `gpu::BruteForceMatcher_GPU_base::matchSingle()` or `gpu::BruteForceMatcher_GPU_base::matchCollection()` to vector with `DMatch`.

**C++:** **static** void gpu::BruteForceMatcher_GPU_base::**matchDownload**(const GpuMat& **trainIdx**, const GpuMat& **distance**, std::vector<DMatch>& **matches**)

**C++:** **static** void gpu::BruteForceMatcher_GPU_base::**matchDownload**(const GpuMat& **trainIdx**, const GpuMat& **imgIdx**, const GpuMat& **distance**, std::vector<DMatch>& **matches**)

## gpu::BruteForceMatcher_GPU_base::matchConvert

Converts matrices obtained via `gpu::BruteForceMatcher_GPU_base::matchSingle()` or `gpu::BruteForceMatcher_GPU_base::matchCollection()` to vector with `DMatch`.

**C++:** void gpu::BruteForceMatcher_GPU_base::**matchConvert**(const Mat& **trainIdx**, const Mat& **distance**, std::vector<DMatch>& **matches**)

**C++:** void gpu::BruteForceMatcher_GPU_base::**matchConvert**(const Mat& **trainIdx**, const Mat& **imgIdx**, const Mat& **distance**, std::vector<DMatch>& **matches**)

## gpu::BruteForceMatcher_GPU_base::knnMatch

Finds the k best matches for each descriptor from a query set with train descriptors.

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatch**(const GpuMat& **query**, const GpuMat& **train**, std::vector<std::vector<DMatch>>& **matches**, int **k**, const GpuMat& **mask**=GpuMat(), bool **compactResult**=false )

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatchSingle**(const GpuMat& **query**, const GpuMat& **train**, GpuMat& **trainIdx**, GpuMat& **distance**, GpuMat& **allDist**, int **k**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatch**(const GpuMat& **query**, std::vector<std::vector<DMatch>>& **matches**, int **k**, const std::vector<GpuMat>& **masks**=std::vector<GpuMat>(), bool **compactResult**=false )

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatch2Collection**(const GpuMat& **query**, const GpuMat& **trainCollection**, GpuMat& **trainIdx**, GpuMat& **imgIdx**, GpuMat& **distance**, const GpuMat& **maskCollection**=GpuMat(), Stream& **stream**=Stream::Null())

> **Parameters**
>> **query** – Query set of descriptors.
>>
>> **train** – Training set of descriptors. It is not be added to train descriptors collection stored in the class object.
>>
>> **k** – Number of the best matches per each query descriptor (or less if it is not possible).
>>
>> **mask** – Mask specifying permissible matches between the input query and train matrices of descriptors.
>>
>> **compactResult** – If `compactResult` is `true`, the `matches` vector does not contain matches for fully masked-out query descriptors.
>>
>> **stream** – Stream for the asynchronous version.

The function returns detected k (or less if not possible) matches in the increasing order by distance.

The third variant of the method stores the results in GPU memory.

**See Also:**

`DescriptorMatcher::knnMatch()`

## gpu::BruteForceMatcher_GPU_base::knnMatchDownload

Downloads matrices obtained via `gpu::BruteForceMatcher_GPU_base::knnMatchSingle()` or `gpu::BruteForceMatcher_GPU_base::knnMatch2Collection()` to vector with `DMatch`.

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatchDownload**(const GpuMat& **trainIdx**, const GpuMat& **distance**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false)

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatch2Download**(const GpuMat& **trainIdx**, const GpuMat& **imgIdx**, const GpuMat& **distance**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false)

If `compactResult` is `true`, the `matches` vector does not contain matches for fully masked-out query descriptors.

## gpu::BruteForceMatcher_GPU_base::knnMatchConvert

Converts matrices obtained via `gpu::BruteForceMatcher_GPU_base::knnMatchSingle()` or `gpu::BruteForceMatcher_GPU_base::knnMatch2Collection()` to CPU vector with `DMatch`.

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatchConvert**(const Mat& **trainIdx**, const Mat& **distance**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false)

**C++:** void gpu::BruteForceMatcher_GPU_base::**knnMatch2Convert**(const Mat& **trainIdx**, const Mat& **imgIdx**, const Mat& **distance**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false)

If compactResult is true , the matches vector does not contain matches for fully masked-out query descriptors.

## gpu::BruteForceMatcher_GPU_base::radiusMatch

For each query descriptor, finds the best matches with a distance less than a given threshold.

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatch**(const GpuMat& **query**, const GpuMat& **train**, std::vector<std::vector<DMatch>>& **matches**, float **maxDistance**, const GpuMat& **mask**=GpuMat(), bool **compactResult**=false)

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchSingle**(const GpuMat& **query**, const GpuMat& **train**, GpuMat& **trainIdx**, GpuMat& **distance**, GpuMat& **nMatches**, float **maxDistance**, const GpuMat& **mask**=GpuMat(), Stream& **stream**=Stream::Null())

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatch**(const GpuMat& **query**, std::vector<std::vector<DMatch>>& **matches**, float **maxDistance**, const std::vector<GpuMat>& **masks**=std::vector<GpuMat>(), bool **compactResult**=false)

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchCollection**(const GpuMat& **query**, GpuMat& **trainIdx**, GpuMat& **imgIdx**, GpuMat& **distance**, GpuMat& **nMatches**, float **maxDistance**, const std::vector<GpuMat>& **masks**=std::vector<GpuMat>(), Stream& **stream**=Stream::Null())

Parameters

**query** – Query set of descriptors.

**train** – Training set of descriptors. It is not added to train descriptors collection stored in the class object.

**maxDistance** – Distance threshold.

> **mask** – Mask specifying permissible matches between the input query and train matrices of descriptors.

> **compactResult** – If `compactResult` is `true` , the `matches` vector does not contain matches for fully masked-out query descriptors.

> **stream** – Stream for the asynchronous version.

The function returns detected matches in the increasing order by distance.

The methods work only on devices with the compute capability $>= 1.1$.

The third variant of the method stores the results in GPU memory and does not store the points by the distance.

**See Also:**

`DescriptorMatcher::radiusMatch()`

## gpu::BruteForceMatcher_GPU_base::radiusMatchDownload

Downloads matrices obtained via `gpu::BruteForceMatcher_GPU_base::radiusMatchSingle()` or `gpu::BruteForceMatcher_GPU_base::radiusMatchCollection()` to vector with `DMatch`.

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchDownload**(const GpuMat& **trainIdx**, const GpuMat& **distance**, const GpuMat& **nMatches**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false )

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchDownload**(const GpuMat& **trainIdx**, const GpuMat& **imgIdx**, const GpuMat& **distance**, const GpuMat& **nMatches**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false )

If `compactResult` is `true` , the `matches` vector does not contain matches for fully masked-out query descriptors.

## gpu::BruteForceMatcher_GPU_base::radiusMatchConvert

Converts matrices obtained via `gpu::BruteForceMatcher_GPU_base::radiusMatchSingle()` or `gpu::BruteForceMatcher_GPU_base::radiusMatchCollection()` to vector with `DMatch`.

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchConvert**(const Mat& **trainIdx**, const Mat& **distance**, const Mat& **nMatches**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false )

**C++:** void gpu::BruteForceMatcher_GPU_base::**radiusMatchConvert**(const Mat& **trainIdx**, const Mat& **imgIdx**, const Mat& **distance**, const Mat& **nMatches**, std::vector<std::vector<DMatch>>& **matches**, bool **compactResult**=false )

If `compactResult` is `true` , the `matches` vector does not contain matches for fully masked-out query descriptors.

# 11.10 Image Filtering

Functions and classes described in this section are used to perform various linear or non-linear filtering operations on 2D images.

## gpu::BaseRowFilter_GPU

**class** gpu::**BaseRowFilter_GPU**

Base class for linear or non-linear filters that processes rows of 2D arrays. Such filters are used for the "horizontal" filtering passes in separable filters.

```cpp
class BaseRowFilter_GPU
{
public:
    BaseRowFilter_GPU(int ksize_, int anchor_);
    virtual ~BaseRowFilter_GPU() {}
    virtual void operator()(const GpuMat& src, GpuMat& dst, Stream& stream = Stream::Null()) = 0;
    int ksize, anchor;
};
```

**Note:** This class does not allocate memory for a destination image. Usually this class is used inside `gpu::FilterEngine_GPU`.

## gpu::BaseColumnFilter_GPU

**class** gpu::**BaseColumnFilter_GPU**

Base class for linear or non-linear filters that processes columns of 2D arrays. Such filters are used for the "vertical" filtering passes in separable filters.

```cpp
class BaseColumnFilter_GPU
{
public:
    BaseColumnFilter_GPU(int ksize_, int anchor_);
    virtual ~BaseColumnFilter_GPU() {}
    virtual void operator()(const GpuMat& src, GpuMat& dst, Stream& stream = Stream::Null()) = 0;
    int ksize, anchor;
};
```

**Note:** This class does not allocate memory for a destination image. Usually this class is used inside `gpu::FilterEngine_GPU`.

## gpu::BaseFilter_GPU

**class** gpu::**BaseFilter_GPU**

Base class for non-separable 2D filters.

```
class CV_EXPORTS BaseFilter_GPU
{
public:
    BaseFilter_GPU(const Size& ksize_, const Point& anchor_);
    virtual ~BaseFilter_GPU() {}
    virtual void operator()(const GpuMat& src, GpuMat& dst, Stream& stream = Stream::Null()) = 0;
    Size ksize;
    Point anchor;
};
```

**Note:** This class does not allocate memory for a destination image. Usually this class is used inside gpu::FilterEngine_GPU.

## gpu::FilterEngine_GPU

**class** gpu::**FilterEngine_GPU**

Base class for the Filter Engine.

```
class CV_EXPORTS FilterEngine_GPU
{
public:
    virtual ~FilterEngine_GPU() {}

    virtual void apply(const GpuMat& src, GpuMat& dst,
                       Rect roi = Rect(0,0,-1,-1), Stream& stream = Stream::Null()) = 0;
};
```

The class can be used to apply an arbitrary filtering operation to an image. It contains all the necessary intermediate buffers. Pointers to the initialized FilterEngine_GPU instances are returned by various create*Filter_GPU functions (see below), and they are used inside high-level functions such as gpu::filter2D(), gpu::erode(), gpu::Sobel() , and others.

By using FilterEngine_GPU instead of functions you can avoid unnecessary memory allocation for intermediate buffers and get better performance:

```
while (...)
{
    gpu::GpuMat src = getImg();
    gpu::GpuMat dst;
    // Allocate and release buffers at each iterations
    gpu::GaussianBlur(src, dst, ksize, sigma1);
}

// Allocate buffers only once
cv::Ptr<gpu::FilterEngine_GPU> filter =
    gpu::createGaussianFilter_GPU(CV_8UC4, ksize, sigma1);
while (...)
{
    gpu::GpuMat src = getImg();
    gpu::GpuMat dst;
    filter->apply(src, dst, cv::Rect(0, 0, src.cols, src.rows));
}
// Release buffers only once
filter.release();
```

FilterEngine_GPU can process a rectangular sub-region of an image. By default, if roi == Rect(0,0,-1,-1) , FilterEngine_GPU processes the inner region of an image ( Rect(anchor.x, anchor.y, src_size.width - ksize.width, src_size.height - ksize.height) ) because some filters do not check whether indices are outside the image for better performance. See below to understand which filters support processing the whole image and which do not and identify image type limitations.

**Note:** The GPU filters do not support the in-place mode.

**See Also:**

gpu::BaseRowFilter_GPU, gpu::BaseColumnFilter_GPU, gpu::BaseFilter_GPU, gpu::createFilter2D_GPU(), gpu::createSeparableFilter_GPU(), gpu::createBoxFilter_GPU(), gpu::createMorphologyFilter_GPU(),gpu::createLinearFilter_GPU(),gpu::createSeparableLinearFilter_GPU(), gpu::createDerivFilter_GPU(), gpu::createGaussianFilter_GPU()

## gpu::createFilter2D_GPU

Creates a non-separable filter engine with the specified filter.

**C++:** Ptr<FilterEngine_GPU> gpu::**createFilter2D_GPU**(const Ptr<BaseFilter_GPU>& **filter2D**, int **srcType**, int **dstType**)

> **Parameters**
>
> > **filter2D** – Non-separable 2D filter.
> >
> > **srcType** – Input image type. It must be supported by filter2D .
> >
> > **dstType** – Output image type. It must be supported by filter2D .

Usually this function is used inside such high-level functions as gpu::createLinearFilter_GPU(), gpu::createBoxFilter_GPU().

## gpu::createSeparableFilter_GPU

Creates a separable filter engine with the specified filters.

**C++:** Ptr<FilterEngine_GPU> gpu::**createSeparableFilter_GPU**(const Ptr<BaseRowFilter_GPU>& **rowFilter**, const Ptr<BaseColumnFilter_GPU>& **columnFilter**, int **srcType**, int **bufType**, int **dstType**)

> **Parameters**
>
> > **rowFilter** – "Horizontal" 1D filter.
> >
> > **columnFilter** – "Vertical" 1D filter.
> >
> > **srcType** – Input image type. It must be supported by rowFilter .
> >
> > **bufType** – Buffer image type. It must be supported by rowFilter and columnFilter .
> >
> > **dstType** – Output image type. It must be supported by columnFilter .

Usually this function is used inside such high-level functions as gpu::createSeparableLinearFilter_GPU().

## gpu::getRowSumFilter_GPU

Creates a horizontal 1D box filter.

**C++:** Ptr<BaseRowFilter_GPU> gpu::**getRowSumFilter_GPU**(int **srcType**, int **sumType**, int **ksize**, int **anchor**=-1)

> **Parameters**
>
>> **srcType** – Input image type. Only `CV_8UC1` type is supported for now.
>>
>> **sumType** – Output image type. Only `CV_32FC1` type is supported for now.
>>
>> **ksize** – Kernel size.
>>
>> **anchor** – Anchor point. The default value (-1) means that the anchor is at the kernel center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

## gpu::getColumnSumFilter_GPU

Creates a vertical 1D box filter.

**C++:** Ptr<BaseColumnFilter_GPU> gpu::**getColumnSumFilter_GPU**(int **sumType**, int **dstType**, int **ksize**, int **anchor**=-1)

> **Parameters**
>
>> **sumType** – Input image type. Only `CV_8UC1` type is supported for now.
>>
>> **dstType** – Output image type. Only `CV_32FC1` type is supported for now.
>>
>> **ksize** – Kernel size.
>>
>> **anchor** – Anchor point. The default value (-1) means that the anchor is at the kernel center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

## gpu::createBoxFilter_GPU

Creates a normalized 2D box filter.

**C++:** Ptr<FilterEngine_GPU> gpu::**createBoxFilter_GPU**(int **srcType**, int **dstType**, const Size& **ksize**, const Point& **anchor**=Point(-1,-1))

**C++:** Ptr<BaseFilter_GPU> gpu::**getBoxFilter_GPU**(int **srcType**, int **dstType**, const Size& **ksize**, Point **anchor**=Point(-1, -1))

> **Parameters**
>
>> **srcType** – Input image type supporting `CV_8UC1` and `CV_8UC4` .
>>
>> **dstType** – Output image type. It supports only the same values as the source type.
>>
>> **ksize** – Kernel size.
>>
>> **anchor** – Anchor point. The default value `Point(-1, -1)` means that the anchor is at the kernel center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`boxFilter()`

## gpu::boxFilter

Smooths the image using the normalized box filter.

**C++:** void gpu::**boxFilter**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, Size **ksize**, Point **anchor**=Point(-1,-1), Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Input image. `CV_8UC1` and `CV_8UC4` source types are supported.
> >
> > **dst** – Output image type. The size and type is the same as `src` .
> >
> > **ddepth** – Output image depth. If -1, the output image has the same depth as the input one. The only values allowed here are `CV_8U` and -1.
> >
> > **ksize** – Kernel size.
> >
> > **anchor** – Anchor point. The default value `Point(-1, -1)` means that the anchor is at the kernel center.
> >
> > **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`boxFilter()`

## gpu::blur

Acts as a synonym for the normalized box filter.

**C++:** void gpu::**blur**(const GpuMat& **src**, GpuMat& **dst**, Size **ksize**, Point **anchor**=Point(-1,-1), Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **src** – Input image. `CV_8UC1` and `CV_8UC4` source types are supported.
> >
> > **dst** – Output image type with the same size and type as `src` .
> >
> > **ksize** – Kernel size.
> >
> > **anchor** – Anchor point. The default value Point(-1, -1) means that the anchor is at the kernel center.
> >
> > **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`blur()`, `gpu::boxFilter()`

## gpu::createMorphologyFilter_GPU

Creates a 2D morphological filter.

**C++:** Ptr<FilterEngine_GPU> gpu::**createMorphologyFilter_GPU**(int **op**, int **type**, const Mat& **kernel**, const Point& **anchor**=Point(-1,-1), int **iterations**=1 )

**C++:** Ptr<BaseFilter_GPU> gpu::**getMorphologyFilter_GPU**(int **op**, int **type**, const Mat& **kernel**, const Size& **ksize**, Point **anchor**=Point(-1,-1))

> **Parameters**
>
> > **op** – Morphology operation id. Only `MORPH_ERODE` and `MORPH_DILATE` are supported.
> >
> > **type** – Input/output image type. Only `CV_8UC1` and `CV_8UC4` are supported.
> >
> > **kernel** – 2D 8-bit structuring element for the morphological operation.
> >
> > **ksize** – Size of a horizontal or vertical structuring element used for separable morphological operations.
> >
> > **anchor** – Anchor position within the structuring element. Negative values mean that the anchor is at the center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`createMorphologyFilter()`

## gpu::erode

Erodes an image by using a specific structuring element.

**C++:** void gpu::**erode**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **kernel**, Point **anchor**=Point(-1, -1), int **iterations**=1 )

**C++:** void gpu::**erode**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **kernel**, GpuMat& **buf**, Point **anchor**=Point(-1, -1), int **iterations**=1, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. Only `CV_8UC1` and `CV_8UC4` types are supported.
> >
> > **dst** – Destination image with the same size and type as `src` .
> >
> > **kernel** – Structuring element used for erosion. If `kernel=Mat()`, a 3x3 rectangular structuring element is used.

> **anchor** – Position of an anchor within the element. The default value (`-1, -1`) means that the anchor is at the element center.
>
> **iterations** – Number of times erosion to be applied.
>
> **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`erode()`

## gpu::dilate

Dilates an image by using a specific structuring element.

**C++:** void gpu::**dilate**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **kernel**, Point **anchor**=Point(-1, -1), int **iterations**=1 )

**C++:** void gpu::**dilate**(const GpuMat& **src**, GpuMat& **dst**, const Mat& **kernel**, GpuMat& **buf**, Point **anchor**=Point(-1, -1), int **iterations**=1, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **src** – Source image. `CV_8UC1` and `CV_8UC4` source types are supported.
> >
> > **dst** – Destination image with the same size and type as `src`.
> >
> > **kernel** – Structuring element used for dilation. If `kernel=Mat()`, a 3x3 rectangular structuring element is used.
> >
> > **anchor** – Position of an anchor within the element. The default value (`-1, -1`) means that the anchor is at the element center.
> >
> > **iterations** – Number of times dilation to be applied.
> >
> > **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`dilate()`

## gpu::morphologyEx

Applies an advanced morphological operation to an image.

**C++:** void gpu::**morphologyEx**(const GpuMat& **src**, GpuMat& **dst**, int **op**, const Mat& **kernel**, Point **anchor**=Point(-1, -1), int **iterations**=1 )

**C++:** void gpu::**morphologyEx**(const GpuMat& **src**, GpuMat& **dst**, int **op**, const Mat& **kernel**, GpuMat& **buf1**, GpuMat& **buf2**, Point **anchor**=Point(-1, -1), int **iterations**=1, Stream& **stream**=Stream::Null() )

> **Parameters**

> > **src** – Source image. `CV_8UC1` and `CV_8UC4` source types are supported.
>
> > **dst** – Destination image with the same size and type as `src` .
>
> > **op** – Type of morphological operation. The following types are possible:
>
> > > – **MORPH_OPEN** opening
> > >
> > > – **MORPH_CLOSE** closing
> > >
> > > – **MORPH_GRADIENT** morphological gradient
> > >
> > > – **MORPH_TOPHAT** "top hat"
> > >
> > > – **MORPH_BLACKHAT** "black hat"
>
> > **kernel** – Structuring element.
>
> > **anchor** – Position of an anchor within the element. The default value `Point(-1, -1)` means that the anchor is at the element center.
>
> > **iterations** – Number of times erosion and dilation to be applied.
>
> > **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`morphologyEx()`

## gpu::createLinearFilter_GPU

Creates a non-separable linear filter.

**C++:** Ptr<FilterEngine_GPU> gpu::**createLinearFilter_GPU**(int **srcType**, int **dstType**, const Mat& **kernel**, Point **anchor**=Point(-1,-1), int **borderType**=BORDER_DEFAULT )

> > **Parameters**
>
> > > **srcType** – Input image type. Supports `CV_8U` , `CV_16U` and `CV_32F` one and four channel image.
> > >
> > > **dstType** – Output image type. The same type as `src` is supported.
> > >
> > > **kernel** – 2D array of filter coefficients. Floating-point coefficients will be converted to fixed-point representation before the actual processing. Supports size up to 16. For larger kernels use `gpu::convolve()`.
> > >
> > > **anchor** – Anchor point. The default value Point(-1, -1) means that the anchor is at the kernel center.
> > >
> > > **borderType** – Pixel extrapolation method. For details, see `borderInterpolate()` .

**See Also:**

`createLinearFilter()`

## gpu::filter2D

Applies the non-separable 2D linear filter to an image.

**C++:** void gpu::**filter2D**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, const Mat& **kernel**, Point **anchor**=Point(-1,-1), int **borderType**=BORDER_DEFAULT, Stream& **stream**=Stream::Null())

>     Parameters

>> **src** – Source image. Supports `CV_8U` , `CV_16U` and `CV_32F` one and four channel image.

>> **dst** – Destination image. The size and the number of channels is the same as `src` .

>> **ddepth** – Desired depth of the destination image. If it is negative, it is the same as `src.depth()` . It supports only the same depth as the source image depth.

>> **kernel** – 2D array of filter coefficients.

>> **anchor** – Anchor of the kernel that indicates the relative position of a filtered point within the kernel. The anchor resides within the kernel. The special default value (-1,-1) means that the anchor is at the kernel center.

>> **borderType** – Pixel extrapolation method. For details, see `borderInterpolate()` .

>> **stream** – Stream for the asynchronous version.

**See Also:**

`filter2D()`, `gpu::convolve()`

## gpu::Laplacian

Applies the Laplacian operator to an image.

**C++:** void gpu::**Laplacian**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, int **ksize**=1, double **scale**=1, int **borderType**=BORDER_DEFAULT, Stream& **stream**=Stream::Null())

>     Parameters

>> **src** – Source image. `CV_8UC1` and `CV_8UC4` source types are supported.

>> **dst** – Destination image. The size and number of channels is the same as `src` .

>> **ddepth** – Desired depth of the destination image. It supports only the same depth as the source image depth.

>> **ksize** – Aperture size used to compute the second-derivative filters (see `getDerivKernels()`). It must be positive and odd. Only `ksize = 1` and `ksize = 3` are supported.

>> **scale** – Optional scale factor for the computed Laplacian values. By default, no scaling is applied (see `getDerivKernels()` ).

>> **borderType** – Pixel extrapolation method. For details, see `borderInterpolate()` .

>> **stream** – Stream for the asynchronous version.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

**See Also:**

`Laplacian()`, `gpu::filter2D()`

## gpu::getLinearRowFilter_GPU

Creates a primitive row filter with the specified kernel.

**C++:** Ptr<BaseRowFilter_GPU> gpu::**getLinearRowFilter_GPU**(int **srcType**, int **bufType**, const Mat& **rowKernel**, int **anchor**=-1, int **border-Type**=BORDER_DEFAULT )

>  Parameters

>>  **srcType** – Source array type. Only `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` source types are supported.

>>  **bufType** – Intermediate buffer type with as many channels as `srcType` .

>>  **rowKernel** – Filter coefficients. Support kernels with `size <= 16` .

>>  **anchor** – Anchor position within the kernel. Negative values mean that the anchor is positioned at the aperture center.

>>  **borderType** – Pixel extrapolation method. For details, see `borderInterpolate()`. For details on limitations, see below.

There are two versions of the algorithm: NPP and OpenCV.

- NPP version is called when `srcType == CV_8UC1` or `srcType == CV_8UC4` and `bufType == srcType` . Otherwise, the OpenCV version is called. NPP supports only `BORDER_CONSTANT` border type and does not check indices outside the image.

- OpenCV version supports only `CV_32F` buffer depth and `BORDER_REFLECT101` , `BORDER_REPLICATE` , and `BORDER_CONSTANT` border types. It checks indices outside the image.

**See Also:**

`createSeparableLinearFilter()` .

## gpu::getLinearColumnFilter_GPU

Creates a primitive column filter with the specified kernel.

**C++:** Ptr<BaseColumnFilter_GPU> gpu::**getLinearColumnFilter_GPU**(int **bufType**, int **dstType**, const Mat& **columnKernel**, int **anchor**=-1, int **border-Type**=BORDER_DEFAULT )

>  Parameters

>>  **bufType** – Intermediate buffer type with as many channels as `dstType` .

>>  **dstType** – Destination array type. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` destination types are supported.

>>  **columnKernel** – Filter coefficients. Support kernels with `size <= 16` .

>>  **anchor** – Anchor position within the kernel. Negative values mean that the anchor is positioned at the aperture center.

>>  **borderType** – Pixel extrapolation method. For details, see `borderInterpolate()` . For details on limitations, see below.

There are two versions of the algorithm: NPP and OpenCV.

- NPP version is called when `dstType == CV_8UC1` or `dstType == CV_8UC4` and `bufType == dstType`. Otherwise, the OpenCV version is called. NPP supports only `BORDER_CONSTANT` border type and does not check indices outside the image.

- OpenCV version supports only `CV_32F` buffer depth and `BORDER_REFLECT101`, `BORDER_REPLICATE`, and `BORDER_CONSTANT` border types. It checks indices outside image.

**See Also:**

`gpu::getLinearRowFilter_GPU()`, `createSeparableLinearFilter()`

## gpu::createSeparableLinearFilter_GPU

Creates a separable linear filter engine.

**C++:** Ptr<FilterEngine_GPU> gpu::**createSeparableLinearFilter_GPU**(int **srcType**, int **dstType**, const Mat& **rowKernel**, const Mat& **columnKernel**, const Point& **anchor**=Point(-1,-1), int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1)

> **Parameters**
>
> > **srcType** – Source array type. `CV_8UC1`, `CV_8UC4`, `CV_16SC1`, `CV_16SC2`, `CV_16SC3`, `CV_32SC1`, `CV_32FC1` source types are supported.
> >
> > **dstType** – Destination array type. `CV_8UC1`, `CV_8UC4`, `CV_16SC1`, `CV_16SC2`, `CV_16SC3`, `CV_32SC1`, `CV_32FC1` destination types are supported.
> >
> > **rowKernel** – Horizontal filter coefficients. Support kernels with `size <= 16`.
> >
> > **columnKernel** – Vertical filter coefficients. Support kernels with `size <= 16`.
> >
> > **anchor** – Anchor position within the kernel. Negative values mean that anchor is positioned at the aperture center.
> >
> > **rowBorderType** – Pixel extrapolation method in the vertical direction For details, see `borderInterpolate()`. For details on limitations, see `gpu::getLinearRowFilter_GPU()`, cpp:ocv:func:*gpu::getLinearColumnFilter_GPU*.
> >
> > **columnBorderType** – Pixel extrapolation method in the horizontal direction.

**See Also:**

`gpu::getLinearRowFilter_GPU()`, `gpu::getLinearColumnFilter_GPU()`, `createSeparableLinearFilter()`

## gpu::sepFilter2D

Applies a separable 2D linear filter to an image.

**C++:** void gpu::**sepFilter2D**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, const Mat& **kernelX**, const Mat& **kernelY**, Point **anchor**=Point(-1,-1), int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1 )

**C++:** void gpu::**sepFilter2D**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, const Mat& **kernelX**, const Mat& **kernelY**, GpuMat& **buf**, Point **anchor**=Point(-1,-1), int **row-BorderType**=BORDER_DEFAULT, int **columnBorderType**=-1, Stream& **stream**=Stream::Null() )

> Parameters
>> **src** – Source image. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_32SC1` , `CV_32FC1` source types are supported.
>>
>> **dst** – Destination image with the same size and number of channels as `src` .
>>
>> **ddepth** – Destination image depth. `CV_8U` , `CV_16S` , `CV_32S` , and `CV_32F` are supported.
>>
>> **kernelX** – Horizontal filter coefficients.
>>
>> **kernelY** – Vertical filter coefficients.
>>
>> **anchor** – Anchor position within the kernel. The default value (`-1, 1`) means that the anchor is at the kernel center.
>>
>> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.
>>
>> **columnBorderType** – Pixel extrapolation method in the horizontal direction.
>>
>> **stream** – Stream for the asynchronous version.

See Also:

`gpu::createSeparableLinearFilter_GPU()`, `sepFilter2D()`

## gpu::createDerivFilter_GPU

Creates a filter engine for the generalized Sobel operator.

**C++:** Ptr<FilterEngine_GPU> gpu::**createDerivFilter_GPU**(int **srcType**, int **dstType**, int **dx**, int **dy**, int **ksize**, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1)

> Parameters
>> **srcType** – Source image type. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` source types are supported.
>>
>> **dstType** – Destination image type with as many channels as `srcType` , `CV_8U` , `CV_16S` , `CV_32S` , and `CV_32F` depths are supported.
>>
>> **dx** – Derivative order in respect of x.
>>
>> **dy** – Derivative order in respect of y.
>>
>> **ksize** – Aperture size. See `getDerivKernels()` for details.
>>
>> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.
>>
>> **columnBorderType** – Pixel extrapolation method in the horizontal direction.

See Also:

`gpu::createSeparableLinearFilter_GPU()`, `createDerivFilter()`

## gpu::Sobel

Applies the generalized Sobel operator to an image.

**C++:** void gpu::**Sobel**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, int **dx**, int **dy**, int **ksize**=3, double **scale**=1, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1 )

**C++:** void gpu::**Sobel**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, int **dx**, int **dy**, GpuMat& **buf**, int **ksize**=3, double **scale**=1, int **rowBorderType**=BORDER_DEFAULT, int **column-BorderType**=-1, Stream& **stream**=Stream::Null() )

> **Parameters**

>> **src** – Source image. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` source types are supported.

>> **dst** – Destination image with the same size and number of channels as source image.

>> **ddepth** – Destination image depth. `CV_8U` , `CV_16S` , `CV_32S` , and `CV_32F` are supported.

>> **dx** – Derivative order in respect of x.

>> **dy** – Derivative order in respect of y.

>> **ksize** – Size of the extended Sobel kernel. Possible values are 1, 3, 5 or 7.

>> **scale** – Optional scale factor for the computed derivative values. By default, no scaling is applied. For details, see `getDerivKernels()` .

>> **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.

>> **columnBorderType** – Pixel extrapolation method in the horizontal direction.

>> **stream** – Stream for the asynchronous version.

**See Also:**

`gpu::createSeparableLinearFilter_GPU()`, `Sobel()`

## gpu::Scharr

Calculates the first x- or y- image derivative using the Scharr operator.

**C++:** void gpu::**Scharr**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, int **dx**, int **dy**, double **scale**=1, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1 )

**C++:** void gpu::**Scharr**(const GpuMat& **src**, GpuMat& **dst**, int **ddepth**, int **dx**, int **dy**, GpuMat& **buf**, double **scale**=1, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1, Stream& **stream**=Stream::Null() )

> **Parameters**

>> **src** – Source image. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` source types are supported.

>> **dst** – Destination image with the same size and number of channels as `src` has.

>> **ddepth** – Destination image depth. `CV_8U` , `CV_16S` , `CV_32S` , and `CV_32F` are supported.

>> **dx** – Order of the derivative x.

>> **dy** – Order of the derivative y.

>> **scale** – Optional scale factor for the computed derivative values. By default, no scaling is applied. See `getDerivKernels()` for details.

    **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.

    **columnBorderType** – Pixel extrapolation method in the horizontal direction.

    **stream** – Stream for the asynchronous version.

**See Also:**

`gpu::createSeparableLinearFilter_GPU()`, `Scharr()`

## gpu::createGaussianFilter_GPU

Creates a Gaussian filter engine.

**C++:** Ptr<FilterEngine_GPU> gpu::**createGaussianFilter_GPU**(int **type**, Size **ksize**, double **sigma1**, double **sigma2**=0, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1 )

    **Parameters**

        **type** – Source and destination image type. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` are supported.

        **ksize** – Aperture size. See `getGaussianKernel()` for details.

        **sigma1** – Gaussian sigma in the horizontal direction. See `getGaussianKernel()` for details.

        **sigma2** – Gaussian sigma in the vertical direction. If 0, then $sigma2 \leftarrow sigma1$ .

        **rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.

        **columnBorderType** – Pixel extrapolation method in the horizontal direction.

**See Also:**

`gpu::createSeparableLinearFilter_GPU()`, `createGaussianFilter()`

## gpu::GaussianBlur

Smooths an image using the Gaussian filter.

**C++:** void gpu::**GaussianBlur**(const GpuMat& **src**, GpuMat& **dst**, Size **ksize**, double **sigma1**, double **sigma2**=0, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1 )

**C++:** void gpu::**GaussianBlur**(const GpuMat& **src**, GpuMat& **dst**, Size **ksize**, GpuMat& **buf**, double **sigma1**, double **sigma2**=0, int **rowBorderType**=BORDER_DEFAULT, int **columnBorderType**=-1, Stream& **stream**=Stream::Null() )

    **Parameters**

        **src** – Source image. `CV_8UC1` , `CV_8UC4` , `CV_16SC1` , `CV_16SC2` , `CV_16SC3` , `CV_32SC1` , `CV_32FC1` source types are supported.

        **dst** – Destination image with the same size and type as `src` .

        **ksize** – Gaussian kernel size. `ksize.width` and `ksize.height` can differ but they both must be positive and odd. If they are zeros, they are computed from `sigma1` and `sigma2` .

>**sigma1** – Gaussian kernel standard deviation in X direction.

>**sigma2** – Gaussian kernel standard deviation in Y direction. If `sigma2` is zero, it is set to be equal to `sigma1`. If they are both zeros, they are computed from `ksize.width` and `ksize.height`, respectively. See `getGaussianKernel()` for details. To fully control the result regardless of possible future modification of all this semantics, you are recommended to specify all of `ksize`, `sigma1`, and `sigma2`.

>**rowBorderType** – Pixel extrapolation method in the vertical direction. For details, see `borderInterpolate()`.

>**columnBorderType** – Pixel extrapolation method in the horizontal direction.

>**stream** – Stream for the asynchronous version.

**See Also:**

`gpu::createGaussianFilter_GPU()`, `GaussianBlur()`

## gpu::getMaxFilter_GPU

Creates the maximum filter.

**C++:** Ptr<BaseFilter_GPU> gpu::**getMaxFilter_GPU**(int **srcType**, int **dstType**, const Size& **ksize**, Point **anchor**=Point(-1,-1))

>    **Parameters**

>>    **srcType** – Input image type. Only `CV_8UC1` and `CV_8UC4` are supported.

>>    **dstType** – Output image type. It supports only the same type as the source type.

>>    **ksize** – Kernel size.

>>    **anchor** – Anchor point. The default value (-1) means that the anchor is at the kernel center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

## gpu::getMinFilter_GPU

Creates the minimum filter.

**C++:** Ptr<BaseFilter_GPU> gpu::**getMinFilter_GPU**(int **srcType**, int **dstType**, const Size& **ksize**, Point **anchor**=Point(-1,-1))

>    **Parameters**

>>    **srcType** – Input image type. Only `CV_8UC1` and `CV_8UC4` are supported.

>>    **dstType** – Output image type. It supports only the same type as the source type.

>>    **ksize** – Kernel size.

>>    **anchor** – Anchor point. The default value (-1) means that the anchor is at the kernel center.

**Note:** This filter does not check out-of-border accesses, so only a proper sub-matrix of a bigger matrix has to be passed to it.

# 11.11 Camera Calibration and 3D Reconstruction

## gpu::StereoBM_GPU

**class** gpu::**StereoBM_GPU**

Class computing stereo correspondence (disparity map) using the block matching algorithm.

```cpp
class StereoBM_GPU
{
public:
    enum { BASIC_PRESET = 0, PREFILTER_XSOBEL = 1 };

    enum { DEFAULT_NDISP = 64, DEFAULT_WINSZ = 19 };

    StereoBM_GPU();
    StereoBM_GPU(int preset, int ndisparities = DEFAULT_NDISP,
                int winSize = DEFAULT_WINSZ);

    void operator() (const GpuMat& left, const GpuMat& right,
                     GpuMat& disparity, Stream& stream = Stream::Null());

    static bool checkIfGpuCallReasonable();

    int preset;
    int ndisp;
    int winSize;

    float avergeTexThreshold;

    ...
};
```

The class also performs pre- and post-filtering steps: Sobel pre-filtering (if PREFILTER_XSOBEL flag is set) and low textureness filtering (if averageTexThreshols > 0 ). If avergeTexThreshold = 0 , low textureness filtering is disabled. Otherwise, the disparity is set to 0 in each point (x, y) , where for the left image

$$\sum HorizontalGradiensInWindow(x, y, winSize) < (winSize \cdot winSize) \cdot avergeTexThreshold$$

This means that the input left image is low textured.

## gpu::StereoBM_GPU::StereoBM_GPU

Enables gpu::StereoBM_GPU constructors.

**C++:** gpu::StereoBM_GPU::**StereoBM_GPU**()

**C++:** gpu::StereoBM_GPU::**StereoBM_GPU**(int **preset**, int **ndisparities**=DEFAULT_NDISP, int **winSize**=DEFAULT_WINSZ)

> **Parameters**
>
> > **preset** – Parameter presetting:
> >
> > – **BASIC_PRESET** Basic mode without pre-processing.
> >
> > – **PREFILTER_XSOBEL** Sobel pre-filtering mode.

**ndisparities** – Number of disparities. It must be a multiple of 8 and less or equal to 256.

**winSize** – Block size.

## gpu::StereoBM_GPU::operator ()

Enables the stereo correspondence operator that finds the disparity for the specified rectified stereo pair.

**C++:** void gpu::StereoBM_GPU::**operator()**(const GpuMat& **left**, const GpuMat& **right**, GpuMat& **disparity**, Stream& **stream**=Stream::Null())

> Parameters

> > **left** – Left image. Only CV_8UC1 type is supported.

> > **right** – Right image with the same size and the same type as the left one.

> > **disparity** – Output disparity map. It is a CV_8UC1 image with the same size as the input images.

> > **stream** – Stream for the asynchronous version.

## gpu::StereoBM_GPU::checkIfGpuCallReasonable

Uses a heuristic method to estimate whether the current GPU is faster than the CPU in this algorithm. It queries the currently active device.

**C++:** bool gpu::StereoBM_GPU::**checkIfGpuCallReasonable**()

## gpu::StereoBeliefPropagation

**class** gpu::**StereoBeliefPropagation**

Class computing stereo correspondence using the belief propagation algorithm.

```cpp
class StereoBeliefPropagation
{
public:
    enum { DEFAULT_NDISP  = 64 };
    enum { DEFAULT_ITERS  = 5  };
    enum { DEFAULT_LEVELS = 5  };

    static void estimateRecommendedParams(int width, int height,
        int& ndisp, int& iters, int& levels);

    explicit StereoBeliefPropagation(int ndisp = DEFAULT_NDISP,
        int iters  = DEFAULT_ITERS,
        int levels = DEFAULT_LEVELS,
        int msg_type = CV_32F);
    StereoBeliefPropagation(int ndisp, int iters, int levels,
        float max_data_term, float data_weight,
        float max_disc_term, float disc_single_jump,
        int msg_type = CV_32F);

    void operator()(const GpuMat& left, const GpuMat& right,
                    GpuMat& disparity, Stream& stream = Stream::Null());
    void operator()(const GpuMat& data, GpuMat& disparity, Stream& stream = Stream::Null());
```

```
    int ndisp;

    int iters;
    int levels;

    float max_data_term;
    float data_weight;
    float max_disc_term;
    float disc_single_jump;

    int msg_type;


    ...
};
```

The class implements algorithm described in [Felzenszwalb2006] . It can compute own data cost (using a truncated linear model) or use a user-provided data cost.

---

**Note:** StereoBeliefPropagation requires a lot of memory for message storage:

$$width\_step \cdot height \cdot ndisp \cdot 4 \cdot (1 + 0.25)$$

and for data cost storage:

$$width\_step \cdot height \cdot ndisp \cdot (1 + 0.25 + 0.0625 + \cdots + \frac{1}{4^{levels}})$$

width_step is the number of bytes in a line including padding.

---

## gpu::StereoBeliefPropagation::StereoBeliefPropagation

Enables the gpu::StereoBeliefPropagation constructors.

**C++:** gpu::StereoBeliefPropagation::**StereoBeliefPropagation**(int     **ndisp**=DEFAULT_NDISP, int **iters**=DEFAULT_ITERS, int **levels**=DEFAULT_LEVELS, int **msg_type**=CV_32F)

**C++:** gpu::StereoBeliefPropagation::**StereoBeliefPropagation**(int **ndisp**, int **iters**, int **levels**, float     **max_data_term**, float     **data_weight**,     float **max_disc_term**,     float **disc_single_jump**,     int **msg_type**=CV_32F)

        **Parameters**

                **ndisp** – Number of disparities.

                **iters** – Number of BP iterations on each level.

                **levels** – Number of levels.

                **max_data_term** – Threshold for data cost truncation.

                **data_weight** – Data weight.

                **max_disc_term** – Threshold for discontinuity truncation.

                **disc_single_jump** – Discontinuity single jump.

---

**msg_type** – Type for messages. `CV_16SC1` and `CV_32FC1` types are supported.

`StereoBeliefPropagation` uses a truncated linear model for the data cost and discontinuity terms:

$$DataCost = data\_weight \cdot \min(|I_2 - I_1|, max\_data\_term)$$

$$DiscTerm = \min(disc\_single\_jump \cdot |f_1 - f_2|, max\_disc\_term)$$

For more details, see [Felzenszwalb2006].

By default, `gpu::StereoBeliefPropagation` uses floating-point arithmetics and the `CV_32FC1` type for messages. But it can also use fixed-point arithmetics and the `CV_16SC1` message type for better performance. To avoid an overflow in this case, the parameters must satisfy the following requirement:

$$10 \cdot 2^{levels-1} \cdot max\_data\_term < SHRT\_MAX$$

## gpu::StereoBeliefPropagation::estimateRecommendedParams

Uses a heuristic method to compute the recommended parameters ( `ndisp`, `iters` and `levels` ) for the specified image size ( `width` and `height` ).

**C++:** void gpu::StereoBeliefPropagation::**estimateRecommendedParams**(int **width**, int **height**, int& **ndisp**, int& **iters**, int& **levels**)

## gpu::StereoBeliefPropagation::operator ()

Enables the stereo correspondence operator that finds the disparity for the specified rectified stereo pair or data cost.

**C++:** void gpu::StereoBeliefPropagation::**operator()**(const GpuMat& **left**, const GpuMat& **right**, GpuMat& **disparity**, Stream& **stream**=Stream::Null())

**C++:** void gpu::StereoBeliefPropagation::**operator()**(const GpuMat& **data**, GpuMat& **disparity**, Stream& **stream**=Stream::Null())

>   **Parameters**

>>   **left** – Left image. `CV_8UC1` , `CV_8UC3` and `CV_8UC4` types are supported.

>>   **right** – Right image with the same size and the same type as the left one.

>>   **data** – User-specified data cost, a matrix of `msg_type` type and `Size(<image columns>*ndisp, <image rows>)` size.

>>   **disparity** – Output disparity map. If `disparity` is empty, the output type is `CV_16SC1` . Otherwise, the type is retained.

>>   **stream** – Stream for the asynchronous version.

## gpu::StereoConstantSpaceBP

**class** gpu::**StereoConstantSpaceBP**

Class computing stereo correspondence using the constant space belief propagation algorithm.

```
class StereoConstantSpaceBP
{
public:
    enum { DEFAULT_NDISP    = 128 };
    enum { DEFAULT_ITERS    = 8   };
    enum { DEFAULT_LEVELS   = 4   };
    enum { DEFAULT_NR_PLANE = 4   };

    static void estimateRecommendedParams(int width, int height,
        int& ndisp, int& iters, int& levels, int& nr_plane);

    explicit StereoConstantSpaceBP(int ndisp = DEFAULT_NDISP,
        int iters   = DEFAULT_ITERS,
        int levels  = DEFAULT_LEVELS,
        int nr_plane = DEFAULT_NR_PLANE,
        int msg_type = CV_32F);
    StereoConstantSpaceBP(int ndisp, int iters, int levels, int nr_plane,
        float max_data_term, float data_weight,
        float max_disc_term, float disc_single_jump,
        int min_disp_th = 0,
        int msg_type = CV_32F);

    void operator()(const GpuMat& left, const GpuMat& right,
                    GpuMat& disparity, Stream& stream = Stream::Null());

    int ndisp;

    int iters;
    int levels;

    int nr_plane;

    float max_data_term;
    float data_weight;
    float max_disc_term;
    float disc_single_jump;

    int min_disp_th;

    int msg_type;

    bool use_local_init_data_cost;

    ...
};
```

The class implements algorithm described in [Yang2010]. `StereoConstantSpaceBP` supports both local minimum and global minimum data cost initialization algorithms. For more details, see the paper mentioned above. By default, a local algorithm is used. To enable a global algorithm, set `use_local_init_data_cost` to `false` .

## gpu::StereoConstantSpaceBP::StereoConstantSpaceBP

Enables the `gpu::StereoConstantSpaceBP` constructors.

**C++:** `gpu::StereoConstantSpaceBP::`**`StereoConstantSpaceBP`**(int **ndisp**=DEFAULT_NDISP, int **iters**=DEFAULT_ITERS, int **levels**=DEFAULT_LEVELS, int **nr_plane**=DEFAULT_NR_PLANE, int **msg_type**=CV_32F)

**C++:** `gpu::StereoConstantSpaceBP::`**`StereoConstantSpaceBP`**(int **ndisp**, int **iters**, int **levels**, int **nr_plane**, float **max_data_term**, float **data_weight**, float **max_disc_term**, float **disc_single_jump**, int **min_disp_th**=0, int **msg_type**=CV_32F)

> **Parameters**

>> **ndisp** – Number of disparities.

>> **iters** – Number of BP iterations on each level.

>> **levels** – Number of levels.

>> **nr_plane** – Number of disparity levels on the first level.

>> **max_data_term** – Truncation of data cost.

>> **data_weight** – Data weight.

>> **max_disc_term** – Truncation of discontinuity.

>> **disc_single_jump** – Discontinuity single jump.

>> **min_disp_th** – Minimal disparity threshold.

>> **msg_type** – Type for messages. `CV_16SC1` and `CV_32FC1` types are supported.

`StereoConstantSpaceBP` uses a truncated linear model for the data cost and discontinuity terms:

$$\mathrm{DataCost} = data\_weight \cdot \min(|I_2 - I_1|, max\_data\_term)$$

$$\mathrm{DiscTerm} = \min(disc\_single\_jump \cdot |f_1 - f_2|, max\_disc\_term)$$

For more details, see [Yang2010].

By default, `StereoConstantSpaceBP` uses floating-point arithmetics and the `CV_32FC1` type for messages. But it can also use fixed-point arithmetics and the `CV_16SC1` message type for better performance. To avoid an overflow in this case, the parameters must satisfy the following requirement:

$$10 \cdot 2^{levels-1} \cdot max\_data\_term < \mathrm{SHRT\_MAX}$$

## gpu::StereoConstantSpaceBP::estimateRecommendedParams

Uses a heuristic method to compute parameters (ndisp, iters, levelsand nrplane) for the specified image size (widthand height).

**C++:** void `gpu::StereoConstantSpaceBP::`**`estimateRecommendedParams`**(int **width**, int **height**, int& **ndisp**, int& **iters**, int& **levels**, int& **nr_plane**)

## gpu::StereoConstantSpaceBP::operator ()

Enables the stereo correspondence operator that finds the disparity for the specified rectified stereo pair.

**C++:** void gpu::StereoConstantSpaceBP::**operator()**(const GpuMat& **left**, const GpuMat& **right**, GpuMat& **disparity**, Stream& **stream**=Stream::Null())

>        **Parameters**
>
>> **left** – Left image. `CV_8UC1` , `CV_8UC3` and `CV_8UC4` types are supported.
>>
>> **right** – Right image with the same size and the same type as the left one.
>>
>> **disparity** – Output disparity map. If `disparity` is empty, the output type is `CV_16SC1` . Otherwise, the output type is `disparity.type()` .
>>
>> **stream** – Stream for the asynchronous version.

## gpu::DisparityBilateralFilter

**class** gpu::**DisparityBilateralFilter**

Class refining a disparity map using joint bilateral filtering.

```
class CV_EXPORTS DisparityBilateralFilter
{
public:
    enum { DEFAULT_NDISP  = 64 };
    enum { DEFAULT_RADIUS = 3 };
    enum { DEFAULT_ITERS  = 1 };

    explicit DisparityBilateralFilter(int ndisp = DEFAULT_NDISP,
        int radius = DEFAULT_RADIUS, int iters = DEFAULT_ITERS);

    DisparityBilateralFilter(int ndisp, int radius, int iters,
        float edge_threshold, float max_disc_threshold,
        float sigma_range);

    void operator()(const GpuMat& disparity, const GpuMat& image,
                    GpuMat& dst, Stream& stream = Stream::Null());

    ...
};
```

The class implements [Yang2010] algorithm.

## gpu::DisparityBilateralFilter::DisparityBilateralFilter

Enables the gpu::DisparityBilateralFilter constructors.

**C++:** gpu::DisparityBilateralFilter::**DisparityBilateralFilter**(int **ndisp**=DEFAULT_NDISP, int **radius**=DEFAULT_RADIUS, int **iters**=DEFAULT_ITERS)

**C++:** gpu::DisparityBilateralFilter::**DisparityBilateralFilter**(int **ndisp**, int **radius**, int **iters**, float **edge_threshold**, float **max_disc_threshold**, float **sigma_range**)

**Parameters**

**ndisp** – Number of disparities.

**radius** – Filter radius.

**iters** – Number of iterations.

**edge_threshold** – Threshold for edges.

**max_disc_threshold** – Constant to reject outliers.

**sigma_range** – Filter range.

## gpu::DisparityBilateralFilter::operator ()

Refines a disparity map using joint bilateral filtering.

**C++:** void gpu::DisparityBilateralFilter::**operator()**(const GpuMat& **disparity**, const GpuMat& **image**, GpuMat& **dst**, Stream& **stream**=Stream::Null())

**Parameters**

**disparity** – Input disparity map. `CV_8UC1` and `CV_16SC1` types are supported.

**image** – Input image. `CV_8UC1` and `CV_8UC3` types are supported.

**dst** – Destination disparity map. It has the same size and type as `disparity` .

**stream** – Stream for the asynchronous version.

## gpu::drawColorDisp

Colors a disparity image.

**C++:** void gpu::**drawColorDisp**(const GpuMat& **src_disp**, GpuMat& **dst_disp**, int **ndisp**, Stream& **stream**=Stream::Null())

**Parameters**

**src_disp** – Source disparity image. `CV_8UC1` and `CV_16SC1` types are supported.

**dst_disp** – Output disparity image. It has the same size as `src_disp` . The type is `CV_8UC4` in BGRA format (alpha = 255).

**ndisp** – Number of disparities.

**stream** – Stream for the asynchronous version.

This function draws a colored disparity map by converting disparity values from `[0..ndisp)` interval first to `HSV` color space (where different disparity values correspond to different hues) and then converting the pixels to `RGB` for visualization.

## gpu::reprojectImageTo3D

Reprojects a disparity image to 3D space.

**C++:** void gpu::**reprojectImageTo3D**(const GpuMat& **disp**, GpuMat& **xyzw**, const Mat& **Q**, int **dst_cn**=4, Stream& **stream**=Stream::Null())

**Parameters**

**disp** – Input disparity image. `CV_8U` and `CV_16S` types are supported.

**xyzw** – Output 3- or 4-channel floating-point image of the same size as `disp` . Each element of `xyzw(x,y)` contains 3D coordinates `(x,y,z)` or `(x,y,z,1)` of the point `(x,y)` , computed from the disparity map.

**Q** – $4 \times 4$ perspective transformation matrix that can be obtained via `stereoRectify()` .

**dst_cn** – The number of channels for output image. Can be 3 or 4.

**stream** – Stream for the asynchronous version.

**See Also:**

`reprojectImageTo3D()`

## gpu::solvePnPRansac

Finds the object pose from 3D-2D point correspondences.

**C++:** void gpu::**solvePnPRansac**(const Mat& **object**, const Mat& **image**, const Mat& **camera_mat**, const Mat& **dist_coef**, Mat& **rvec**, Mat& **tvec**, bool **use_extrinsic_guess**=false, int **num_iters**=100, float **max_dist**=8.0, int **min_inlier_count**=100, vector<int>* **inliers**=NULL)

> **Parameters**
>
> > **object** – Single-row matrix of object points.
> >
> > **image** – Single-row matrix of image points.
> >
> > **camera_mat** – 3x3 matrix of intrinsic camera parameters.
> >
> > **dist_coef** – Distortion coefficients. See `undistortPoints()` for details.
> >
> > **rvec** – Output 3D rotation vector.
> >
> > **tvec** – Output 3D translation vector.
> >
> > **use_extrinsic_guess** – Flag to indicate that the function must use `rvec` and `tvec` as an initial transformation guess. It is not supported for now.
> >
> > **num_iters** – Maximum number of RANSAC iterations.
> >
> > **max_dist** – Euclidean distance threshold to detect whether point is inlier or not.
> >
> > **min_inlier_count** – Flag to indicate that the function must stop if greater or equal number of inliers is achieved. It is not supported for now.
> >
> > **inliers** – Output vector of inlier indices.

**See Also:**

`solvePnPRansac()`

## 11.12 Video Analysis

### gpu::BroxOpticalFlow

class gpu::**BroxOpticalFlow**

Class computing the optical flow for two images using Brox et al Optical Flow algorithm ([Brox2004]).

```cpp
class BroxOpticalFlow
{
public:
    BroxOpticalFlow(float alpha_, float gamma_, float scale_factor_, int inner_iterations_, int outer_iterations_, int

    //! Compute optical flow
    //! frame0 - source frame (supports only CV_32FC1 type)
    //! frame1 - frame to track (with the same size and type as frame0)
    //! u      - flow horizontal component (along x axis)
    //! v      - flow vertical component (along y axis)
    void operator ()(const GpuMat& frame0, const GpuMat& frame1, GpuMat& u, GpuMat& v, Stream& stream = Stream::Null()

    //! flow smoothness
    float alpha;

    //! gradient constancy importance
    float gamma;

    //! pyramid scale factor
    float scale_factor;

    //! number of lagged non-linearity iterations (inner loop)
    int inner_iterations;

    //! number of warping iterations (number of pyramid levels)
    int outer_iterations;

    //! number of linear system solver iterations
    int solver_iterations;

    GpuMat buf;
};
```

## gpu::GoodFeaturesToTrackDetector_GPU

**class** gpu::**GoodFeaturesToTrackDetector_GPU**

Class used for strong corners detection on an image.

```cpp
class GoodFeaturesToTrackDetector_GPU
{
public:
    explicit GoodFeaturesToTrackDetector_GPU(int maxCorners_ = 1000, double qualityLevel_ = 0.01, double minDistance_
        int blockSize_ = 3, bool useHarrisDetector_ = false, double harrisK_ = 0.04);

    void operator ()(const GpuMat& image, GpuMat& corners, const GpuMat& mask = GpuMat());

    int maxCorners;
    double qualityLevel;
    double minDistance;

    int blockSize;
    bool useHarrisDetector;
    double harrisK;

    void releaseMemory();
};
```

The class finds the most prominent corners in the image.

**See Also:**

`goodFeaturesToTrack()`

## gpu::GoodFeaturesToTrackDetector_GPU::GoodFeaturesToTrackDetector_GPU

Constructor.

**C++:** `gpu::GoodFeaturesToTrackDetector_GPU::`**`GoodFeaturesToTrackDetector_GPU`**(int **maxCorners**=1000, double **qualityLevel**=0.01, double **minDistance**=0.0, int **blockSize**=3, bool **useHarrisDetector**=false, double **harrisK**=0.04)

> **Parameters**
>
> > **maxCorners** – Maximum number of corners to return. If there are more corners than are found, the strongest of them is returned.
> >
> > **qualityLevel** – Parameter characterizing the minimal accepted quality of image corners. The parameter value is multiplied by the best corner quality measure, which is the minimal eigenvalue (see `gpu::cornerMinEigenVal()` ) or the Harris function response (see `gpu::cornerHarris()` ). The corners with the quality measure less than the product are rejected. For example, if the best corner has the quality measure = 1500, and the `qualityLevel=0.01` , then all the corners with the quality measure less than 15 are rejected.
> >
> > **minDistance** – Minimum possible Euclidean distance between the returned corners.
> >
> > **blockSize** – Size of an average block for computing a derivative covariation matrix over each pixel neighborhood. See `cornerEigenValsAndVecs()` .
> >
> > **useHarrisDetector** – Parameter indicating whether to use a Harris detector (see `gpu::cornerHarris()`) or `gpu::cornerMinEigenVal()`.
> >
> > **harrisK** – Free parameter of the Harris detector.

## gpu::GoodFeaturesToTrackDetector_GPU::operator ()

Finds the most prominent corners in the image.

**C++:** void `gpu::GoodFeaturesToTrackDetector_GPU::`**`operator()`**(const GpuMat& **image**, GpuMat& **corners**, const GpuMat& **mask**=GpuMat())

> **Parameters**
>
> > **image** – Input 8-bit, single-channel image.

> **corners** – Output vector of detected corners (it will be one row matrix with CV_32FC2 type).
>
> **mask** – Optional region of interest. If the image is not empty (it needs to have the type CV_8UC1 and the same size as `image` ), it specifies the region in which the corners are detected.

**See Also:**

goodFeaturesToTrack()

## gpu::GoodFeaturesToTrackDetector_GPU::releaseMemory

Releases inner buffers memory.

**C++:** void gpu::GoodFeaturesToTrackDetector_GPU::**releaseMemory**()

## gpu::FarnebackOpticalFlow

**class** gpu::**FarnebackOpticalFlow**

Class computing a dense optical flow using the Gunnar Farneback's algorithm.

```cpp
class CV_EXPORTS FarnebackOpticalFlow
{
public:
    FarnebackOpticalFlow()
    {
        numLevels = 5;
        pyrScale = 0.5;
        fastPyramids = false;
        winSize = 13;
        numIters = 10;
        polyN = 5;
        polySigma = 1.1;
        flags = 0;
    }

    int numLevels;
    double pyrScale;
    bool fastPyramids;
    int winSize;
    int numIters;
    int polyN;
    double polySigma;
    int flags;

    void operator ()(const GpuMat &frame0, const GpuMat &frame1, GpuMat &flowx, GpuMat &flowy, Stream &s = Stream::Nul

    void releaseMemory();

private:
    /* hidden */
};
```

## gpu::FarnebackOpticalFlow::operator ()

Computes a dense optical flow using the Gunnar Farneback's algorithm.

**C++:** void gpu::FarnebackOpticalFlow::**operator()**(const GpuMat& **frame0**, const GpuMat& **frame1**, GpuMat& **flowx**, GpuMat& **flowy**, Stream& **s**=Stream::Null())

> Parameters
>
>> **frame0** – First 8-bit gray-scale input image
>>
>> **frame1** – Second 8-bit gray-scale input image
>>
>> **flowx** – Flow horizontal component
>>
>> **flowy** – Flow vertical component
>>
>> **s** – Stream

**See Also:**

`calcOpticalFlowFarneback()`

## gpu::FarnebackOpticalFlow::releaseMemory

Releases unused auxiliary memory buffers.

**C++:** void gpu::FarnebackOpticalFlow::**releaseMemory**()

## gpu::PyrLKOpticalFlow

**class** gpu::**PyrLKOpticalFlow**

Class used for calculating an optical flow.

```cpp
class PyrLKOpticalFlow
{
public:
    PyrLKOpticalFlow();

    void sparse(const GpuMat& prevImg, const GpuMat& nextImg, const GpuMat& prevPts, GpuMat& nextPts,
        GpuMat& status, GpuMat* err = 0);

    void dense(const GpuMat& prevImg, const GpuMat& nextImg, GpuMat& u, GpuMat& v, GpuMat* err = 0);

    Size winSize;
    int maxLevel;
    int iters;
    bool useInitialFlow;

    void releaseMemory();
};
```

The class can calculate an optical flow for a sparse feature set or dense optical flow using the iterative Lucas-Kanade method with pyramids.

**See Also:**

`calcOpticalFlowPyrLK()`

## gpu::PyrLKOpticalFlow::sparse

Calculate an optical flow for a sparse feature set.

**C++:** void gpu::PyrLKOpticalFlow::**sparse**(const GpuMat& **prevImg**, const GpuMat& **nextImg**, const GpuMat& **prevPts**, GpuMat& **nextPts**, GpuMat& **status**, GpuMat* **err**=0)

> **Parameters**

>> **prevImg** – First 8-bit input image (supports both grayscale and color images).

>> **nextImg** – Second input image of the same size and the same type as `prevImg` .

>> **prevPts** – Vector of 2D points for which the flow needs to be found. It must be one row matrix with CV_32FC2 type.

>> **nextPts** – Output vector of 2D points (with single-precision floating-point coordinates) containing the calculated new positions of input features in the second image. When `useInitialFlow` is true, the vector must have the same size as in the input.

>> **status** – Output status vector (CV_8UC1 type). Each element of the vector is set to 1 if the flow for the corresponding features has been found. Otherwise, it is set to 0.

>> **err** – Output vector (CV_32FC1 type) that contains the difference between patches around the original and moved points or min eigen value if `getMinEigenVals` is checked. It can be NULL, if not needed.

**See Also:**

`calcOpticalFlowPyrLK()`

## gpu::PyrLKOpticalFlow::dense

Calculate dense optical flow.

**C++:** void gpu::PyrLKOpticalFlow::**dense**(const GpuMat& **prevImg**, const GpuMat& **nextImg**, GpuMat& **u**, GpuMat& **v**, GpuMat* **err**=0)

> **Parameters**

>> **prevImg** – First 8-bit grayscale input image.

>> **nextImg** – Second input image of the same size and the same type as `prevImg` .

>> **u** – Horizontal component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

>> **v** – Vertical component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

>> **err** – Output vector (CV_32FC1 type) that contains the difference between patches around the original and moved points or min eigen value if `getMinEigenVals` is checked. It can be NULL, if not needed.

## gpu::PyrLKOpticalFlow::releaseMemory

Releases inner buffers memory.

**C++:** void gpu::PyrLKOpticalFlow::**releaseMemory**()

---

## gpu::interpolateFrames

Interpolates frames (images) using provided optical flow (displacement field).

**C++:** void gpu::**interpolateFrames**(const GpuMat& **frame0**, const GpuMat& **frame1**, const GpuMat& **fu**, const GpuMat& **fv**, const GpuMat& **bu**, const GpuMat& **bv**, float **pos**, GpuMat& **newFrame**, GpuMat& **buf**, Stream& **stream**=Stream::Null())

> **Parameters**
>
>> **frame0** – First frame (32-bit floating point images, single channel).
>>
>> **frame1** – Second frame. Must have the same type and size as `frame0` .
>>
>> **fu** – Forward horizontal displacement.
>>
>> **fv** – Forward vertical displacement.
>>
>> **bu** – Backward horizontal displacement.
>>
>> **bv** – Backward vertical displacement.
>>
>> **pos** – New frame position.
>>
>> **newFrame** – Output image.
>>
>> **buf** – Temporary buffer, will have width x 6*height size, CV_32FC1 type and contain 6 GpuMat: occlusion masks for first frame, occlusion masks for second, interpolated forward horizontal flow, interpolated forward vertical flow, interpolated backward horizontal flow, interpolated backward vertical flow.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::FGDStatModel

**class** gpu::**FGDStatModel**

Class used for background/foreground segmentation.

```
class FGDStatModel
{
public:
    struct Params
    {
        ...
    };

    explicit FGDStatModel(int out_cn = 3);
    explicit FGDStatModel(const cv::gpu::GpuMat& firstFrame, const Params& params = Params(), int out_cn = 3);

    ~FGDStatModel();

    void create(const cv::gpu::GpuMat& firstFrame, const Params& params = Params());
    void release();

    int update(const cv::gpu::GpuMat& curFrame);

    //8UC3 or 8UC4 reference background image
    cv::gpu::GpuMat background;

    //8UC1 foreground image
```

```
    cv::gpu::GpuMat foreground;

    std::vector< std::vector<cv::Point> > foreground_regions;
};
```

The class discriminates between foreground and background pixels by building and maintaining a model of the background

The results are available through the class fields:

  .. ocv:member:: cv::gpu::GpuMat background

    The output background image.

  .. ocv:member:: cv::gpu::GpuMat foreground

    The output foreground mask as an 8-bit binary image.

  .. ocv:member:: cv::gpu::GpuMat foreground_regions

    The output foreground regions calculated by :ocv:func:'findContours'.

## gpu::FGDStatModel::FGDStatModel

Constructors.

**C++:** gpu::FGDStatModel::**FGDStatModel**(int **out_cn**=3)

**C++:** gpu::FGDStatModel::**FGDStatModel**(const cv::gpu::GpuMat& **firstFrame**, const Params& **params**=Params(), int **out_cn**=3)

      **Parameters**

           **firstFrame** – First frame from video stream. Supports 3- and 4-channels input ( CV_8UC3 and CV_8UC4 ).

           **params** – Algorithm's parameters. See [FGD2003] for explanation.

           **out_cn** – Channels count in output result and inner buffers. Can be 3 or 4. 4-channels version requires more memory, but works a bit faster.

**See Also:**

gpu::FGDStatModel::create()

## gpu::FGDStatModel::create

Initializes background model.

**C++:** void gpu::FGDStatModel::**create**(const cv::gpu::GpuMat& **firstFrame**, const Params& **params**=Params())

      **Parameters**

           **firstFrame** – First frame from video stream. Supports 3- and 4-channels input ( CV_8UC3 and CV_8UC4 ).

           **params** – Algorithm's parameters. See [FGD2003] for explanation.

## gpu::FGDStatModel::release

Releases all inner buffer's memory.

**C++:** void gpu::FGDStatModel::**release**()

## gpu::FGDStatModel::update

Updates the background model and returns foreground regions count.

**C++:** int gpu::FGDStatModel::**update**(const cv::gpu::GpuMat& **curFrame**)

>> **Parameters**

>>> **curFrame** – Next video frame.

## gpu::MOG_GPU

**class** gpu::**MOG_GPU**

Gaussian Mixture-based Backbround/Foreground Segmentation Algorithm.

```cpp
class MOG_GPU
{
public:
    MOG_GPU(int nmixtures = -1);

    void initialize(Size frameSize, int frameType);

    void operator()(const GpuMat& frame, GpuMat& fgmask, float learningRate = 0.0f, Stream& stream = Stream::Null());

    void getBackgroundImage(GpuMat& backgroundImage, Stream& stream = Stream::Null()) const;

    void release();

    int history;
    float varThreshold;
    float backgroundRatio;
    float noiseSigma;
};
```

The class discriminates between foreground and background pixels by building and maintaining a model of the background. Any pixel which does not fit this model is then deemed to be foreground. The class implements algorithm described in [MOG2001].

**See Also:**

BackgroundSubtractorMOG

## gpu::MOG_GPU::MOG_GPU

The constructor.

**C++:** gpu::MOG_GPU::**MOG_GPU**(int **nmixtures**=-1)

>> **Parameters**

>>> **nmixtures** – Number of Gaussian mixtures.

Default constructor sets all parameters to default values.

## gpu::MOG_GPU::operator()

Updates the background model and returns the foreground mask.

**C++:** void gpu::MOG_GPU::**operator()**(const GpuMat& **frame**, GpuMat& **fgmask**, float **learningRate**=0.0f, Stream& **stream**=Stream::Null())

> **Parameters**
>
> > **frame** – Next video frame.
> >
> > **fgmask** – The output foreground mask as an 8-bit binary image.
> >
> > **stream** – Stream for the asynchronous version.

## gpu::MOG_GPU::getBackgroundImage

Computes a background image.

**C++:** void gpu::MOG_GPU::**getBackgroundImage**(GpuMat& **backgroundImage**, Stream& **stream**=Stream::Null()) const

> **Parameters**
>
> > **backgroundImage** – The output background image.
> >
> > **stream** – Stream for the asynchronous version.

## gpu::MOG_GPU::release

Releases all inner buffer's memory.

**C++:** void gpu::MOG_GPU::**release**()

## gpu::MOG2_GPU

**class** gpu::**MOG2_GPU**

Gaussian Mixture-based Background/Foreground Segmentation Algorithm.

```
class MOG2_GPU
{
public:
    MOG2_GPU(int nmixtures = -1);

    void initialize(Size frameSize, int frameType);

    void operator()(const GpuMat& frame, GpuMat& fgmask, float learningRate = 0.0f, Stream& stream = Stream::Null())

    void getBackgroundImage(GpuMat& backgroundImage, Stream& stream = Stream::Null()) const;

    void release();

    // parameters
    ...
};
```

The class discriminates between foreground and background pixels by building and maintaining a model of the background

Here are important members of the class that control the algorithm, which you can set after constructing the class ins

  .. ocv:member:: float backgroundRatio

    Threshold defining whether the component is significant enough to be included into the background model ( corres

  .. ocv:member:: float varThreshold

    Threshold for the squared Mahalanobis distance that helps decide when a sample is close to the existing componen

  .. ocv:member:: float fVarInit

    Initial variance for the newly generated components. It affects the speed of adaptation. The parameter value is

  .. ocv:member:: float fVarMin

    Parameter used to further control the variance.

  .. ocv:member:: float fVarMax

    Parameter used to further control the variance.

  .. ocv:member:: float fCT

    Complexity reduction parameter. This parameter defines the number of samples needed to accept to prove the compo

  .. ocv:member:: uchar nShadowDetection

    The value for marking shadow pixels in the output foreground mask. Default value is 127.

  .. ocv:member:: float fTau

    Shadow threshold. The shadow is detected if the pixel is a darker version of the background. ''Tau'' is a thresh

  .. ocv:member:: bool bShadowDetection

    Parameter defining whether shadow detection should be enabled.

**See Also:**

`BackgroundSubtractorMOG2`

## gpu::MOG2_GPU::MOG2_GPU

The constructor.

**C++:** gpu::MOG2_GPU::**MOG2_GPU**(int **nmixtures**=-1)

> **Parameters**
>
> > **nmixtures** – Number of Gaussian mixtures.

Default constructor sets all parameters to default values.

## gpu::MOG2_GPU::operator()

Updates the background model and returns the foreground mask.

**C++:** void gpu::MOG2_GPU::**operator()**(const GpuMat& **frame**, GpuMat& **fgmask**, float **learningRate**=-1.0f, Stream& **stream**=Stream::Null() )

> **Parameters**
>
>> **frame** – Next video frame.
>>
>> **fgmask** – The output foreground mask as an 8-bit binary image.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::MOG2_GPU::getBackgroundImage

Computes a background image.

**C++:** void gpu::MOG2_GPU::**getBackgroundImage**(GpuMat& **backgroundImage**, Stream& **stream**=Stream::Null()) const

> **Parameters**
>
>> **backgroundImage** – The output background image.
>>
>> **stream** – Stream for the asynchronous version.

## gpu::MOG2_GPU::release

Releases all inner buffer's memory.

**C++:** void gpu::MOG2_GPU::**release**()

## gpu::VIBE_GPU

**class** gpu::**VIBE_GPU**

Class used for background/foreground segmentation.

```
class VIBE_GPU
{
public:
    explicit VIBE_GPU(unsigned long rngSeed = 1234567);

    void initialize(const GpuMat& firstFrame, Stream& stream = Stream::Null());

    void operator()(const GpuMat& frame, GpuMat& fgmask, Stream& stream = Stream::Null());

    void release();

    ...
};
```

The class discriminates between foreground and background pixels by building and maintaining a model of the background. Any pixel which does not fit this model is then deemed to be foreground. The class implements algorithm described in [VIBE2011].

## gpu::VIBE_GPU::VIBE_GPU

The constructor.

**C++:** gpu::VIBE_GPU::**VIBE_GPU**(unsigned long **rngSeed**=1234567)

>      Parameters

>           **rngSeed** – Value used to initiate a random sequence.

Default constructor sets all parameters to default values.

## gpu::VIBE_GPU::initialize

Initialize background model and allocates all inner buffers.

**C++:** void gpu::VIBE_GPU::**initialize**(const GpuMat& **firstFrame**, Stream& **stream**=Stream::Null())

>      Parameters

>           **firstFrame** – First frame from video sequence.

>           **stream** – Stream for the asynchronous version.

## gpu::VIBE_GPU::operator()

Updates the background model and returns the foreground mask

**C++:** void gpu::VIBE_GPU::**operator()**(const GpuMat& **frame**, GpuMat& **fgmask**, Stream& **stream**=Stream::Null())

>      Parameters

>           **frame** – Next video frame.

>           **fgmask** – The output foreground mask as an 8-bit binary image.

>           **stream** – Stream for the asynchronous version.

## gpu::VIBE_GPU::release

Releases all inner buffer's memory.

**C++:** void gpu::VIBE_GPU::**release**()

## gpu::GMG_GPU

**class** gpu::**GMG_GPU**

>     Class used for background/foreground segmentation.

```
class GMG_GPU_GPU
{
public:
    GMG_GPU();

    void initialize(Size frameSize, float min = 0.0f, float max = 255.0f);

    void operator ()(const GpuMat& frame, GpuMat& fgmask, float learningRate = -1.0f, Stream& stream = Stream::Nu
```

```
    void release();

    int    maxFeatures;
    float  learningRate;
    int    numInitializationFrames;
    int    quantizationLevels;
    float  backgroundPrior;
    float  decisionThreshold;
    int    smoothingRadius;

    ...
};
```

The class discriminates between foreground and background pixels by building and maintaining a model of the background. Any pixel which does not fit this model is then deemed to be foreground. The class implements algorithm described in [GMG2012].

Here are important members of the class that control the algorithm, which you can set after constructing the class instance:

> int **maxFeatures**
>> Total number of distinct colors to maintain in histogram.

> float **learningRate**
>> Set between 0.0 and 1.0, determines how quickly features are "forgotten" from histograms.

> int **numInitializationFrames**
>> Number of frames of video to use to initialize histograms.

> int **quantizationLevels**
>> Number of discrete levels in each channel to be used in histograms.

> float **backgroundPrior**
>> Prior probability that any given pixel is a background pixel. A sensitivity parameter.

> float **decisionThreshold**
>> Value above which pixel is determined to be FG.

> float **smoothingRadius**
>> Smoothing radius, in pixels, for cleaning up FG image.

## gpu::GMG_GPU::GMG_GPU

The default constructor.

**C++:** gpu::GMG_GPU::**GMG_GPU**()

Default constructor sets all parameters to default values.

## gpu::GMG_GPU::initialize

Initialize background model and allocates all inner buffers.

**C++:** void gpu::GMG_GPU::**initialize**(Size **frameSize**, float **min**=0.0f, float **max**=255.0f)

> **Parameters**

>> **frameSize** – Input frame size.

>> **min** – Minimum value taken on by pixels in image sequence. Usually 0.

> **max** – Maximum value taken on by pixels in image sequence, e.g. 1.0 or 255.

## gpu::GMG_GPU::operator()

Updates the background model and returns the foreground mask

**C++:** void gpu::GMG_GPU::**operator()**(const GpuMat& **frame**, GpuMat& **fgmask**, float **learningRate**=-1.0f, Stream& **stream**=Stream::Null() )

> **Parameters**
>
> > **frame** – Next video frame.
> >
> > **fgmask** – The output foreground mask as an 8-bit binary image.
> >
> > **stream** – Stream for the asynchronous version.

## gpu::GMG_GPU::release

Releases all inner buffer's memory.

**C++:** void gpu::GMG_GPU::**release**()

## gpu::VideoWriter_GPU

Video writer class.

**class** gpu::**VideoWriter_GPU**

The class uses H264 video codec.

---

**Note:** Currently only Windows platform is supported.

---

## gpu::VideoWriter_GPU::VideoWriter_GPU

Constructors.

**C++:** gpu::VideoWriter_GPU::**VideoWriter_GPU**()

**C++:** gpu::VideoWriter_GPU::**VideoWriter_GPU**(const std::string& **fileName**, cv::Size **frameSize**, double **fps**, SurfaceFormat **format**=SF_BGR )

**C++:** gpu::VideoWriter_GPU::**VideoWriter_GPU**(const std::string& **fileName**, cv::Size **frameSize**, double **fps**, const EncoderParams& **params**, SurfaceFormat **format**=SF_BGR )

**C++:** gpu::VideoWriter_GPU::**VideoWriter_GPU**(const cv::Ptr<EncoderCallBack>& **encoderCallback**, cv::Size **frameSize**, double **fps**, SurfaceFormat **format**=SF_BGR )

**C++:** gpu::VideoWriter_GPU::**VideoWriter_GPU**(const cv::Ptr<EncoderCallBack>& **encoderCallback**, cv::Size **frameSize**, double **fps**, const EncoderParams& **params**, SurfaceFormat **format**=SF_BGR )

> **Parameters**
>
> > **fileName** – Name of the output video file. Only AVI file format is supported.

> **frameSize** – Size of the input video frames.
>
> **fps** – Framerate of the created video stream.
>
> **params** – Encoder parameters. See `gpu::VideoWriter_GPU::EncoderParams` .
>
> **format** – Surface format of input frames ( `SF_UYVY` , `SF_YUY2` , `SF_YV12` , `SF_NV12` , `SF_IYUV` , `SF_BGR` or `SF_GRAY`). BGR or gray frames will be converted to YV12 format before encoding, frames with other formats will be used as is.
>
> **encoderCallback** – Callbacks for video encoder. See `gpu::VideoWriter_GPU::EncoderCallBack` . Use it if you want to work with raw video stream.

The constructors initialize video writer. FFMPEG is used to write videos. User can implement own multiplexing with `gpu::VideoWriter_GPU::EncoderCallBack` .

## gpu::VideoWriter_GPU::open

Initializes or reinitializes video writer.

**C++:** void gpu::VideoWriter_GPU::**open**(const std::string& **fileName**, cv::Size **frameSize**, double **fps**, SurfaceFormat **format**=SF_BGR)

**C++:** void gpu::VideoWriter_GPU::**open**(const std::string& **fileName**, cv::Size **frameSize**, double **fps**, const EncoderParams& **params**, SurfaceFormat **format**=SF_BGR)

**C++:** void gpu::VideoWriter_GPU::**open**(const cv::Ptr<EncoderCallBack>& **encoderCallback**, cv::Size **frameSize**, double **fps**, SurfaceFormat **format**=SF_BGR)

**C++:** void gpu::VideoWriter_GPU::**open**(const cv::Ptr<EncoderCallBack>& **encoderCallback**, cv::Size **frameSize**, double **fps**, const EncoderParams& **params**, SurfaceFormat **format**=SF_BGR)

The method opens video writer. Parameters are the same as in the constructor `gpu::VideoWriter_GPU::VideoWriter_GPU()` . The method throws `Exception` if error occurs.

## gpu::VideoWriter_GPU::isOpened

Returns true if video writer has been successfully initialized.

**C++:** bool gpu::VideoWriter_GPU::**isOpened**() const

## gpu::VideoWriter_GPU::close

Releases the video writer.

**C++:** void gpu::VideoWriter_GPU::**close**()

## gpu::VideoWriter_GPU::write

Writes the next video frame.

**C++:** void gpu::VideoWriter_GPU::**write**(const cv::gpu::GpuMat& **image**, bool **lastFrame**=false)

> **Parameters**
>
> > **image** – The written frame.

> **lastFrame** – Indicates that it is end of stream. The parameter can be ignored.

The method write the specified image to video file. The image must have the same size and the same surface format as has been specified when opening the video writer.

## gpu::VideoWriter_GPU::EncoderParams

**struct** gpu::VideoWriter_GPU::**EncoderParams**

Different parameters for CUDA video encoder.

```
struct EncoderParams
{
    int       P_Interval;      //   NVVE_P_INTERVAL,
    int       IDR_Period;      //   NVVE_IDR_PERIOD,
    int       DynamicGOP;      //   NVVE_DYNAMIC_GOP,
    int       RCType;          //   NVVE_RC_TYPE,
    int       AvgBitrate;      //   NVVE_AVG_BITRATE,
    int       PeakBitrate;     //   NVVE_PEAK_BITRATE,
    int       QP_Level_Intra;  //   NVVE_QP_LEVEL_INTRA,
    int       QP_Level_InterP; //   NVVE_QP_LEVEL_INTER_P,
    int       QP_Level_InterB; //   NVVE_QP_LEVEL_INTER_B,
    int       DeblockMode;     //   NVVE_DEBLOCK_MODE,
    int       ProfileLevel;    //   NVVE_PROFILE_LEVEL,
    int       ForceIntra;      //   NVVE_FORCE_INTRA,
    int       ForceIDR;        //   NVVE_FORCE_IDR,
    int       ClearStat;       //   NVVE_CLEAR_STAT,
    int       DIMode;          //   NVVE_SET_DEINTERLACE,
    int       Presets;         //   NVVE_PRESETS,
    int       DisableCabac;    //   NVVE_DISABLE_CABAC,
    int       NaluFramingType; //   NVVE_CONFIGURE_NALU_FRAMING_TYPE
    int       DisableSPSPPS;   //   NVVE_DISABLE_SPS_PPS

    EncoderParams();
    explicit EncoderParams(const std::string& configFile);

    void load(const std::string& configFile);
    void save(const std::string& configFile) const;
};
```

## gpu::VideoWriter_GPU::EncoderParams::EncoderParams

Constructors.

**C++:** gpu::VideoWriter_GPU::EncoderParams::**EncoderParams**()

**C++:** gpu::VideoWriter_GPU::EncoderParams::**EncoderParams**(const std::string& **configFile**)

> **Parameters**

> > **configFile** – Config file name.

Creates default parameters or reads parameters from config file.

## gpu::VideoWriter_GPU::EncoderParams::load

Reads parameters from config file.

**C++:** void gpu::VideoWriter_GPU::EncoderParams::**load**(const std::string& **configFile**)

> Parameters
>
> > **configFile** – Config file name.

## gpu::VideoWriter_GPU::EncoderParams::save

Saves parameters to config file.

**C++:** void gpu::VideoWriter_GPU::EncoderParams::**save**(const std::string& **configFile**) const

> Parameters
>
> > **configFile** – Config file name.

## gpu::VideoWriter_GPU::EncoderCallBack

**class** gpu::VideoWriter_GPU::**EncoderCallBack**

Callbacks for CUDA video encoder.

```cpp
class EncoderCallBack
{
public:
    enum PicType
    {
        IFRAME = 1,
        PFRAME = 2,
        BFRAME = 3
    };

    virtual ~EncoderCallBack() {}

    virtual unsigned char* acquireBitStream(int* bufferSize) = 0;
    virtual void releaseBitStream(unsigned char* data, int size) = 0;
    virtual void onBeginFrame(int frameNumber, PicType picType) = 0;
    virtual void onEndFrame(int frameNumber, PicType picType) = 0;
};
```

## gpu::VideoWriter_GPU::EncoderCallBack::acquireBitStream

Callback function to signal the start of bitstream that is to be encoded.

**C++: virtual** uchar* gpu::VideoWriter_GPU::EncoderCallBack::**acquireBitStream**(int* **buffer-Size = 0**)

Callback must allocate buffer for CUDA encoder and return pointer to it and it's size.

## gpu::VideoWriter_GPU::EncoderCallBack::releaseBitStream

Callback function to signal that the encoded bitstream is ready to be written to file.

**C++: virtual** void gpu::VideoWriter_GPU::EncoderCallBack::**releaseBitStream**(unsigned char* **data**, int **size = 0**)

## gpu::VideoWriter_GPU::EncoderCallBack::onBeginFrame

Callback function to signal that the encoding operation on the frame has started.

**C++: virtual** void gpu::VideoWriter_GPU::EncoderCallBack::**onBeginFrame**(int **frameNumber**, Pic-
Type **picType = 0**)

> **Parameters**
>
> > **picType** – Specify frame type (I-Frame, P-Frame or B-Frame).

## gpu::VideoWriter_GPU::EncoderCallBack::onEndFrame

Callback function signals that the encoding operation on the frame has finished.

**C++: virtual** void gpu::VideoWriter_GPU::EncoderCallBack::**onEndFrame**(int **frameNumber**, Pic-
Type **picType = 0**)

> **Parameters**
>
> > **picType** – Specify frame type (I-Frame, P-Frame or B-Frame).

## gpu::VideoReader_GPU

Class for reading video from files.

**class** gpu::**VideoReader_GPU**

---

**Note:** Currently only Windows and Linux platforms are supported.

---

## gpu::VideoReader_GPU::Codec

Video codecs supported by gpu::VideoReader_GPU .

```
enum Codec
{
    MPEG1 = 0,
    MPEG2,
    MPEG4,
    VC1,
    H264,
    JPEG,
    H264_SVC,
    H264_MVC,

    Uncompressed_YUV420 = (('I'<<24)|('Y'<<16)|('U'<<8)|('V')),   // Y,U,V (4:2:0)
    Uncompressed_YV12   = (('Y'<<24)|('V'<<16)|('1'<<8)|('2')),   // Y,V,U (4:2:0)
    Uncompressed_NV12   = (('N'<<24)|('V'<<16)|('1'<<8)|('2')),   // Y,UV  (4:2:0)
    Uncompressed_YUYV   = (('Y'<<24)|('U'<<16)|('Y'<<8)|('V')),   // YUYV/YUY2 (4:2:2)
    Uncompressed_UYVY   = (('U'<<24)|('Y'<<16)|('V'<<8)|('Y')),   // UYVY (4:2:2)
};
```

## gpu::VideoReader_GPU::ChromaFormat

Chroma formats supported by `gpu::VideoReader_GPU`.

```
enum ChromaFormat
{
    Monochrome=0,
    YUV420,
    YUV422,
    YUV444,
};
```

## gpu::VideoReader_GPU::FormatInfo

**struct** gpu::VideoReader_GPU::**FormatInfo**

Struct providing information about video file format.

```
struct FormatInfo
{
    Codec codec;
    ChromaFormat chromaFormat;
    int width;
    int height;
};
```

## gpu::VideoReader_GPU::VideoReader_GPU

Constructors.

**C++:** gpu::VideoReader_GPU::**VideoReader_GPU**()

**C++:** gpu::VideoReader_GPU::**VideoReader_GPU**(const std::string& **filename**)

**C++:** gpu::VideoReader_GPU::**VideoReader_GPU**(const cv::Ptr<VideoSource>& **source**)

> **Parameters**
>
> > **filename** – Name of the input video file.
> >
> > **source** – Video file parser implemented by user.

The constructors initialize video reader. FFMPEG is used to read videos. User can implement own demultiplexing with `gpu::VideoReader_GPU::VideoSource`.

## gpu::VideoReader_GPU::open

Initializes or reinitializes video reader.

**C++:** void gpu::VideoReader_GPU::**open**(const std::string& **filename**)

**C++:** void gpu::VideoReader_GPU::**open**(const cv::Ptr<VideoSource>& **source**)

The method opens video reader. Parameters are the same as in the constructor `gpu::VideoReader_GPU::VideoReader_GPU()`. The method throws `Exception` if error occurs.

## gpu::VideoReader_GPU::isOpened

Returns true if video reader has been successfully initialized.

**C++:** bool gpu::VideoReader_GPU::**isOpened**() const


## gpu::VideoReader_GPU::close

Releases the video reader.

**C++:** void gpu::VideoReader_GPU::**close**()


## gpu::VideoReader_GPU::read

Grabs, decodes and returns the next video frame.

**C++:** bool gpu::VideoReader_GPU::**read**(GpuMat& **image**)

If no frames has been grabbed (there are no more frames in video file), the methods return `false` . The method throws `Exception` if error occurs.


## gpu::VideoReader_GPU::format

Returns information about video file format.

**C++:** FormatInfo gpu::VideoReader_GPU::**format**() const

The method throws `Exception` if video reader wasn't initialized.


## gpu::VideoReader_GPU::dumpFormat

Dump information about video file format to specified stream.

**C++:** void gpu::VideoReader_GPU::**dumpFormat**(std::ostream& **st**)

> **Parameters**
>
>> **st** – Output stream.

The method throws `Exception` if video reader wasn't initialized.


## gpu::VideoReader_GPU::VideoSource

**class** gpu::VideoReader_GPU::**VideoSource**

Interface for video demultiplexing.

```
class VideoSource
{
public:
    VideoSource();
    virtual ~VideoSource() {}

    virtual FormatInfo format() const = 0;
    virtual void start() = 0;
    virtual void stop() = 0;
```

```
    virtual bool isStarted() const = 0;
    virtual bool hasError() const = 0;

protected:
    bool parseVideoData(const unsigned char* data, size_t size, bool endOfStream = false);
};
```

User can implement own demultiplexing by implementing this interface.

## gpu::VideoReader_GPU::VideoSource::format

Returns information about video file format.

**C++: virtual** FormatInfo gpu::VideoReader_GPU::VideoSource::**format**( const = 0)

## gpu::VideoReader_GPU::VideoSource::start

Starts processing.

**C++: virtual** void gpu::VideoReader_GPU::VideoSource::**start**( = 0)

Implementation must create own thread with video processing and call periodic `gpu::VideoReader_GPU::VideoSource::parseVideoData()`.

## gpu::VideoReader_GPU::VideoSource::stop

Stops processing.

**C++: virtual** void gpu::VideoReader_GPU::VideoSource::**stop**( = 0)

## gpu::VideoReader_GPU::VideoSource::isStarted

Returns `true` if processing was successfully started.

**C++: virtual** bool gpu::VideoReader_GPU::VideoSource::**isStarted**( const = 0)

## gpu::VideoReader_GPU::VideoSource::hasError

Returns `true` if error occured during processing.

**C++: virtual** bool gpu::VideoReader_GPU::VideoSource::**hasError**( const = 0)

## gpu::VideoReader_GPU::VideoSource::parseVideoData

Parse next video frame. Implementation must call this method after new frame was grabbed.

**C++:** bool gpu::VideoReader_GPU::VideoSource::**parseVideoData**(const uchar\* **data**, size_t **size**, bool
**endOfStream**=false)

>        **Parameters**

>                **data** – Pointer to frame data. Can be NULL if `endOfStream` if `true` .

>                **size** – Size in bytes of current frame.

> **endOfStream** – Indicates that it is end of stream.

# PHOTO. COMPUTATIONAL PHOTOGRAPHY

## 12.1 Inpainting

### inpaint

Restores the selected region in an image using the region neighborhood.

**C++:** void **inpaint**(InputArray **src**, InputArray **inpaintMask**, OutputArray **dst**, double **inpaintRadius**, int **flags**)

**Python:** cv2.**inpaint**(src, inpaintMask, inpaintRadius, flags[, dst]) → dst

**C:** void **cvInpaint**(const CvArr* **src**, const CvArr* **inpaint_mask**, CvArr* **dst**, double **inpaintRange**, int **flags**)

**Python:** cv.**Inpaint**(src, mask, dst, inpaintRadius, flags) → None

> **Parameters**
>
> > **src** – Input 8-bit 1-channel or 3-channel image.
> >
> > **inpaintMask** – Inpainting mask, 8-bit 1-channel image. Non-zero pixels indicate the area that needs to be inpainted.
> >
> > **dst** – Output image with the same size and type as `src` .
> >
> > **inpaintRadius** – Radius of a circular neighborhood of each point inpainted that is considered by the algorithm.
> >
> > **flags** – Inpainting method that could be one of the following:
> >
> > – **INPAINT_NS** Navier-Stokes based method.
> >
> > – **INPAINT_TELEA** Method by Alexandru Telea [Telea04].

The function reconstructs the selected image area from the pixel near the area boundary. The function may be used to remove dust and scratches from a scanned photo, or to remove undesirable objects from still images or video. See http://en.wikipedia.org/wiki/Inpainting for more details.

## 12.2 Denoising

### fastNlMeansDenoising

Perform image denoising using Non-local Means Denoising algorithm http://www.ipol.im/pub/algo/bcm_non_local_means_denoising/ with several computational optimizations. Noise expected to be a gaussian white noise

**C++:** void **fastNlMeansDenoising**(InputArray **src**, OutputArray **dst**, float **h**=3, int **templateWindowSize**=7, int **searchWindowSize**=21 )

>    **Parameters**

>>    **src** – Input 8-bit 1-channel, 2-channel or 3-channel image.

>>    **dst** – Output image with the same size and type as `src` .

>>    **templateWindowSize** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels

>>    **searchWindowSize** – Size in pixels of the window that is used to compute weighted average for given pixel. Should be odd. Affect performance linearly: greater searchWindowsSize - greater denoising time. Recommended value 21 pixels

>>    **h** – Parameter regulating filter strength. Big h value perfectly removes noise but also removes image details, smaller h value preserves details but also preserves some noise

This function expected to be applied to grayscale images. For colored images look at `fastNlMeansDenoisingColored`. Advanced usage of this functions can be manual denoising of colored image in different colorspaces. Such approach is used in `fastNlMeansDenoisingColored` by converting image to CIELAB colorspace and then separately denoise L and AB components with different h parameter.

### fastNlMeansDenoisingColored

Modification of `fastNlMeansDenoising` function for colored images

**C++:** void **fastNlMeansDenoisingColored**(InputArray **src**, OutputArray **dst**, float **h**=3, float **hColor**=3, int **templateWindowSize**=7, int **searchWindowSize**=21 )

>    **Parameters**

>>    **src** – Input 8-bit 3-channel image.

>>    **dst** – Output image with the same size and type as `src` .

>>    **templateWindowSize** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels

>>    **searchWindowSize** – Size in pixels of the window that is used to compute weighted average for given pixel. Should be odd. Affect performance linearly: greater searchWindowsSize - greater denoising time. Recommended value 21 pixels

>>    **h** – Parameter regulating filter strength for luminance component. Bigger h value perfectly removes noise but also removes image details, smaller h value preserves details but also preserves some noise

>>    **hForColorComponents** – The same as h but for color components. For most images value equals 10 will be enought to remove colored noise and do not distort colors

The function converts image to CIELAB colorspace and then separately denoise L and AB components with given h parameters using `fastNlMeansDenoising` function.

## fastNlMeansDenoisingMulti

Modification of `fastNlMeansDenoising` function for images sequence where consecutive images have been captured in small period of time. For example video. This version of the function is for grayscale images or for manual manipulation with colorspaces. For more details see http://citeseerx.ist.psu.edu/viewdoc/summary?doi=10.1.1.131.6394

**C++:** void **fastNlMeansDenoisingMulti**(InputArrayOfArrays **srcImgs**, OutputArray **dst**, int **imgToDenoiseIndex**, int **temporalWindowSize**, float **h**=3, int **templateWindowSize**=7, int **searchWindowSize**=21 )

> **Parameters**
>
> > **srcImgs** – Input 8-bit 1-channel, 2-channel or 3-channel images sequence. All images should have the same type and size.
> >
> > **imgToDenoiseIndex** – Target image to denoise index in `srcImgs` sequence
> >
> > **temporalWindowSize** – Number of surrounding images to use for target image denoising. Should be odd. Images from imgToDenoiseIndex - `temporalWindowSize` / 2 to imgToDenoiseIndex - `temporalWindowSize` / 2 from `srcImgs` will be used to denoise `srcImgs[imgToDenoiseIndex]` image.
> >
> > **dst** – Output image with the same size and type as `srcImgs` images.
> >
> > **templateWindowSize** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels
> >
> > **searchWindowSize** – Size in pixels of the window that is used to compute weighted average for given pixel. Should be odd. Affect performance linearly: greater searchWindowsSize - greater denoising time. Recommended value 21 pixels
> >
> > **h** – Parameter regulating filter strength for luminance component. Bigger h value perfectly removes noise but also removes image details, smaller h value preserves details but also preserves some noise

## fastNlMeansDenoisingColoredMulti

Modification of `fastNlMeansDenoisingMulti` function for colored images sequences

**C++:** void **fastNlMeansDenoisingColoredMulti**(InputArrayOfArrays **srcImgs**, OutputArray **dst**, int **imgToDenoiseIndex**, int **temporalWindowSize**, float **h**=3, float **hColor**=3, int **templateWindowSize**=7, int **searchWindowSize**=21 )

> **Parameters**
>
> > **srcImgs** – Input 8-bit 3-channel images sequence. All images should have the same type and size.
> >
> > **imgToDenoiseIndex** – Target image to denoise index in `srcImgs` sequence
> >
> > **temporalWindowSize** – Number of surrounding images to use for target image denoising. Should be odd. Images from imgToDenoiseIndex - `temporalWindowSize` / 2 to imgToDenoiseIndex - `temporalWindowSize` / 2 from `srcImgs` will be used to denoise `srcImgs[imgToDenoiseIndex]` image.
> >
> > **dst** – Output image with the same size and type as `srcImgs` images.
> >
> > **templateWindowSize** – Size in pixels of the template patch that is used to compute weights. Should be odd. Recommended value 7 pixels

>   > **searchWindowSize** – Size in pixels of the window that is used to compute weighted average
>   > for given pixel. Should be odd. Affect performance linearly: greater searchWindowsSize -
>   > greater denoising time. Recommended value 21 pixels
>
>   > **h** – Parameter regulating filter strength for luminance component. Bigger h value perfectly
>   > removes noise but also removes image details, smaller h value preserves details but also
>   > preserves some noise.
>
>   > **hForColorComponents** – The same as h but for color components.

The function converts images to CIELAB colorspace and then separately denoise L and AB components with given h
parameters using `fastNlMeansDenoisingMulti` function.

# STITCHING. IMAGES STITCHING

## 13.1 Stitching Pipeline

This figure illustrates the stitching module pipeline implemented in the `Stitcher` class. Using that class it's possible to configure/remove some steps, i.e. adjust the stitching pipeline according to the particular needs. All building blocks from the pipeline are available in the `detail` namespace, one can combine and use them separately.

The implemented stitching pipeline is very similar to the one proposed in [BL07].

## 13.2 References

## 13.3 High Level Functionality

### Stitcher

**class Stitcher**

High level image stitcher. It's possible to use this class without being aware of the entire stitching pipeline. However, to be able to achieve higher stitching stability and quality of the final images at least being familiar with the theory is recommended (see *Stitching Pipeline*).

```cpp
class CV_EXPORTS Stitcher
{
public:
    enum { ORIG_RESOL = -1 };
    enum Status { OK, ERR_NEED_MORE_IMGS };

    // Creates stitcher with default parameters
    static Stitcher createDefault(bool try_use_gpu = false);

    Status estimateTransform(InputArray images);
    Status estimateTransform(InputArray images, const std::vector<std::vector<Rect> > &rois);

    Status composePanorama(OutputArray pano);
    Status composePanorama(InputArray images, OutputArray pano);

    Status stitch(InputArray images, OutputArray pano);
    Status stitch(InputArray images, const std::vector<std::vector<Rect> > &rois, OutputArray pano);

    double registrationResol() const { return registr_resol_; }
    void setRegistrationResol(double resol_mpx) { registr_resol_ = resol_mpx; }

    double seamEstimationResol() const { return seam_est_resol_; }
    void setSeamEstimationResol(double resol_mpx) { seam_est_resol_ = resol_mpx; }

    double compositingResol() const { return compose_resol_; }
    void setCompositingResol(double resol_mpx) { compose_resol_ = resol_mpx; }

    double panoConfidenceThresh() const { return conf_thresh_; }
    void setPanoConfidenceThresh(double conf_thresh) { conf_thresh_ = conf_thresh; }

    bool waveCorrection() const { return do_wave_correct_; }
    void setWaveCorrection(bool flag) { do_wave_correct_ = flag; }

    detail::WaveCorrectKind waveCorrectKind() const { return wave_correct_kind_; }
    void setWaveCorrectKind(detail::WaveCorrectKind kind) { wave_correct_kind_ = kind; }

    Ptr<detail::FeaturesFinder> featuresFinder() { return features_finder_; }
    const Ptr<detail::FeaturesFinder> featuresFinder() const { return features_finder_; }
    void setFeaturesFinder(Ptr<detail::FeaturesFinder> features_finder)
        { features_finder_ = features_finder; }

    Ptr<detail::FeaturesMatcher> featuresMatcher() { return features_matcher_; }
    const Ptr<detail::FeaturesMatcher> featuresMatcher() const { return features_matcher_; }
    void setFeaturesMatcher(Ptr<detail::FeaturesMatcher> features_matcher)
        { features_matcher_ = features_matcher; }
```

```
        const cv::Mat& matchingMask() const { return matching_mask_; }
        void setMatchingMask(const cv::Mat &mask)
        {
            CV_Assert(mask.type() == CV_8U && mask.cols == mask.rows);
            matching_mask_ = mask.clone();
        }

        Ptr<detail::BundleAdjusterBase> bundleAdjuster() { return bundle_adjuster_; }
        const Ptr<detail::BundleAdjusterBase> bundleAdjuster() const { return bundle_adjuster_; }
        void setBundleAdjuster(Ptr<detail::BundleAdjusterBase> bundle_adjuster)
            { bundle_adjuster_ = bundle_adjuster; }

        Ptr<WarperCreator> warper() { return warper_; }
        const Ptr<WarperCreator> warper() const { return warper_; }
        void setWarper(Ptr<WarperCreator> warper) { warper_ = warper; }

        Ptr<detail::ExposureCompensator> exposureCompensator() { return exposure_comp_; }
        const Ptr<detail::ExposureCompensator> exposureCompensator() const { return exposure_comp_; }
        void setExposureCompensator(Ptr<detail::ExposureCompensator> exposure_comp)
            { exposure_comp_ = exposure_comp; }

        Ptr<detail::SeamFinder> seamFinder() { return seam_finder_; }
        const Ptr<detail::SeamFinder> seamFinder() const { return seam_finder_; }
        void setSeamFinder(Ptr<detail::SeamFinder> seam_finder) { seam_finder_ = seam_finder; }

        Ptr<detail::Blender> blender() { return blender_; }
        const Ptr<detail::Blender> blender() const { return blender_; }
        void setBlender(Ptr<detail::Blender> blender) { blender_ = blender; }

private:
    /* hidden */
};
```

# Stitcher::createDefault

Creates a stitcher with the default parameters.

**C++:** Stitcher Stitcher::**createDefault**(bool **try_use_gpu**=false)

> **Parameters**
>
> > **try_use_gpu** – Flag indicating whether GPU should be used whenever it's possible.
>
> **Returns** Stitcher class instance.

# Stitcher::estimateTransform

These functions try to match the given images and to estimate rotations of each camera.

---

**Note:** Use the functions only if you're aware of the stitching pipeline, otherwise use `Stitcher::stitch()`.

---

**C++:** Status Stitcher::**estimateTransform**(InputArray **images**)

**C++:** Status Stitcher::**estimateTransform**(InputArray **images**, const std::vector<std::vector<Rect>>& **rois**)

> **Parameters**
>
> > **images** – Input images.
> >
> > **rois** – Region of interest rectangles.
>
> **Returns** Status code.

## Stitcher::composePanorama

These functions try to compose the given images (or images stored internally from the other function calls) into the final pano under the assumption that the image transformations were estimated before.

---

**Note:** Use the functions only if you're aware of the stitching pipeline, otherwise use `Stitcher::stitch()`.

---

**C++:** Status `Stitcher::`**`composePanorama`**(OutputArray **pano**)

**C++:** Status `Stitcher::`**`composePanorama`**(InputArray **images**, OutputArray **pano**)

> **Parameters**
>
> > **images** – Input images.
> >
> > **pano** – Final pano.
>
> **Returns** Status code.

## Stitcher::stitch

These functions try to stitch the given images.

**C++:** Status `Stitcher::`**`stitch`**(InputArray **images**, OutputArray **pano**)

**C++:** Status `Stitcher::`**`stitch`**(InputArray **images**, const std::vector<std::vector<Rect>>& **rois**, OutputArray **pano**)

> **Parameters**
>
> > **images** – Input images.
> >
> > **rois** – Region of interest rectangles.
> >
> > **pano** – Final pano.
>
> **Returns** Status code.

## WarperCreator

class `WarperCreator`

Image warper factories base class.

```cpp
class WarperCreator
{
public:
    virtual ~WarperCreator() {}
    virtual Ptr<detail::RotationWarper> create(float scale) const = 0;
};
```

---

## PlaneWarper

**class PlaneWarper** : **public WarperCreator**

Plane warper factory class.

```
class PlaneWarper : public WarperCreator
{
public:
    Ptr<detail::RotationWarper> create(float scale) const { return new detail::PlaneWarper(scale); }
};
```

**See Also:**

detail::PlaneWarper

## CylindricalWarper

**class CylindricalWarper** : **public WarperCreator**

Cylindrical warper factory class.

```
class CylindricalWarper: public WarperCreator
{
public:
    Ptr<detail::RotationWarper> create(float scale) const { return new detail::CylindricalWarper(scale); }
};
```

**See Also:**

detail::CylindricalWarper

## SphericalWarper

**class SphericalWarper** : **public WarperCreator**

Spherical warper factory class.

```
class SphericalWarper: public WarperCreator
{
public:
    Ptr<detail::RotationWarper> create(float scale) const { return new detail::SphericalWarper(scale); }
};
```

**See Also:**

detail::SphericalWarper

# 13.4 Camera

## detail::CameraParams

**struct** detail::**CameraParams**

Describes camera parameters.

---

**Note:** Translation is assumed to be zero during the whole stitching pipeline.

---

```
struct CV_EXPORTS CameraParams
{
    CameraParams();
    CameraParams(const CameraParams& other);
    const CameraParams& operator =(const CameraParams& other);
    Mat K() const;

    double focal; // Focal length
    double aspect; // Aspect ratio
    double ppx; // Principal point X
    double ppy; // Principal point Y
    Mat R; // Rotation
    Mat t; // Translation
};
```

## 13.5 Features Finding and Images Matching

### detail::ImageFeatures

**struct** detail::**ImageFeatures**

Structure containing image keypoints and descriptors.

```
struct CV_EXPORTS ImageFeatures
{
    int img_idx;
    Size img_size;
    std::vector<KeyPoint> keypoints;
    Mat descriptors;
};
```

### detail::FeaturesFinder

**class** detail::**FeaturesFinder**

Feature finders base class.

```
class CV_EXPORTS FeaturesFinder
{
public:
    virtual ~FeaturesFinder() {}
    void operator ()(const Mat &image, ImageFeatures &features);
    void operator ()(const Mat &image, ImageFeatures &features, const std::vector<cv::Rect> &rois);
    virtual void collectGarbage() {}

protected:
    virtual void find(const Mat &image, ImageFeatures &features) = 0;
};
```

---

## detail::FeaturesFinder::operator()

Finds features in the given image.

**C++:** void detail::FeaturesFinder::**operator()**(const Mat& **image**, ImageFeatures& **features**)

**C++:** void detail::FeaturesFinder::**operator()**(const Mat& **image**, ImageFeatures& **features**, const std::vector<cv::Rect>& **rois**)

> Parameters
>
>> **image** – Source image
>>
>> **features** – Found features
>>
>> **rois** – Regions of interest

**See Also:**

detail::ImageFeatures, Rect_

## detail::FeaturesFinder::collectGarbage

Frees unused memory allocated before if there is any.

**C++:** void detail::FeaturesFinder::**collectGarbage**()

## detail::FeaturesFinder::find

This method must implement features finding logic in order to make the wrappers detail::FeaturesFinder::operator() work.

**C++:** void detail::FeaturesFinder::**find**(const Mat& **image**, ImageFeatures& **features**)

> Parameters
>
>> **image** – Source image
>>
>> **features** – Found features

**See Also:**

detail::ImageFeatures

## detail::SurfFeaturesFinder

**class** detail::**SurfFeaturesFinder** : **public** detail::**FeaturesFinder**

SURF features finder.

```cpp
class CV_EXPORTS SurfFeaturesFinder : public FeaturesFinder
{
public:
    SurfFeaturesFinder(double hess_thresh = 300., int num_octaves = 3, int num_layers = 4,
                       int num_octaves_descr = /*4*/3, int num_layers_descr = /*2*/4);

private:
    /* hidden */
};
```

**See Also:**

detail::FeaturesFinder, SURF

## detail::OrbFeaturesFinder

**class** detail::**OrbFeaturesFinder** : **public** detail::**FeaturesFinder**

ORB features finder.

```
class CV_EXPORTS OrbFeaturesFinder : public FeaturesFinder
{
public:
    OrbFeaturesFinder(Size _grid_size = Size(3,1), size_t n_features = 1500,
                      const ORB::CommonParams &detector_params = ORB::CommonParams(1.3f, 5));

private:
    /* hidden */
};
```

**See Also:**

detail::FeaturesFinder, ORB

## detail::MatchesInfo

**struct** detail::**MatchesInfo**

Structure containing information about matches between two images. It's assumed that there is a homography between those images.

```
struct CV_EXPORTS MatchesInfo
{
    MatchesInfo();
    MatchesInfo(const MatchesInfo &other);
    const MatchesInfo& operator =(const MatchesInfo &other);

    int src_img_idx, dst_img_idx;       // Images indices (optional)
    std::vector<DMatch> matches;
    std::vector<uchar> inliers_mask;    // Geometrically consistent matches mask
    int num_inliers;                    // Number of geometrically consistent matches
    Mat H;                              // Estimated homography
    double confidence;                  // Confidence two images are from the same panorama
};
```

## detail::FeaturesMatcher

**class** detail::**FeaturesMatcher**

Feature matchers base class.

```
class CV_EXPORTS FeaturesMatcher
{
public:
    virtual ~FeaturesMatcher() {}

    void operator ()(const ImageFeatures &features1, const ImageFeatures &features2,
```

```
                        MatchesInfo& matches_info) { match(features1, features2, matches_info); }

    void operator ()(const std::vector<ImageFeatures> &features, std::vector<MatchesInfo> &pairwise_matches,
                     const Mat &mask = cv::Mat());

    bool isThreadSafe() const { return is_thread_safe_; }

    virtual void collectGarbage() {}

protected:
    FeaturesMatcher(bool is_thread_safe = false) : is_thread_safe_(is_thread_safe) {}

    virtual void match(const ImageFeatures &features1, const ImageFeatures &features2,
                       MatchesInfo& matches_info) = 0;

    bool is_thread_safe_;
};
```

## detail::FeaturesMatcher::operator()

Performs images matching.

**C++:** void `detail::FeaturesMatcher::`**`operator()`**(const ImageFeatures& **features1**, const ImageFeatures& **features2**, MatchesInfo& **matches_info**)

> **Parameters**
>
>> **features1** – First image features
>>
>> **features2** – Second image features
>>
>> **matches_info** – Found matches

**C++:** void `detail::FeaturesMatcher::`**`operator()`**(const std::vector<ImageFeatures>& **features**, std::vector<MatchesInfo>& **pairwise_matches**, const Mat& **mask**=Mat() )

> **Parameters**
>
>> **features** – Features of the source images
>>
>> **pairwise_matches** – Found pairwise matches
>>
>> **mask** – Mask indicating which image pairs must be matched

The function is parallelized with the TBB library.

**See Also:**

`detail::MatchesInfo`

## detail::FeaturesMatcher::isThreadSafe

**C++:** bool `detail::FeaturesMatcher::`**`isThreadSafe`**() const

> **Returns** True, if it's possible to use the same matcher instance in parallel, false otherwise

## detail::FeaturesMatcher::collectGarbage

Frees unused memory allocated before if there is any.

**C++:** void detail::FeaturesMatcher::**collectGarbage**()

## detail::FeaturesMatcher::match

This method must implement matching logic in order to make the wrappers detail::FeaturesMatcher::operator() work.

**C++:** void detail::FeaturesMatcher::**match**(const ImageFeatures& **features1**, const ImageFeatures& **features2**, MatchesInfo& **matches_info**)

> Parameters
>
> > **features1** – first image features
> >
> > **features2** – second image features
> >
> > **matches_info** – found matches

## detail::BestOf2NearestMatcher

class detail::**BestOf2NearestMatcher** : **public** detail::**FeaturesMatcher**

Features matcher which finds two best matches for each feature and leaves the best one only if the ratio between descriptor distances is greater than the threshold match_conf.

```
class CV_EXPORTS BestOf2NearestMatcher : public FeaturesMatcher
{
public:
    BestOf2NearestMatcher(bool try_use_gpu = false, float match_conf = 0.65f,
                          int num_matches_thresh1 = 6, int num_matches_thresh2 = 6);

    void collectGarbage();

protected:
    void match(const ImageFeatures &features1, const ImageFeatures &features2, MatchesInfo &matches_info);

    int num_matches_thresh1_;
    int num_matches_thresh2_;
    Ptr<FeaturesMatcher> impl_;
};
```

**See Also:**

detail::FeaturesMatcher

## detail::BestOf2NearestMatcher::BestOf2NearestMatcher

Constructs a "best of 2 nearest" matcher.

**C++:** detail::BestOf2NearestMatcher::**BestOf2NearestMatcher**(bool **try_use_gpu**=false, float **match_conf**=0.3f, int **num_matches_thresh1**=6, int **num_matches_thresh2**=6)

> Parameters

**try_use_gpu** – Should try to use GPU or not

**match_conf** – Match distances ration threshold

**num_matches_thresh1** – Minimum number of matches required for the 2D projective transform estimation used in the inliers classification step

**num_matches_thresh2** – Minimum number of matches required for the 2D projective transform re-estimation on inliers

## 13.6 Rotation Estimation

### detail::Estimator

**class** detail::**Estimator**

Rotation estimator base class. It takes features of all images, pairwise matches between all images and estimates rotations of all cameras.

---

**Note:** The coordinate system origin is implementation-dependent, but you can always normalize the rotations in respect to the first camera, for instance.

---

```cpp
class CV_EXPORTS Estimator
{
public:
    virtual ~Estimator() {}

    void operator ()(const std::vector<ImageFeatures> &features, const std::vector<MatchesInfo> &pairwise_matches,
                     std::vector<CameraParams> &cameras)
        { estimate(features, pairwise_matches, cameras); }

protected:
    virtual void estimate(const std::vector<ImageFeatures> &features, const std::vector<MatchesInfo> &pairwise_matches,
                          std::vector<CameraParams> &cameras) = 0;
};
```

### detail::Estimator::operator()

Estimates camera parameters.

**C++:** detail::Estimator::**operator()**(const std::vector<ImageFeatures>& **features**, const std::vector<MatchesInfo>& **pairwise_matches**, std::vector<CameraParams>& **cameras**)

>   **Parameters**

>>      **features** – Features of images

>>      **pairwise_matches** – Pairwise matches of images

>>      **cameras** – Estimated camera parameters

## detail::Estimator::estimate

This method must implement camera parameters estimation logic in order to make the wrapper detail::Estimator::operator() work.

**C++:** void detail::Estimator::**estimate**(const std::vector<ImageFeatures>& **features**, const std::vector<MatchesInfo>& **pairwise_matches**, std::vector<CameraParams>& **cameras**)

> **Parameters**
>
> > **features** – Features of images
> >
> > **pairwise_matches** – Pairwise matches of images
> >
> > **cameras** – Estimated camera parameters

## detail::HomographyBasedEstimator

**class** detail::**HomographyBasedEstimator** : **public** detail::**Estimator**

Homography based rotation estimator.

```
class CV_EXPORTS HomographyBasedEstimator : public Estimator
{
public:
    HomographyBasedEstimator(bool is_focals_estimated = false)
        : is_focals_estimated_(is_focals_estimated) {}

private:
    /* hidden */
};
```

## detail::BundleAdjusterBase

**class** detail::**BundleAdjusterBase** : **public** detail::**Estimator**

Base class for all camera parameters refinement methods.

```
class CV_EXPORTS BundleAdjusterBase : public Estimator
{
public:
    const Mat refinementMask() const { return refinement_mask_.clone(); }
    void setRefinementMask(const Mat &mask)
    {
        CV_Assert(mask.type() == CV_8U && mask.size() == Size(3, 3));
        refinement_mask_ = mask.clone();
    }

    double confThresh() const { return conf_thresh_; }
    void setConfThresh(double conf_thresh) { conf_thresh_ = conf_thresh; }

    CvTermCriteria termCriteria() { return term_criteria_; }
    void setTermCriteria(const CvTermCriteria& term_criteria) { term_criteria_ = term_criteria; }

protected:
    BundleAdjusterBase(int num_params_per_cam, int num_errs_per_measurement)
        : num_params_per_cam_(num_params_per_cam),
```

```
            num_errs_per_measurement_(num_errs_per_measurement)
    {
        setRefinementMask(Mat::ones(3, 3, CV_8U));
        setConfThresh(1.);
        setTermCriteria(cvTermCriteria(CV_TERMCRIT_EPS + CV_TERMCRIT_ITER, 1000, DBL_EPSILON));
    }

    // Runs bundle adjustment
    virtual void estimate(const std::vector<ImageFeatures> &features,
                          const std::vector<MatchesInfo> &pairwise_matches,
                          std::vector<CameraParams> &cameras);

    virtual void setUpInitialCameraParams(const std::vector<CameraParams> &cameras) = 0;
    virtual void obtainRefinedCameraParams(std::vector<CameraParams> &cameras) const = 0;
    virtual void calcError(Mat &err) = 0;
    virtual void calcJacobian(Mat &jac) = 0;

    // 3x3 8U mask, where 0 means don't refine respective parameter, != 0 means refine
    Mat refinement_mask_;

    int num_images_;
    int total_num_matches_;

    int num_params_per_cam_;
    int num_errs_per_measurement_;

    const ImageFeatures *features_;
    const MatchesInfo *pairwise_matches_;

    // Threshold to filter out poorly matched image pairs
    double conf_thresh_;

    //Levenberg-Marquardt algorithm termination criteria
    CvTermCriteria term_criteria_;

    // Camera parameters matrix (CV_64F)
    Mat cam_params_;

    // Connected images pairs
    std::vector<std::pair<int,int> > edges_;
};
```

**See Also:**

detail::Estimator

## detail::BundleAdjusterBase::BundleAdjusterBase

Construct a bundle adjuster base instance.

**C++:** detail::BundleAdjusterBase::**BundleAdjusterBase**(int **num_params_per_cam**, int **num_errs_per_measurement**)

> **Parameters**
>
> > **num_params_per_cam** – Number of parameters per camera
> >
> > **num_errs_per_measurement** – Number of error terms (components) per match

## detail::BundleAdjusterBase::setUpInitialCameraParams

Sets initial camera parameter to refine.

**C++:** void detail::BundleAdjusterBase::**setUpInitialCameraParams**(const std::vector<CameraParams>& **cameras**)

> Parameters
>
> > **cameras** – Camera parameters

## detail::BundleAdjusterBase::calcError

Calculates error vector.

**C++:** void detail::BundleAdjusterBase::**calcError**(Mat& **err**)

> Parameters
>
> > **err** – Error column-vector of length `total_num_matches` ∗ `num_errs_per_measurement`

## detail::BundleAdjusterBase::calcJacobian

Calculates the cost function jacobian.

**C++:** void detail::BundleAdjusterBase::**calcJacobian**(Mat& **jac**)

> Parameters
>
> > **jac** – Jacobian matrix of dimensions (`total_num_matches` ∗ `num_errs_per_measurement`) x (`num_images` ∗ `num_params_per_cam`)

## detail::BundleAdjusterBase::obtainRefinedCameraParams

Gets the refined camera parameters.

**C++:** void detail::BundleAdjusterBase::**obtainRefinedCameraParams**(std::vector<CameraParams>& **cameras**) const

> Parameters
>
> > **cameras** – Refined camera parameters

## detail::BundleAdjusterReproj

class detail::**BundleAdjusterReproj** : **public** detail::**BundleAdjusterBase**

Implementation of the camera parameters refinement algorithm which minimizes sum of the reprojection error squares.

```
class CV_EXPORTS BundleAdjusterReproj : public BundleAdjusterBase
{
public:
    BundleAdjusterReproj() : BundleAdjusterBase(7, 2) {}

private:
```

```
    /* hidden */
};
```

**See Also:**

detail::BundleAdjusterBase, detail::Estimator

## detail::BundleAdjusterRay

**class** detail::**BundleAdjusterRay** : **public** detail::**BundleAdjusterBase**

Implementation of the camera parameters refinement algorithm which minimizes sum of the distances between the rays passing through the camera center and a feature.

```
class CV_EXPORTS BundleAdjusterRay : public BundleAdjusterBase
{
public:
    BundleAdjusterRay() : BundleAdjusterBase(4, 3) {}

private:
    /* hidden */
};
```

**See Also:**

detail::BundleAdjusterBase

## detail::WaveCorrectKind

Wave correction kind.

```
enum CV_EXPORTS WaveCorrectKind
{
    WAVE_CORRECT_HORIZ,
    WAVE_CORRECT_VERT
};
```

## detail::waveCorrect

Tries to make panorama more horizontal (or vertical).

**C++:** void detail::**waveCorrect**(std::vector<Mat>& **rmats**, WaveCorrectKind **kind**)

>   **Parameters**

>>      **rmats** – Camera rotation matrices.

>>      **kind** – Correction kind, see detail::WaveCorrectKind.

## 13.7 Autocalibration

## detail::focalsFromHomography

Tries to estimate focal lengths from the given homography under the assumption that the camera undergoes rotations around its centre only.

**C++:** void detail::**focalsFromHomography**(const Mat& **H**, double& **f0**, double& **f1**, bool& **f0_ok**, bool& **f1_ok**)

> **Parameters**
>> **H** – Homography.
>>
>> **f0** – Estimated focal length along X axis.
>>
>> **f1** – Estimated focal length along Y axis.
>>
>> **f0_ok** – True, if f0 was estimated successfully, false otherwise.
>>
>> **f1_ok** – True, if f1 was estimated successfully, false otherwise.

## detail::estimateFocal

Estimates focal lengths for each given camera.

**C++:** void detail::**estimateFocal**(const std::vector<ImageFeatures>& **features**, const std::vector<MatchesInfo>& **pairwise_matches**, std::vector<double>& **focals**)

> **Parameters**
>> **features** – Features of images.
>>
>> **pairwise_matches** – Matches between all image pairs.
>>
>> **focals** – Estimated focal lengths for each camera.

# 13.8 Images Warping

## detail::RotationWarper

**class** detail::**RotationWarper**

Rotation-only model image warper interface.

```
class CV_EXPORTS RotationWarper
{
public:
    virtual ~RotationWarper() {}

    virtual Point2f warpPoint(const Point2f &pt, const Mat &K, const Mat &R) = 0;

    virtual Rect buildMaps(Size src_size, const Mat &K, const Mat &R, Mat &xmap, Mat &ymap) = 0;

    virtual Point warp(const Mat &src, const Mat &K, const Mat &R, int interp_mode, int border_mode,
                       Mat &dst) = 0;

    virtual void warpBackward(const Mat &src, const Mat &K, const Mat &R, int interp_mode, int border_mode,
                              Size dst_size, Mat &dst) = 0;

    virtual Rect warpRoi(Size src_size, const Mat &K, const Mat &R) = 0;
};
```

## detail::RotationWarper::warpPoint

Projects the image point.

**C++:** Point2f detail::RotationWarper::**warpPoint**(const Point2f& **pt**, const Mat& **K**, const Mat& **R**)

> **Parameters**
>
> > **pt** – Source point
> >
> > **K** – Camera intrinsic parameters
> >
> > **R** – Camera rotation matrix
>
> **Returns** Projected point

## detail::RotationWarper::buildMaps

Builds the projection maps according to the given camera data.

**C++:** Rect detail::RotationWarper::**buildMaps**(Size **src_size**, const Mat& **K**, const Mat& **R**, Mat& **xmap**, Mat& **ymap**)

> **Parameters**
>
> > **src_size** – Source image size
> >
> > **K** – Camera intrinsic parameters
> >
> > **R** – Camera rotation matrix
> >
> > **xmap** – Projection map for the x axis
> >
> > **ymap** – Projection map for the y axis
>
> **Returns** Projected image minimum bounding box

## detail::RotationWarper::warp

Projects the image.

**C++:** Point detail::RotationWarper::**warp**(const Mat& **src**, const Mat& **K**, const Mat& **R**, int **interp_mode**, int **border_mode**, Mat& **dst**)

> **Parameters**
>
> > **src** – Source image
> >
> > **K** – Camera intrinsic parameters
> >
> > **R** – Camera rotation matrix
> >
> > **interp_mode** – Interpolation mode
> >
> > **border_mode** – Border extrapolation mode
> >
> > **dst** – Projected image
>
> **Returns** Project image top-left corner

## detail::RotationWarper::warpBackward

Projects the image backward.

**C++:** void detail::RotationWarper::**warpBackward**(const Mat& **src**, const Mat& **K**, const Mat& **R**, int **interp_mode**, int **border_mode**, Size **dst_size**, Mat& **dst**)

>    **Parameters**

>        **src** – Projected image

>        **K** – Camera intrinsic parameters

>        **R** – Camera rotation matrix

>        **interp_mode** – Interpolation mode

>        **border_mode** – Border extrapolation mode

>        **dst_size** – Backward-projected image size

>        **dst** – Backward-projected image

## detail::RotationWarper::warpRoi

**C++:** Rect detail::RotationWarper::**warpRoi**(Size **src_size**, const Mat& **K**, const Mat& **R**)

>    **Parameters**

>        **src_size** – Source image bounding box

>        **K** – Camera intrinsic parameters

>        **R** – Camera rotation matrix

>    **Returns** Projected image minimum bounding box

## detail::ProjectorBase

**struct** detail::**ProjectorBase**

Base class for warping logic implementation.

```
struct CV_EXPORTS ProjectorBase
{
    void setCameraParams(const Mat &K = Mat::eye(3, 3, CV_32F),
                         const Mat &R = Mat::eye(3, 3, CV_32F),
                         const Mat &T = Mat::zeros(3, 1, CV_32F));

    float scale;
    float k[9];
    float rinv[9];
    float r_kinv[9];
    float k_rinv[9];
    float t[3];
};
```

## detail::RotationWarperBase

**class** detail::**RotationWarperBase**

Base class for rotation-based warper using a detail::ProjectorBase derived class.

```cpp
template <class P>
class CV_EXPORTS RotationWarperBase : public RotationWarper
{
public:
    Point2f warpPoint(const Point2f &pt, const Mat &K, const Mat &R);

    Rect buildMaps(Size src_size, const Mat &K, const Mat &R, Mat &xmap, Mat &ymap);

    Point warp(const Mat &src, const Mat &K, const Mat &R, int interp_mode, int border_mode,
            Mat &dst);

    void warpBackward(const Mat &src, const Mat &K, const Mat &R, int interp_mode, int border_mode,
                    Size dst_size, Mat &dst);

    Rect warpRoi(Size src_size, const Mat &K, const Mat &R);

protected:

    // Detects ROI of the destination image. It's correct for any projection.
    virtual void detectResultRoi(Size src_size, Point &dst_tl, Point &dst_br);

    // Detects ROI of the destination image by walking over image border.
    // Correctness for any projection isn't guaranteed.
    void detectResultRoiByBorder(Size src_size, Point &dst_tl, Point &dst_br);

    P projector_;
};
```

## detail::PlaneWarper

**class** detail::**PlaneWarper** : **public** detail::**RotationWarperBase<PlaneProjector>**

Warper that maps an image onto the z = 1 plane.

```cpp
class CV_EXPORTS PlaneWarper : public RotationWarperBase<PlaneProjector>
{
public:
    PlaneWarper(float scale = 1.f) { projector_.scale = scale; }

    void setScale(float scale) { projector_.scale = scale; }

    Point2f warpPoint(const Point2f &pt, const Mat &K, const Mat &R, const Mat &T);

    Rect buildMaps(Size src_size, const Mat &K, const Mat &R, const Mat &T, Mat &xmap, Mat &ymap);

    Point warp(const Mat &src, const Mat &K, const Mat &R, const Mat &T, int interp_mode, int border_mode,
            Mat &dst);

    Rect warpRoi(Size src_size, const Mat &K, const Mat &R, const Mat &T);

protected:
```

```
    void detectResultRoi(Size src_size, Point &dst_tl, Point &dst_br);
};
```

**See Also:**

detail::RotationWarper

## detail::PlaneWarper::PlaneWarper

Construct an instance of the plane warper class.

**C++:** void detail::PlaneWarper::**PlaneWarper**(float **scale**=1.f)

> **Parameters**
>
> > **scale** – Projected image scale multiplier

## detail::SphericalWarper

**class** detail::**SphericalWarper** : **public** detail::**RotationWarperBase<SphericalProjector>**

Warper that maps an image onto the unit sphere located at the origin.

```
class CV_EXPORTS SphericalWarper : public RotationWarperBase<SphericalProjector>
{
public:
    SphericalWarper(float scale) { projector_.scale = scale; }

protected:
    void detectResultRoi(Size src_size, Point &dst_tl, Point &dst_br);
};
```

**See Also:**

detail::RotationWarper

## detail::SphericalWarper::SphericalWarper

Construct an instance of the spherical warper class.

**C++:** void detail::SphericalWarper::**SphericalWarper**(float **scale**)

> **Parameters**
>
> > **scale** – Projected image scale multiplier

## detail::CylindricalWarper

**class** detail::**CylindricalWarper** : **public** detail::**RotationWarperBase<CylindricalProjector>**

Warper that maps an image onto the x*x + z*z = 1 cylinder.

```
class CV_EXPORTS CylindricalWarper : public RotationWarperBase<CylindricalProjector>
{
public:
    CylindricalWarper(float scale) { projector_.scale = scale; }
```

```
protected:
    void detectResultRoi(Size src_size, Point &dst_tl, Point &dst_br)
    {
        RotationWarperBase<CylindricalProjector>::detectResultRoiByBorder(src_size, dst_tl, dst_br);
    }
};
```

See Also:

detail::RotationWarper

## detail::CylindricalWarper::CylindricalWarper

Construct an instance of the cylindrical warper class.

**C++:** void detail::CylindricalWarper::**CylindricalWarper**(float **scale**)

>   **Parameters**

>>   **scale** – Projected image scale multiplier

# 13.9 Seam Estimation

## detail::SeamFinder

**class** detail::**SeamFinder**

Base class for a seam estimator.

```
class CV_EXPORTS SeamFinder
{
public:
    virtual ~SeamFinder() {}
    virtual void find(const std::vector<Mat> &src, const std::vector<Point> &corners,
                      std::vector<Mat> &masks) = 0;
};
```

## detail::SeamFinder::find

Estimates seams.

**C++:** void detail::SeamFinder::**find**(const std::vector<Mat>& **src**, const std::vector<Point>& **corners**, std::vector<Mat>& **masks**)

>   **Parameters**

>>   **src** – Source images

>>   **corners** – Source image top-left corners

>>   **masks** – Source image masks to update

## detail::NoSeamFinder

**class** detail::**NoSeamFinder** : **public** detail::**SeamFinder**

Stub seam estimator which does nothing.

```cpp
class CV_EXPORTS NoSeamFinder : public SeamFinder
{
public:
    void find(const std::vector<Mat>&, const std::vector<Point>&, std::vector<Mat>&) {}
};
```

**See Also:**

detail::SeamFinder

## detail::PairwiseSeamFinder

**class** detail::**PairwiseSeamFinder** : **public** detail::**SeamFinder**

Base class for all pairwise seam estimators.

```cpp
class CV_EXPORTS PairwiseSeamFinder : public SeamFinder
{
public:
    virtual void find(const std::vector<Mat> &src, const std::vector<Point> &corners,
                      std::vector<Mat> &masks);

protected:
    void run();
    virtual void findInPair(size_t first, size_t second, Rect roi) = 0;

    std::vector<Mat> images_;
    std::vector<Size> sizes_;
    std::vector<Point> corners_;
    std::vector<Mat> masks_;
};
```

**See Also:**

detail::SeamFinder

## detail::PairwiseSeamFinder::findInPair

Resolves masks intersection of two specified images in the given ROI.

**C++:** void detail::PairwiseSeamFinder::**findInPair**(size_t **first**, size_t **second**, Rect **roi**)

> **Parameters**
>
> > **first** – First image index
> >
> > **second** – Second image index
> >
> > **roi** – Region of interest

## detail::VoronoiSeamFinder

**class** detail::**VoronoiSeamFinder** : **public** detail::**PairwiseSeamFinder**

Voronoi diagram-based seam estimator.

```
class CV_EXPORTS VoronoiSeamFinder : public PairwiseSeamFinder
{
public:
    virtual void find(const std::vector<Size> &size, const std::vector<Point> &corners,
                      std::vector<Mat> &masks);
private:
    void findInPair(size_t first, size_t second, Rect roi);
};
```

**See Also:**

detail::PairwiseSeamFinder

## detail::GraphCutSeamFinderBase

**class** detail::**GraphCutSeamFinderBase**

Base class for all minimum graph-cut-based seam estimators.

```
class CV_EXPORTS GraphCutSeamFinderBase
{
public:
    enum { COST_COLOR, COST_COLOR_GRAD };
};
```

## detail::GraphCutSeamFinder

**class** detail::**GraphCutSeamFinder** : **public** detail::**GraphCutSeamFinderBase**, **public** detail::**SeamFinder**

Minimum graph cut-based seam estimator. See details in [V03].

```
class CV_EXPORTS GraphCutSeamFinder : public GraphCutSeamFinderBase, public SeamFinder
{
public:
    GraphCutSeamFinder(int cost_type = COST_COLOR_GRAD, float terminal_cost = 10000.f,
                       float bad_region_penalty = 1000.f);

    void find(const std::vector<Mat> &src, const std::vector<Point> &corners,
              std::vector<Mat> &masks);

private:
    /* hidden */
};
```

**See Also:**

detail::GraphCutSeamFinderBase, detail::SeamFinder

## 13.10 Exposure Compensation

### detail::ExposureCompensator

**class** detail::**ExposureCompensator**

Base class for all exposure compensators.

```cpp
class CV_EXPORTS ExposureCompensator
{
public:
    virtual ~ExposureCompensator() {}

    enum { NO, GAIN, GAIN_BLOCKS };
    static Ptr<ExposureCompensator> createDefault(int type);

    void feed(const std::vector<Point> &corners, const std::vector<Mat> &images,
            const std::vector<Mat> &masks);
    virtual void feed(const std::vector<Point> &corners, const std::vector<Mat> &images,
                    const std::vector<std::pair<Mat,uchar> > &masks) = 0;
    virtual void apply(int index, Point corner, Mat &image, const Mat &mask) = 0;
};
```

### detail::ExposureCompensator::feed

**C++:** void detail::ExposureCompensator::**feed**(const std::vector<Point>& **corners**, const std::vector<Mat>& **images**, const std::vector<Mat>& **masks**)

**C++:** void detail::ExposureCompensator::**feed**(const std::vector<Point>& **corners**, const std::vector<Mat>& **images**, const std::vector<std::pair<Mat, uchar>>& **masks**)

> **Parameters**
>
>> **corners** – Source image top-left corners
>>
>> **images** – Source images
>>
>> **masks** – Image masks to update (second value in pair specifies the value which should be used to detect where image is)

### detil::ExposureCompensator::apply

Compensate exposure in the specified image.

**C++:** void detail::ExposureCompensator::**apply**(int **index**, Point **corner**, Mat& **image**, const Mat& **mask**)

> **Parameters**
>
>> **index** – Image index
>>
>> **corner** – Image top-left corner
>>
>> **image** – Image to process
>>
>> **mask** – Image mask

## detail::NoExposureCompensator

**class** detail::**NoExposureCompensator** : **public** detail::**ExposureCompensator**

Stub exposure compensator which does nothing.

```
class CV_EXPORTS NoExposureCompensator : public ExposureCompensator
{
public:
    void feed(const std::vector<Point> &/*corners*/, const std::vector<Mat> &/*images*/,
              const std::vector<std::pair<Mat,uchar> > &/*masks*/) {};
    void apply(int /*index*/, Point /*corner*/, Mat &/*image*/, const Mat &/*mask*/) {};
};
```

**See Also:**

detail::ExposureCompensator

## detail::GainCompensator

**class** detail::**GainCompensator** : **public** detail::**ExposureCompensator**

Exposure compensator which tries to remove exposure related artifacts by adjusting image intensities, see [BL07] and [WJ10] for details.

```
class CV_EXPORTS GainCompensator : public ExposureCompensator
{
public:
    void feed(const std::vector<Point> &corners, const std::vector<Mat> &images,
              const std::vector<std::pair<Mat,uchar> > &masks);
    void apply(int index, Point corner, Mat &image, const Mat &mask);
    std::vector<double> gains() const;

private:
    /* hidden */
};
```

**See Also:**

detail::ExposureCompensator

## detail::BlocksGainCompensator

**class** detail::**BlocksGainCompensator** : **public** detail::**ExposureCompensator**

Exposure compensator which tries to remove exposure related artifacts by adjusting image block intensities, see [UES01] for details.

```
class CV_EXPORTS BlocksGainCompensator : public ExposureCompensator
{
public:
    BlocksGainCompensator(int bl_width = 32, int bl_height = 32)
            : bl_width_(bl_width), bl_height_(bl_height) {}
    void feed(const std::vector<Point> &corners, const std::vector<Mat> &images,
              const std::vector<std::pair<Mat,uchar> > &masks);
    void apply(int index, Point corner, Mat &image, const Mat &mask);

private:
```

```
    /* hidden */
};
```

**See Also:**

detail::ExposureCompensator

# 13.11 Image Blenders

## detail::Blender

**class** detail::**Blender**

Base class for all blenders.

```
class CV_EXPORTS Blender
{
public:
    virtual ~Blender() {}

    enum { NO, FEATHER, MULTI_BAND };
    static Ptr<Blender> createDefault(int type, bool try_gpu = false);

    void prepare(const std::vector<Point> &corners, const std::vector<Size> &sizes);
    virtual void prepare(Rect dst_roi);
    virtual void feed(const Mat &img, const Mat &mask, Point tl);
    virtual void blend(Mat &dst, Mat &dst_mask);

protected:
    Mat dst_, dst_mask_;
    Rect dst_roi_;
};
```

## detail::Blender::prepare

Prepares the blender for blending.

**C++:** void detail::Blender::**prepare**(const std::vector<Point>& **corners**, const std::vector<Size>& **sizes**)

> **Parameters**
>
> > **corners** – Source images top-left corners
> >
> > **sizes** – Source image sizes

## detail::Blender::feed

Processes the image.

**C++:** void detail::Blender::**feed**(const Mat& **img**, const Mat& **mask**, Point **tl**)

> **Parameters**
>
> > **img** – Source image
> >
> > **mask** – Source image mask

> > **tl** – Source image top-left corners

## detail::Blender::blend

Blends and returns the final pano.

**C++:** void detail::Blender::**blend**(Mat& **dst**, Mat& **dst_mask**)

> > **Parameters**

> > > **dst** – Final pano

> > > **dst_mask** – Final pano mask

## detail::FeatherBlender

class detail::**FeatherBlender** : **public** detail::**Blender**

Simple blender which mixes images at its borders.

```cpp
class CV_EXPORTS FeatherBlender : public Blender
{
public:
    FeatherBlender(float sharpness = 0.02f) { setSharpness(sharpness); }

    float sharpness() const { return sharpness_; }
    void setSharpness(float val) { sharpness_ = val; }

    void prepare(Rect dst_roi);
    void feed(const Mat &img, const Mat &mask, Point tl);
    void blend(Mat &dst, Mat &dst_mask);

    // Creates weight maps for fixed set of source images by their masks and top-left corners.
    // Final image can be obtained by simple weighting of the source images.
    Rect createWeightMaps(const std::vector<Mat> &masks, const std::vector<Point> &corners,
                          std::vector<Mat> &weight_maps);

private:
    /* hidden */
};
```

**See Also:**

detail::Blender

## detail::MultiBandBlender

class detail::**MultiBandBlender** : **public** detail::**Blender**

Blender which uses multi-band blending algorithm (see [BA83]).

```cpp
class CV_EXPORTS MultiBandBlender : public Blender
{
public:
    MultiBandBlender(int try_gpu = false, int num_bands = 5);
    int numBands() const { return actual_num_bands_; }
    void setNumBands(int val) { actual_num_bands_ = val; }
```

```
    void prepare(Rect dst_roi);
    void feed(const Mat &img, const Mat &mask, Point tl);
    void blend(Mat &dst, Mat &dst_mask);

private:
    /* hidden */
};
```

**See Also:**

detail::Blender

# NONFREE. NON-FREE FUNCTIONALITY

The module contains algorithms that may be patented in some countries or have some other limitations on the use.

## 14.1 Feature Detection and Description

### SIFT

**class SIFT** : **public Feature2D**

Class for extracting keypoints and computing descriptors using the Scale Invariant Feature Transform (SIFT) algorithm by D. Lowe [Lowe04].

### SIFT::SIFT

The SIFT constructors.

**C++:** SIFT::**SIFT**(int **nfeatures**=0, int **nOctaveLayers**=3, double **contrastThreshold**=0.04, double **edgeThreshold**=10, double **sigma**=1.6)

> **Parameters**

>> **nfeatures** – The number of best features to retain. The features are ranked by their scores (measured in SIFT algorithm as the local contrast)

>> **nOctaveLayers** – The number of layers in each octave. 3 is the value used in D. Lowe paper. The number of octaves is computed automatically from the image resolution.

>> **contrastThreshold** – The contrast threshold used to filter out weak features in semi-uniform (low-contrast) regions. The larger the threshold, the less features are produced by the detector.

>> **edgeThreshold** – The threshold used to filter out edge-like features. Note that the its meaning is different from the contrastThreshold, i.e. the larger the edgeThreshold, the less features are filtered out (more features are retained).

>> **sigma** – The sigma of the Gaussian applied to the input image at the octave #0. If your image is captured with a weak camera with soft lenses, you might want to reduce the number.

## SIFT::operator ()

Extract features and computes their descriptors using SIFT algorithm

**C++:** void SIFT::**operator()**(InputArray **img**, InputArray **mask**, vector<KeyPoint>& **keypoints**, OutputArray **descriptors**, bool **useProvidedKeypoints**=false)

> **Parameters**
>
>> **img** – Input 8-bit grayscale image
>>
>> **mask** – Optional input mask that marks the regions where we should detect features.
>>
>> **keypoints** – The input/output vector of keypoints
>>
>> **descriptors** – The output matrix of descriptors. Pass `cv::noArray()` if you do not need them.
>>
>> **useProvidedKeypoints** – Boolean flag. If it is true, the keypoint detector is not run. Instead, the provided vector of keypoints is used and the algorithm just computes their descriptors.

## SURF

**class SURF** : **public Feature2D**
> Class for extracting Speeded Up Robust Features from an image [Bay06]. The class is derived from `CvSURFParams` structure, which specifies the algorithm parameters:

int **extended**

> • 0 means that the basic descriptors (64 elements each) shall be computed
>
> • 1 means that the extended descriptors (128 elements each) shall be computed

int **upright**

> • 0 means that detector computes orientation of each feature.
>
> • 1 means that the orientation is not computed (which is much, much faster). For example, if you match images from a stereo pair, or do image stitching, the matched features likely have very similar angles, and you can speed up feature extraction by setting `upright=1`.

double **hessianThreshold**
> Threshold for the keypoint detector. Only features, whose hessian is larger than `hessianThreshold` are retained by the detector. Therefore, the larger the value, the less keypoints you will get. A good default value could be from 300 to 500, depending from the image contrast.

int **nOctaves**
> The number of a gaussian pyramid octaves that the detector uses. It is set to 4 by default. If you want to get very large features, use the larger value. If you want just small features, decrease it.

int **nOctaveLayers**
> The number of images within each octave of a gaussian pyramid. It is set to 2 by default.

## SURF::SURF

The SURF extractor constructors.

**C++:** SURF::**SURF**()

**C++:** SURF::**SURF**(double **hessianThreshold**, int **nOctaves**=4, int **nOctaveLayers**=2, bool **extended**=true, bool **upright**=false )

**Python:** cv2.**SURF**([hessianThreshold[, nOctaves[, nOctaveLayers[, extended[, upright ]]]]]) → <SURF object>

>> **Parameters**

>>> **hessianThreshold** – Threshold for hessian keypoint detector used in SURF.

>>> **nOctaves** – Number of pyramid octaves the keypoint detector will use.

>>> **nOctaveLayers** – Number of octave layers within each octave.

>>> **extended** – Extended descriptor flag (true - use extended 128-element descriptors; false - use 64-element descriptors).

>>> **upright** – Up-right or rotated features flag (true - do not compute orientation of features; false - compute orientation).

## SURF::operator()

Detects keypoints and computes SURF descriptors for them.

**C++:** void SURF::**operator()**(InputArray **img**, InputArray **mask**, vector<KeyPoint>& **keypoints**) const

**C++:** void SURF::**operator()**(InputArray **img**, InputArray **mask**, vector<KeyPoint>& **keypoints**, OutputArray **descriptors**, bool **useProvidedKeypoints**=false)

**Python:** cv2.SURF.**detect**(image[, mask ]) → keypoints

**C:** void **cvExtractSURF**(const CvArr* **image**, const CvArr* **mask**, CvSeq** **keypoints**, CvSeq** **descriptors**, CvMemStorage* **storage**, CvSURFParams **params**)

**Python:** cv.**ExtractSURF**(image, mask, storage, params)-> (keypoints, descriptors)

>> **Parameters**

>>> **image** – Input 8-bit grayscale image

>>> **mask** – Optional input mask that marks the regions where we should detect features.

>>> **keypoints** – The input/output vector of keypoints

>>> **descriptors** – The output matrix of descriptors. Pass cv::noArray() if you do not need them.

>>> **useProvidedKeypoints** – Boolean flag. If it is true, the keypoint detector is not run. Instead, the provided vector of keypoints is used and the algorithm just computes their descriptors.

>>> **storage** – Memory storage for the output keypoints and descriptors in OpenCV 1.x API.

>>> **params** – SURF algorithm parameters in OpenCV 1.x API.

The function is parallelized with the TBB library.

If you are using the C version, make sure you call cv::initModule_nonfree() from nonfree/nonfree.hpp.

# CONTRIB. CONTRIBUTED/EXPERIMENTAL STUFF

The module contains some recently added functionality that has not been stabilized, or functionality that is considered optional.

## 15.1 Stereo Correspondence

### StereoVar

**class StereoVar**

Class for computing stereo correspondence using the variational matching algorithm

```cpp
class StereoVar
{
    StereoVar();
    StereoVar(    int levels, double pyrScale,
                            int nIt, int minDisp, int maxDisp,
                            int poly_n, double poly_sigma, float fi,
                            float lambda, int penalization, int cycle,
                            int flags);
    virtual ~StereoVar();

    virtual void operator()(InputArray left, InputArray right, OutputArray disp);

    int       levels;
    double    pyrScale;
    int       nIt;
    int       minDisp;
    int       maxDisp;
    int       poly_n;
    double    poly_sigma;
    float     fi;
    float     lambda;
    int       penalization;
    int       cycle;
    int       flags;

    ...
};
```

The class implements the modified S. G. Kosov algorithm [Publication] that differs from the original one as follows:

- The automatic initialization of method's parameters is added.

- The method of Smart Iteration Distribution (SID) is implemented.

- The support of Multi-Level Adaptation Technique (MLAT) is not included.

- The method of dynamic adaptation of method's parameters is not included.

## StereoVar::StereoVar

**C++:** StereoVar::**StereoVar**()

**C++:** StereoVar::**StereoVar**(int **levels**, double **pyrScale**, int **nIt**, int **minDisp**, int **maxDisp**, int **poly_n**, double **poly_sigma**, float **fi**, float **lambda**, int **penalization**, int **cycle**, int **flags**)

The constructor

**Parameters**

**levels** – The number of pyramid layers, including the initial image. levels=1 means that no extra layers are created and only the original images are used. This parameter is ignored if flag USE_AUTO_PARAMS is set.

**pyrScale** – Specifies the image scale (<1) to build the pyramids for each image. pyrScale=0.5 means the classical pyramid, where each next layer is twice smaller than the previous. (This parameter is ignored if flag USE_AUTO_PARAMS is set).

**nIt** – The number of iterations the algorithm does at each pyramid level. (If the flag USE_SMART_ID is set, the number of iterations will be redistributed in such a way, that more iterations will be done on more coarser levels.)

**minDisp** – Minimum possible disparity value. Could be negative in case the left and right input images change places.

**maxDisp** – Maximum possible disparity value.

**poly_n** – Size of the pixel neighbourhood used to find polynomial expansion in each pixel. The larger values mean that the image will be approximated with smoother surfaces, yielding more robust algorithm and more blurred motion field. Typically, poly_n = 3, 5 or 7

**poly_sigma** – Standard deviation of the Gaussian that is used to smooth derivatives that are used as a basis for the polynomial expansion. For poly_n=5 you can set poly_sigma=1.1 , for poly_n=7 a good value would be poly_sigma=1.5

**fi** – The smoothness parameter, ot the weight coefficient for the smoothness term.

**lambda** – The threshold parameter for edge-preserving smoothness. (This parameter is ignored if PENALIZATION_CHARBONNIER or PENALIZATION_PERONA_MALIK is used.)

**penalization** – Possible values: PENALIZATION_TICHONOV - linear smoothness; PENALIZATION_CHARBONNIER - non-linear edge preserving smoothness; PENALIZATION_PERONA_MALIK - non-linear edge-enhancing smoothness. (This parameter is ignored if flag USE_AUTO_PARAMS is set).

**cycle** – Type of the multigrid cycle. Possible values: CYCLE_O and CYCLE_V for null- and v-cycles respectively. (This parameter is ignored if flag USE_AUTO_PARAMS is set).

**flags** – The operation flags; can be a combination of the following:

– USE_INITIAL_DISPARITY: Use the input flow as the initial flow approximation.

- **USE_EQUALIZE_HIST**: Use the histogram equalization in the pre-processing phase.

- **USE_SMART_ID**: Use the smart iteration distribution (SID).

- **USE_AUTO_PARAMS**: Allow the method to initialize the main parameters.

- **USE_MEDIAN_FILTERING**: Use the median filer of the solution in the post processing phase.

The first constructor initializes `StereoVar` with all the default parameters. So, you only have to set `StereoVar::maxDisp` and / or `StereoVar::minDisp` at minimum. The second constructor enables you to set each parameter to a custom value.

## StereoVar::operator ()

**C++:** void StereoVar::**operator()**(const Mat& **left**, const Mat& **right**, Mat& **disp**)
> Computes disparity using the variational algorithm for a rectified stereo pair.

> **Parameters**

>> **left** – Left 8-bit single-channel or 3-channel image.

>> **right** – Right image of the same size and the same type as the left one.

>> **disp** – Output disparity map. It is a 8-bit signed single-channel image of the same size as the input image.

The method executes the variational algorithm on a rectified stereo pair. See `stereo_match.cpp` OpenCV sample on how to prepare images and call the method.

**Note**:

The method is not constant, so you should not use the same `StereoVar` instance from different threads simultaneously.

# 15.2 FaceRecognizer - Face Recognition with OpenCV

OpenCV 2.4 now comes with the very new `FaceRecognizer` class for face recognition. This documentation is going to explain you *the API* in detail and it will give you a lot of help to get started (full source code examples). *Face Recognition with OpenCV* is the definite guide to the new `FaceRecognizer`. There's also a *tutorial on gender classification*, a *tutorial for face recognition in videos* and it's shown *how to load & save your results*.

These documents are the help I have wished for, when I was working myself into face recognition. I hope you also think the new `FaceRecognizer` is a useful addition to OpenCV.

Please issue any feature requests and/or bugs on the official OpenCV bug tracker at:

- http://code.opencv.org/projects/opencv/issues

## Contents

**FaceRecognizer**

**FaceRecognizer**

class **FaceRecognizer** : **public Algorithm**

All face recognition models in OpenCV are derived from the abstract base class `FaceRecognizer`, which provides a unified access to all face recongition algorithms in OpenCV.

```cpp
class FaceRecognizer : public Algorithm
{
public:
    //! virtual destructor
    virtual ~FaceRecognizer() {}

    // Trains a FaceRecognizer.
    virtual void train(InputArray src, InputArray labels) = 0;

    // Updates a FaceRecognizer.
    virtual void update(InputArrayOfArrays src, InputArray labels);

    // Gets a prediction from a FaceRecognizer.
    virtual int predict(InputArray src) const = 0;

    // Predicts the label and confidence for a given sample.
    virtual void predict(InputArray src, int &label, double &confidence) const = 0;

    // Serializes this object to a given filename.
    virtual void save(const string& filename) const;

    // Deserializes this object from a given filename.
    virtual void load(const string& filename);

    // Serializes this object to a given cv::FileStorage.
    virtual void save(FileStorage& fs) const = 0;

    // Deserializes this object from a given cv::FileStorage.
    virtual void load(const FileStorage& fs) = 0;
};
```

**Description**  I'll go a bit more into detail explaining `FaceRecognizer`, because it doesn't look like a powerful interface at first sight. But: Every `FaceRecognizer` is an `Algorithm`, so you can easily get/set all model internals (if allowed by the implementation). `Algorithm` is a relatively new OpenCV concept, which is available since the 2.4 release. I suggest you take a look at its description.

`Algorithm` provides the following features for all derived classes:

- So called "virtual constructor". That is, each Algorithm derivative is registered at program start and you can get the list of registered algorithms and create instance of a particular algorithm by its name (see `Algorithm::create()`). If you plan to add your own algorithms, it is good practice to add a unique prefix to your algorithms to distinguish them from other algorithms.

- Setting/Retrieving algorithm parameters by name. If you used video capturing functionality from OpenCV highgui module, you are probably familar with `cvSetCaptureProperty()`, `cvGetCaptureProperty()`, `VideoCapture::set()` and `VideoCapture::get()`. `Algorithm` provides similar method where instead of integer id's you specify the parameter names as text strings. See `Algorithm::set()` and `Algorithm::get()` for details.

- Reading and writing parameters from/to XML or YAML files. Every Algorithm derivative can store all its parameters and then read them back. There is no need to re-implement it each time.

Moreover every `FaceRecognizer` supports the:

- **Training** of a `FaceRecognizer` with `FaceRecognizer::train()` on a given set of images (your face database!).

- **Prediction** of a given sample image, that means a face. The image is given as a `Mat`.

- **Loading/Saving** the model state from/to a given XML or YAML.

**Setting the Thresholds**    Sometimes you run into the situation, when you want to apply a threshold on the prediction. A common scenario in face recognition is to tell, wether a face belongs to the training dataset or if it is unknown. You might wonder, why there's no public API in `FaceRecognizer` to set the threshold for the prediction, but rest assured: It's supported.  It just means there's no generic way in an abstract class to provide an interface for setting/getting the thresholds of *every possible* `FaceRecognizer` algorithm.  The appropriate place to set the thresholds is in the constructor of the specific `FaceRecognizer` and since every `FaceRecognizer` is a `Algorithm` (see above), you can get/set the thresholds at runtime!

Here is an example of setting a threshold for the Eigenfaces method, when creating the model:

```
// Let's say we want to keep 10 Eigenfaces and have a threshold value of 10.0
int num_components = 10;
double threshold = 10.0;
// Then if you want to have a cv::FaceRecognizer with a confidence threshold,
// create the concrete implementation with the appropiate parameters:
Ptr<FaceRecognizer> model = createEigenFaceRecognizer(num_components, threshold);
```

Sometimes it's impossible to train the model, just to experiment with threshold values. Thanks to `Algorithm` it's possible to set internal model thresholds during runtime. Let's see how we would set/get the prediction for the Eigenface model, we've created above:

```
// The following line reads the threshold from the Eigenfaces model:
double current_threshold = model->getDouble("threshold");
// And this line sets the threshold to 0.0:
model->set("threshold", 0.0);
```

If you've set the threshold to `0.0` as we did above, then:

```
//
Mat img = imread("person1/3.jpg", CV_LOAD_IMAGE_GRAYSCALE);
// Get a prediction from the model. Note: We've set a threshold of 0.0 above,
// since the distance is almost always larger than 0.0, you'll get -1 as
// label, which indicates, this face is unknown
int predicted_label = model->predict(img);
// ...
```

is going to yield `-1` as predicted label, which states this face is unknown.

**Getting the name of a FaceRecognizer**    Since every `FaceRecognizer` is a `Algorithm`, you can use `Algorithm::name()` to get the name of a `FaceRecognizer`:

```
// Create a FaceRecognizer:
Ptr<FaceRecognizer> model = createEigenFaceRecognizer();
// And here's how to get its name:
std::string name = model->name();
```

### FaceRecognizer::train

Trains a FaceRecognizer with given data and associated labels.

**C++:** void FaceRecognizer::**train**(InputArrayOfArrays **src**, InputArray **labels** = 0)

        **Parameters**

> **src** – The training images, that means the faces you want to learn. The data has to be given as a `vector<Mat>`.
>
> **labels** – The labels corresponding to the images have to be given either as a `vector<int>` or a

The following source code snippet shows you how to learn a Fisherfaces model on a given set of images. The images are read with `imread()` and pushed into a `std::vector<Mat>`. The labels of each image are stored within a `std::vector<int>` (you could also use a `Mat` of type *CV_32SC1*). Think of the label as the subject (the person) this image belongs to, so same subjects (persons) should have the same label. For the available `FaceRecognizer` you don't have to pay any attention to the order of the labels, just make sure same persons have the same label:

```
// holds images and labels
vector<Mat> images;
vector<int> labels;
// images for first person
images.push_back(imread("person0/0.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(0);
images.push_back(imread("person0/1.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(0);
images.push_back(imread("person0/2.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(0);
// images for second person
images.push_back(imread("person1/0.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(1);
images.push_back(imread("person1/1.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(1);
images.push_back(imread("person1/2.jpg", CV_LOAD_IMAGE_GRAYSCALE)); labels.push_back(1);
```

Now that you have read some images, we can create a new `FaceRecognizer`. In this example I'll create a Fisherfaces model and decide to keep all of the possible Fisherfaces:

```
// Create a new Fisherfaces model and retain all available Fisherfaces,
// this is the most common usage of this specific FaceRecognizer:
//
Ptr<FaceRecognizer> model =  createFisherFaceRecognizer();
```

And finally train it on the given dataset (the face images and labels):

```
// This is the common interface to train all of the available cv::FaceRecognizer
// implementations:
//
model->train(images, labels);
```

### FaceRecognizer::update

Updates a FaceRecognizer with given data and associated labels.

**C++:** void FaceRecognizer::**update**(InputArrayOfArrays **src**, InputArray **labels**)

> **Parameters**
>
> > **src** – The training images, that means the faces you want to learn. The data has to be given as a `vector<Mat>`.
> >
> > **labels** – The labels corresponding to the images have to be given either as a `vector<int>` or a

This method updates a (probably trained) `FaceRecognizer`, but only if the algorithm supports it. The Local Binary Patterns Histograms (LBPH) recognizer (see `createLBPHFaceRecognizer()`) can be updated. For the Eigenfaces and Fisherfaces method, this is algorithmically not possible and you have to re-estimate the model with `FaceRecognizer::train()`. In any case, a call to train empties the existing model and learns a new model, while update does not delete any model data.

```
// Create a new LBPH model (it can be updated) and use the default parameters,
// this is the most common usage of this specific FaceRecognizer:
//
Ptr<FaceRecognizer> model =  createLBPHFaceRecognizer();
// This is the common interface to train all of the available cv::FaceRecognizer
// implementations:
//
model->train(images, labels);
// Some containers to hold new image:
vector<Mat> newImages;
vector<int> newLabels;
// You should add some images to the containers:
//
// ...
//
// Now updating the model is as easy as calling:
model->update(newImages,newLabels);
// This will preserve the old model data and extend the existing model
// with the new features extracted from newImages!
```

Calling update on an Eigenfaces model (see `createEigenFaceRecognizer()`), which doesn't support updating, will throw an error similar to:

```
OpenCV Error: The function/feature is not implemented (This FaceRecognizer (FaceRecognizer.Eigenfaces) does not suppor
terminate called after throwing an instance of 'cv::Exception'
```

Please note: The `FaceRecognizer` does not store your training images, because this would be very memory intense and it's not the responsibility of te `FaceRecognizer` to do so. The caller is responsible for maintaining the dataset, he want to work with.

### FaceRecognizer::predict

**C++:** int FaceRecognizer::**predict**(InputArray **src** const = 0)

**C++:** void FaceRecognizer::**predict**(InputArray **src**, int& **label**, double& **confidence** const = 0)
Predicts a label and associated confidence (e.g. distance) for a given input image.

> **Parameters**
>
> > **src** – Sample image to get a prediction from.
> >
> > **label** – The predicted label for the given image.
> >
> > **confidence** – Associated confidence (e.g. distance) for the predicted label.

The suffix `const` means that prediction does not affect the internal model state, so the method can be safely called from within different threads.

The following example shows how to get a prediction from a trained model:

```
using namespace cv;
// Do your initialization here (create the cv::FaceRecognizer model) ...
// ...
// Read in a sample image:
Mat img = imread("person1/3.jpg", CV_LOAD_IMAGE_GRAYSCALE);
// And get a prediction from the cv::FaceRecognizer:
int predicted = model->predict(img);
```

Or to get a prediction and the associated confidence (e.g. distance):

```
using namespace cv;
// Do your initialization here (create the cv::FaceRecognizer model) ...
// ...
Mat img = imread("person1/3.jpg", CV_LOAD_IMAGE_GRAYSCALE);
// Some variables for the predicted label and associated confidence (e.g. distance):
int predicted_label = -1;
double predicted_confidence = 0.0;
// Get the prediction and associated confidence from the model
model->predict(img, predicted_label, predicted_confidence);
```

### FaceRecognizer::save

Saves a FaceRecognizer and its model state.

**C++:** void FaceRecognizer::**save**(const string& **filename**) const
    Saves this model to a given filename, either as XML or YAML.

> **Parameters**
>
> > **filename** – The filename to store this FaceRecognizer to (either XML/YAML).

**C++:** void FaceRecognizer::**save**(FileStorage& **fs**) const
    Saves this model to a given FileStorage.

> **Parameters**
>
> > **fs** – The FileStorage to store this FaceRecognizer to.

Every FaceRecognizer overwrites FaceRecognizer::save(FileStorage& fs) to save the internal model state.
FaceRecognizer::save(const string& filename) saves the state of a model to the given filename.

The suffix const means that prediction does not affect the internal model state, so the method can be safely called
from within different threads.

### FaceRecognizer::load

Loads a FaceRecognizer and its model state.

**C++:** void FaceRecognizer::**load**(const string& **filename**)

**C++:** void FaceRecognizer::**load**(const FileStorage& **fs** = 0)

Loads a persisted model and state from a given XML or YAML file . Every FaceRecognizer
has to overwrite FaceRecognizer::load(FileStorage& fs) to enable loading the model state.
FaceRecognizer::load(FileStorage& fs) in turn gets called by FaceRecognizer::load(const string&
filename), to ease saving a model.

### createEigenFaceRecognizer

**C++:** Ptr<FaceRecognizer> **createEigenFaceRecognizer**(int **num_components**=0, double **thresh-
                                                                  old**=DBL_MAX)

> **Parameters**
>
> > **num_components** – The number of components (read: Eigenfaces) kept for this Prinicpal
> > Component Analysis. As a hint: There's no rule how many components (read: Eigenfaces)
> > should be kept for good reconstruction capabilities. It is based on your input data, so exper-
> > iment with the number. Keeping 80 components should almost always be sufficient.

> **threshold** – The threshold applied in the prediciton.

**Notes:**

- Training and prediction must be done on grayscale images, use `cvtColor()` to convert between the color spaces.

- **THE EIGENFACES METHOD MAKES THE ASSUMPTION, THAT THE TRAINING AND TEST IM-AGES ARE OF EQUAL SIZE.** (caps-lock, because I got so many mails asking for this). You have to make sure your input data has the correct shape, else a meaningful exception is thrown. Use `resize()` to resize the images.

- This model does not support updating.

**Model internal data:**

- `num_components` see `createEigenFaceRecognizer()`.

- `threshold` see `createEigenFaceRecognizer()`.

- `eigenvalues` The eigenvalues for this Principal Component Analysis (ordered descending).

- `eigenvectors` The eigenvectors for this Principal Component Analysis (ordered by their eigenvalue).

- `mean` The sample mean calculated from the training data.

- `projections` The projections of the training data.

- `labels` The threshold applied in the prediction. If the distance to the nearest neighbor is larger than the threshold, this method returns -1.

### createFisherFaceRecognizer

**C++:** Ptr<FaceRecognizer> **createFisherFaceRecognizer**(int **num_components**=0, double **threshold**=DBL_MAX)

> **Parameters**
>
> > **num_components** – The number of components (read: Fisherfaces) kept for this Linear Discriminant Analysis with the Fisherfaces criterion. It's useful to keep all components, that means the number of your classes `c` (read: subjects, persons you want to recognize). If you leave this at the default (`0`) or set it to a value less-equal `0` or greater (`c-1`), it will be set to the correct number (`c-1`) automatically.
> >
> > **threshold** – The threshold applied in the prediction. If the distance to the nearest neighbor is larger than the threshold, this method returns -1.

**Notes:**

- Training and prediction must be done on grayscale images, use `cvtColor()` to convert between the color spaces.

- **THE FISHERFACES METHOD MAKES THE ASSUMPTION, THAT THE TRAINING AND TEST IMAGES ARE OF EQUAL SIZE.** (caps-lock, because I got so many mails asking for this). You have to make sure your input data has the correct shape, else a meaningful exception is thrown. Use `resize()` to resize the images.

- This model does not support updating.

**Model internal data:**

- `num_components` see `createFisherFaceRecognizer()`.

- `threshold` see `createFisherFaceRecognizer()`.

- `eigenvalues` The eigenvalues for this Linear Discriminant Analysis (ordered descending).

- `eigenvectors` The eigenvectors for this Linear Discriminant Analysis (ordered by their eigenvalue).

- `mean` The sample mean calculated from the training data.

- `projections` The projections of the training data.

- `labels` The labels corresponding to the projections.

## createLBPHFaceRecognizer

**C++:** Ptr<FaceRecognizer> **createLBPHFaceRecognizer**(int **radius**=1, int **neighbors**=8, int **grid_x**=8, int **grid_y**=8, double **threshold**=DBL_MAX)

> **Parameters**
>
>> **radius** – The radius used for building the Circular Local Binary Pattern. The greater the radius, the
>>
>> **neighbors** – The number of sample points to build a Circular Local Binary Pattern from. An appropriate value is to use `` 8`` sample points. Keep in mind: the more sample points you include, the higher the computational cost.
>>
>> **grid_x** – The number of cells in the horizontal direction, 8 is a common value used in publications. The more cells, the finer the grid, the higher the dimensionality of the resulting feature vector.
>>
>> **grid_y** – The number of cells in the vertical direction, 8 is a common value used in publications. The more cells, the finer the grid, the higher the dimensionality of the resulting feature vector.
>>
>> **threshold** – The threshold applied in the prediction. If the distance to the nearest neighbor is larger than the threshold, this method returns -1.

**Notes:**

- The Circular Local Binary Patterns (used in training and prediction) expect the data given as grayscale images, use `cvtColor()` to convert between the color spaces.

- This model supports updating.

**Model internal data:**

- `radius` see `createLBPHFaceRecognizer()`.

- `neighbors` see `createLBPHFaceRecognizer()`.

- `grid_x` see `createLBPHFaceRecognizer()`.

- `grid_y` see `createLBPHFaceRecognizer()`.

- `threshold` see `createLBPHFaceRecognizer()`.

- `histograms` Local Binary Patterns Histograms calculated from the given training data (empty if none was given).

• `labels` Labels corresponding to the calculated Local Binary Patterns Histograms.

## Face Recognition with OpenCV

### Introduction

OpenCV (Open Source Computer Vision) is a popular computer vision library started by Intel in 1999. The cross-platform library sets its focus on real-time image processing and includes patent-free implementations of the latest computer vision algorithms. In 2008 Willow Garage took over support and OpenCV 2.3.1 now comes with a programming interface to C, C++, Python and Android. OpenCV is released under a BSD license so it is used in academic projects and commercial products alike.

OpenCV 2.4 now comes with the very new `FaceRecognizer` class for face recognition, so you can start experimenting with face recognition right away. This document is the guide I've wished for, when I was working myself into face recognition. It shows you how to perform face recognition with `FaceRecognizer` in OpenCV (with full source code listings) and gives you an introduction into the algorithms behind. I'll also show how to create the visualizations you can find in many publications, because a lot of people asked for.

The currently available algorithms are:

• Eigenfaces (see `createEigenFaceRecognizer()`)

• Fisherfaces (see `createFisherFaceRecognizer()`)

• Local Binary Patterns Histograms (see `createLBPHFaceRecognizer()`)

You don't need to copy and paste the source code examples from this page, because they are available in the `src` folder coming with this documentation. If you have built OpenCV with the samples turned on, chances are good you have them compiled already! Although it might be interesting for very advanced users, I've decided to leave the implementation details out as I am afraid they confuse new users.

All code in this document is released under the BSD license, so feel free to use it for your projects.

### Face Recognition

Face recognition is an easy task for humans. Experiments in [Tu06] have shown, that even one to three day old babies are able to distinguish between known faces. So how hard could it be for a computer? It turns out we know little about human recognition to date. Are inner features (eyes, nose, mouth) or outer features (head shape, hairline) used for a successful face recognition? How do we analyze an image and how does the brain encode it? It was shown by David Hubel and Torsten Wiesel, that our brain has specialized nerve cells responding to specific local features of a scene, such as lines, edges, angles or movement. Since we don't see the world as scattered pieces, our visual cortex must somehow combine the different sources of information into useful patterns. Automatic face recognition is all about extracting those meaningful features from an image, putting them into a useful representation and performing some kind of classification on them.

Face recognition based on the geometric features of a face is probably the most intuitive approach to face recognition. One of the first automated face recognition systems was described in [Kanade73]: marker points (position of eyes, ears, nose, ...) were used to build a feature vector (distance between the points, angle between them, ...). The recognition was performed by calculating the euclidean distance between feature vectors of a probe and reference image. Such a method is robust against changes in illumination by its nature, but has a huge drawback: the accurate registration of the marker points is complicated, even with state of the art algorithms. Some of the latest work on geometric face recognition was carried out in [Bru92]. A 22-dimensional feature vector was used and experiments on large datasets have shown, that geometrical features alone my not carry enough information for face recognition.

The Eigenfaces method described in [TP91] took a holistic approach to face recognition: A facial image is a point from a high-dimensional image space and a lower-dimensional representation is found, where classification becomes easy. The lower-dimensional subspace is found with Principal Component Analysis, which identifies the axes with maximum variance. While this kind of transformation is optimal from a reconstruction standpoint, it doesn't take any class labels into account. Imagine a situation where the variance is generated from external sources, let it be light. The axes with maximum variance do not necessarily contain any discriminative information at all, hence a classification becomes impossible. So a class-specific projection with a Linear Discriminant Analysis was applied to face recognition in [BHK97]. The basic idea is to minimize the variance within a class, while maximizing the variance between the classes at the same time.

Recently various methods for a local feature extraction emerged. To avoid the high-dimensionality of the input data only local regions of an image are described, the extracted features are (hopefully) more robust against partial occlusion, illumation and small sample size. Algorithms used for a local feature extraction are Gabor Wavelets ([Wiskott97]), Discrete Cosinus Transform ([Messer06]) and Local Binary Patterns ([AHP04]). It's still an open research question what's the best way to preserve spatial information when applying a local feature extraction, because spatial information is potentially useful information.

### Face Database

Let's get some data to experiment with first. I don't want to do a toy example here. We are doing face recognition, so you'll need some face images! You can either create your own dataset or start with one of the available face databases, http://face-rec.org/databases/ gives you an up-to-date overview. Three interesting databases are (parts of the description are quoted from http://face-rec.org):

- AT&T Facedatabase The AT&T Facedatabase, sometimes also referred to as *ORL Database of Faces*, contains ten different images of each of 40 distinct subjects. For some subjects, the images were taken at different times, varying the lighting, facial expressions (open / closed eyes, smiling / not smiling) and facial details (glasses /

no glasses). All the images were taken against a dark homogeneous background with the subjects in an upright, frontal position (with tolerance for some side movement).

• Yale Facedatabase A, also known as Yalefaces. The AT&T Facedatabase is good for initial tests, but it's a fairly easy database. The Eigenfaces method already has a 97% recognition rate on it, so you won't see any great improvements with other algorithms. The Yale Facedatabase A (also known as Yalefaces) is a more appropriate dataset for initial experiments, because the recognition problem is harder. The database consists of 15 people (14 male, 1 female) each with 11 grayscale images sized $320 \times 243$ pixel. There are changes in the light conditions (center light, left light, right light), facial expressions (happy, normal, sad, sleepy, surprised, wink) and glasses (glasses, no-glasses).

The original images are not cropped and aligned. Please look into the *Appendix* for a Python script, that does the job for you.

• Extended Yale Facedatabase B The Extended Yale Facedatabase B contains 2414 images of 38 different people in its cropped version. The focus of this database is set on extracting features that are robust to illumination, the images have almost no variation in emotion/occlusion/... . I personally think, that this dataset is too large for the experiments I perform in this document. You better use the AT&T Facedatabase for intial testing. A first version of the Yale Facedatabase B was used in [BHK97] to see how the Eigenfaces and Fisherfaces method perform under heavy illumination changes. [Lee05] used the same setup to take 16128 images of 28 people. The Extended Yale Facedatabase B is the merge of the two databases, which is now known as Extended Yalefacedatabase B.

**Preparing the data** Once we have acquired some data, we'll need to read it in our program. In the demo applications I have decided to read the images from a very simple CSV file. Why? Because it's the simplest platform-independent approach I can think of. However, if you know a simpler solution please ping me about it. Basically all the CSV file needs to contain are lines composed of a `filename` followed by a `;` followed by the `label` (as *integer number*), making up a line like this:

```
/path/to/image.ext;0
```

Let's dissect the line. `/path/to/image.ext` is the path to an image, probably something like this if you are in Windows: `C:/faces/person0/image0.jpg`. Then there is the separator `;` and finally we assign the label `0` to the image. Think of the label as the subject (the person) this image belongs to, so same subjects (persons) should have the same label.

Download the AT&T Facedatabase from AT&T Facedatabase and the corresponding CSV file from at.txt, which looks like this (file is without ... of course):

```
./at/s1/1.pgm;0
./at/s1/2.pgm;0
...
./at/s2/1.pgm;1
./at/s2/2.pgm;1
...
./at/s40/1.pgm;39
./at/s40/2.pgm;39
```

Imagine I have extracted the files to `D:/data/at` and have downloaded the CSV file to `D:/data/at.txt`. Then you would simply need to Search & Replace `./` with `D:/data/`. You can do that in an editor of your choice, every sufficiently advanced editor can do this. Once you have a CSV file with valid filenames and labels, you can run any of the demos by passing the path to the CSV file as parameter:

```
facerec_demo.exe D:/data/at.txt
```

**Creating the CSV File**   You don't really want to create the CSV file by hand. I have prepared you a little Python script `create_csv.py` (you find it at `src/create_csv.py` coming with this tutorial) that automatically creates you a CSV file. If you have your images in hierarchie like this (`/basepath/<subject>/<image.ext>`):

```
philipp@mango:~/facerec/data/at$ tree
.
|-- s1
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
|-- s2
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
...
|-- s40
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
```

Then simply call create_csv.py with the path to the folder, just like this and you could save the output:

```
philipp@mango:~/facerec/data$ python create_csv.py
at/s13/2.pgm;0
at/s13/7.pgm;0
at/s13/6.pgm;0
at/s13/9.pgm;0
at/s13/5.pgm;0
at/s13/3.pgm;0
at/s13/4.pgm;0
at/s13/10.pgm;0
at/s13/8.pgm;0
at/s13/1.pgm;0
at/s17/2.pgm;1
at/s17/7.pgm;1
at/s17/6.pgm;1
at/s17/9.pgm;1
at/s17/5.pgm;1
at/s17/3.pgm;1
[...]
```

Please see the *Appendix* for additional informations.

### Eigenfaces

The problem with the image representation we are given is its high dimensionality. Two-dimensional $p \times q$ grayscale images span a $m = pq$-dimensional vector space, so an image with $100 \times 100$ pixels lies in a $10,000$-dimensional image space already. The question is: Are all dimensions equally useful for us? We can only make a decision if there's any variance in data, so what we are looking for are the components that account for most of the information. The Principal Component Analysis (PCA) was independently proposed by Karl Pearson (1901) and Harold Hotelling (1933) to turn a set of possibly correlated variables into a smaller set of uncorrelated variables. The idea is, that a high-dimensional dataset is often described by correlated variables and therefore only a few meaningful dimensions account for most of the information. The PCA method finds the directions with the greatest variance in the data, called principal components.

**Algorithmic Description**   Let $X = \{x_1, x_2, \ldots, x_n\}$ be a random vector with observations $x_i \in R^d$.

1. Compute the mean $\mu$

$$\mu = \frac{1}{n} \sum_{i=1}^{n} x_i$$

2. Compute the the Covariance Matrix $S$

$$S = \frac{1}{n} \sum_{i=1}^{n} (x_i - \mu)(x_i - \mu)^T,$$

3. Compute the eigenvalues $\lambda_i$ and eigenvectors $v_i$ of S

$$Sv_i = \lambda_i v_i, i = 1, 2, \ldots, n$$

4. Order the eigenvectors descending by their eigenvalue. The k principal components are the eigenvectors corresponding to the k largest eigenvalues.

The k principal components of the observed vector x are then given by:

$$y = W^T(x - \mu)$$

where $W = (v_1, v_2, \ldots, v_k)$.

The reconstruction from the PCA basis is given by:

$$x = Wy + \mu$$

where $W = (v_1, v_2, \ldots, v_k)$.

The Eigenfaces method then performs face recognition by:

- Projecting all training samples into the PCA subspace.
- Projecting the query image into the PCA subspace.
- Finding the nearest neighbor between the projected training images and the projected query image.

Still there's one problem left to solve. Imagine we are given 400 images sized $100 \times 100$ pixel. The Principal Component Analysis solves the covariance matrix $S = XX^T$, where $size(X) = 10000 \times 400$ in our example. You would end up with a $10000 \times 10000$ matrix, roughly 0.8GB. Solving this problem isn't feasible, so we'll need to apply a trick. From your linear algebra lessons you know that a $M \times N$ matrix with $M > N$ can only have $N - 1$ non-zero eigenvalues. So it's possible to take the eigenvalue decomposition $S = X^TX$ of size $N \times N$ instead:

$$X^TXv_i = \lambda_i vi$$

and get the original eigenvectors of $S = XX^T$ with a left multiplication of the data matrix:

$$XX^T(Xv_i) = \lambda_i(Xv_i)$$

The resulting eigenvectors are orthogonal, to get orthonormal eigenvectors they need to be normalized to unit length. I don't want to turn this into a publication, so please look into [Duda01] for the derivation and proof of the equations.

**Eigenfaces in OpenCV**  For the first source code example, I'll go through it with you. I am first giving you the whole source code listing, and after this we'll look at the most important lines in detail. Please note: every source code listing is commented in detail, so you should have no problems following it.

```
1   /*
2    * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
3    * Released to public domain under terms of the BSD Simplified license.
4    *
5    * Redistribution and use in source and binary forms, with or without
6    * modification, are permitted provided that the following conditions are met:
7    *   * Redistributions of source code must retain the above copyright
8    *     notice, this list of conditions and the following disclaimer.
9    *   * Redistributions in binary form must reproduce the above copyright
10   *     notice, this list of conditions and the following disclaimer in the
11   *     documentation and/or other materials provided with the distribution.
12   *   * Neither the name of the organization nor the names of its contributors
13   *     may be used to endorse or promote products derived from this software
14   *     without specific prior written permission.
15   *
16   *   See <http://www.opensource.org/licenses/bsd-license>
17   */
18
19  #include "opencv2/core/core.hpp"
20  #include "opencv2/contrib/contrib.hpp"
21  #include "opencv2/highgui/highgui.hpp"
22
23  #include <iostream>
24  #include <fstream>
25  #include <sstream>
26
27  using namespace cv;
28  using namespace std;
29
30  static Mat norm_0_255(InputArray _src) {
31      Mat src = _src.getMat();
32      // Create and return normalized image:
33      Mat dst;
34      switch(src.channels()) {
35      case 1:
36          cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC1);
37          break;
38      case 3:
39          cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC3);
40          break;
41      default:
42          src.copyTo(dst);
43          break;
44      }
45      return dst;
46  }
47
48  static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
49      std::ifstream file(filename.c_str(), ifstream::in);
50      if (!file) {
51          string error_message = "No valid input file was given, please check the given filename.";
52          CV_Error(CV_StsBadArg, error_message);
53      }
54      string line, path, classlabel;
55      while (getline(file, line)) {
56          stringstream liness(line);
57          getline(liness, path, separator);
58          getline(liness, classlabel);
```

```
59              if(!path.empty() && !classlabel.empty()) {
60                  images.push_back(imread(path, 0));
61                  labels.push_back(atoi(classlabel.c_str()));
62              }
63          }
64      }
65
66      int main(int argc, const char *argv[]) {
67          // Check for valid command line arguments, print usage
68          // if no arguments were given.
69          if (argc < 2) {
70              cout << "usage: " << argv[0] << " <csv.ext> <output_folder> " << endl;
71              exit(1);
72          }
73          string output_folder;
74          if (argc == 3) {
75              output_folder = string(argv[2]);
76          }
77          // Get the path to your CSV.
78          string fn_csv = string(argv[1]);
79          // These vectors hold the images and corresponding labels.
80          vector<Mat> images;
81          vector<int> labels;
82          // Read in the data. This can fail if no valid
83          // input filename is given.
84          try {
85              read_csv(fn_csv, images, labels);
86          } catch (cv::Exception& e) {
87              cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
88              // nothing more we can do
89              exit(1);
90          }
91          // Quit if there are not enough images for this demo.
92          if(images.size() <= 1) {
93              string error_message = "This demo needs at least 2 images to work. Please add more images to your data set!";
94              CV_Error(CV_StsError, error_message);
95          }
96          // Get the height from the first image. We'll need this
97          // later in code to reshape the images to their original
98          // size:
99          int height = images[0].rows;
100         // The following lines simply get the last images from
101         // your dataset and remove it from the vector. This is
102         // done, so that the training data (which we learn the
103         // cv::FaceRecognizer on) and the test data we test
104         // the model with, do not overlap.
105         Mat testSample = images[images.size() - 1];
106         int testLabel = labels[labels.size() - 1];
107         images.pop_back();
108         labels.pop_back();
109         // The following lines create an Eigenfaces model for
110         // face recognition and train it with the images and
111         // labels read from the given CSV file.
112         // This here is a full PCA, if you just want to keep
113         // 10 principal components (read Eigenfaces), then call
114         // the factory method like this:
115         //
116         //      cv::createEigenFaceRecognizer(10);
```

```
117      //
118      // If you want to create a FaceRecognizer with a
119      // confidence threshold (e.g. 123.0), call it with:
120      //
121      //      cv::createEigenFaceRecognizer(10, 123.0);
122      //
123      // If you want to use _all_ Eigenfaces and have a threshold,
124      // then call the method like this:
125      //
126      //      cv::createEigenFaceRecognizer(0, 123.0);
127      //
128      Ptr<FaceRecognizer> model = createEigenFaceRecognizer();
129      model->train(images, labels);
130      // The following line predicts the label of a given
131      // test image:
132      int predictedLabel = model->predict(testSample);
133      //
134      // To get the confidence of a prediction call the model with:
135      //
136      //      int predictedLabel = -1;
137      //      double confidence = 0.0;
138      //      model->predict(testSample, predictedLabel, confidence);
139      //
140      string result_message = format("Predicted class = %d / Actual class = %d.", predictedLabel, testLabel);
141      cout << result_message << endl;
142      // Here is how to get the eigenvalues of this Eigenfaces model:
143      Mat eigenvalues = model->getMat("eigenvalues");
144      // And we can do the same to display the Eigenvectors (read Eigenfaces):
145      Mat W = model->getMat("eigenvectors");
146      // Get the sample mean from the training data
147      Mat mean = model->getMat("mean");
148      // Display or save:
149      if(argc == 2) {
150          imshow("mean", norm_0_255(mean.reshape(1, images[0].rows)));
151      } else {
152          imwrite(format("%s/mean.png", output_folder.c_str()), norm_0_255(mean.reshape(1, images[0].rows)));
153      }
154      // Display or save the Eigenfaces:
155      for (int i = 0; i < min(10, W.cols); i++) {
156          string msg = format("Eigenvalue #%d = %.5f", i, eigenvalues.at<double>(i));
157          cout << msg << endl;
158          // get eigenvector #i
159          Mat ev = W.col(i).clone();
160          // Reshape to original size & normalize to [0...255] for imshow.
161          Mat grayscale = norm_0_255(ev.reshape(1, height));
162          // Show the image & apply a Jet colormap for better sensing.
163          Mat cgrayscale;
164          applyColorMap(grayscale, cgrayscale, COLORMAP_JET);
165          // Display or save:
166          if(argc == 2) {
167              imshow(format("eigenface_%d", i), cgrayscale);
168          } else {
169              imwrite(format("%s/eigenface_%d.png", output_folder.c_str(), i), norm_0_255(cgrayscale));
170          }
171      }
172
173      // Display or save the image reconstruction at some predefined steps:
174      for(int num_components = min(W.cols, 10); num_components < min(W.cols, 300); num_components+=15) {
```
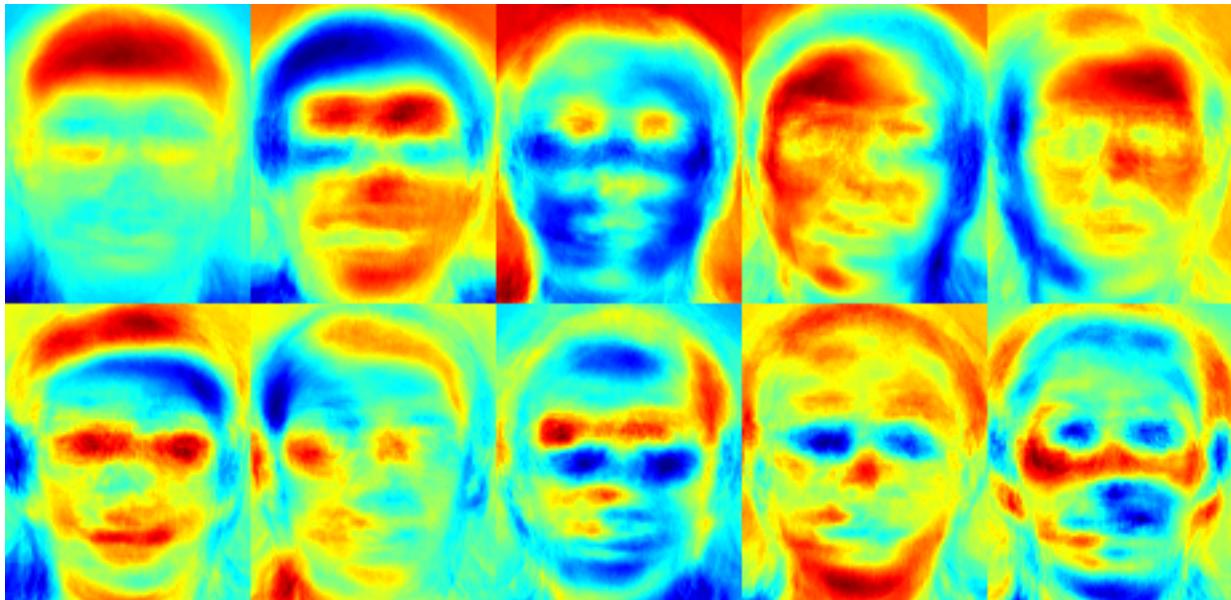
```
175         // slice the eigenvectors from the model
176         Mat evs = Mat(W, Range::all(), Range(0, num_components));
177         Mat projection = subspaceProject(evs, mean, images[0].reshape(1,1));
178         Mat reconstruction = subspaceReconstruct(evs, mean, projection);
179         // Normalize the result:
180         reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
181         // Display or save:
182         if(argc == 2) {
183             imshow(format("eigenface_reconstruction_%d", num_components), reconstruction);
184         } else {
185             imwrite(format("%s/eigenface_reconstruction_%d.png", output_folder.c_str(), num_components), reconstructio
186         }
187     }
188     // Display if we are not writing to an output folder:
189     if(argc == 2) {
190         waitKey(0);
191     }
192     return 0;
193 }
```

The source code for this demo application is also available in the `src` folder coming with this documentation:

- `src/facerec_eigenfaces.cpp`

I've used the jet colormap, so you can see how the grayscale values are distributed within the specific Eigenfaces. You can see, that the Eigenfaces do not only encode facial features, but also the illumination in the images (see the left light in Eigenface #4, right light in Eigenfaces #5):



We've already seen, that we can reconstruct a face from its lower dimensional approximation. So let's see how many Eigenfaces are needed for a good reconstruction. I'll do a subplot with $10, 30, \ldots, 310$ Eigenfaces:

```
// Display or save the image reconstruction at some predefined steps:
for(int num_components = 10; num_components < 300; num_components+=15) {
    // slice the eigenvectors from the model
    Mat evs = Mat(W, Range::all(), Range(0, num_components));
    Mat projection = subspaceProject(evs, mean, images[0].reshape(1,1));
    Mat reconstruction = subspaceReconstruct(evs, mean, projection);
    // Normalize the result:
```

```
    reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
    // Display or save:
    if(argc == 2) {
        imshow(format("eigenface_reconstruction_%d", num_components), reconstruction);
    } else {
        imwrite(format("%s/eigenface_reconstruction_%d.png", output_folder.c_str(), num_components), reconstruction);
    }
}
```

10 Eigenvectors are obviously not sufficient for a good image reconstruction, 50 Eigenvectors may already be sufficient to encode important facial features. You'll get a good reconstruction with approximately 300 Eigenvectors for the AT&T Facedatabase. There are rule of thumbs how many Eigenfaces you should choose for a successful face recognition, but it heavily depends on the input data. [Zhao03] is the perfect point to start researching for this:

**Fisherfaces**

The Principal Component Analysis (PCA), which is the core of the Eigenfaces method, finds a linear combination of features that maximizes the total variance in data. While this is clearly a powerful way to represent data, it doesn't consider any classes and so a lot of discriminative information *may* be lost when throwing components away. Imagine a situation where the variance in your data is generated by an external source, let it be the light. The components identified by a PCA do not necessarily contain any discriminative information at all, so the projected samples are smeared together and a classification becomes impossible (see http://www.bytefish.de/wiki/pca_lda_with_gnu_octave for an example).

The Linear Discriminant Analysis performs a class-specific dimensionality reduction and was invented by the great statistician Sir R. A. Fisher. He successfully used it for classifying flowers in his 1936 paper *The use of multiple measurements in taxonomic problems* [Fisher36]. In order to find the combination of features that separates best between classes the Linear Discriminant Analysis maximizes the ratio of between-classes to within-classes scatter, instead of maximizing the overall scatter. The idea is simple: same classes should cluster tightly together, while different classes are as far away as possible from each other in the lower-dimensional representation. This was also recognized by Belhumeur, Hespanha and Kriegman and so they applied a Discriminant Analysis to face recognition in [BHK97].

**Algorithmic Description** Let X be a random vector with samples drawn from c classes:

$$X = \{X_1, X_2, \ldots, X_c\}$$
$$X_i = \{x_1, x_2, \ldots, x_n\}$$

The scatter matrices $S_B$ and $S\_\{W\}$ are calculated as:

$$S_B = \sum_{i=1}^{c} N_i (\mu_i - \mu)(\mu_i - \mu)^\mathsf{T}$$
$$S_W = \sum_{i=1}^{c} \sum_{x_j \in X_i} (x_j - \mu_i)(x_j - \mu_i)^\mathsf{T}$$

, where $\mu$ is the total mean:

$$\mu = \frac{1}{N} \sum_{i=1}^{N} x_i$$

And $\mu_i$ is the mean of class $i \in \{1, \ldots, c\}$:

$$\mu_i = \frac{1}{|X_i|} \sum_{x_j \in X_i} x_j$$

Fisher's classic algorithm now looks for a projection $W$, that maximizes the class separability criterion:

$$W_{opt} = \arg\max_W \frac{|W^\mathsf{T} S_B W|}{|W^\mathsf{T} S_W W|}$$

Following [BHK97], a solution for this optimization problem is given by solving the General Eigenvalue Problem:

$$S_B v_i = \lambda_i S_w v_i$$
$$S_W^{-1} S_B v_i = \lambda_i v_i$$

There's one problem left to solve: The rank of $S_W$ is at most $(N - c)$, with N samples and c classes. In pattern recognition problems the number of samples N is almost always samller than the dimension of the input data (the

number of pixels), so the scatter matrix $S_W$ becomes singular (see [RJ91]). In [BHK97] this was solved by performing a Principal Component Analysis on the data and projecting the samples into the $(N - c)$-dimensional space. A Linear Discriminant Analysis was then performed on the reduced data, because $S_W$ isn't singular anymore.

The optimization problem can then be rewritten as:

$$W_{pca} = \arg\max_W |W^\mathsf{T} S_\mathsf{T} W|$$

$$W_{fld} = \arg\max_W \frac{|W^\mathsf{T} W_{pca}^\mathsf{T} S_B W_{pca} W|}{|W^\mathsf{T} W_{pca}^\mathsf{T} S_W W_{pca} W|}$$

The transformation matrix $W$, that projects a sample into the $(c - 1)$-dimensional space is then given by:

$$W = W_{fld}^\mathsf{T} W_{pca}^\mathsf{T}$$

### Fisherfaces in OpenCV

```
/*
 * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
 * Released to public domain under terms of the BSD Simplified license.
 *
 * Redistribution and use in source and binary forms, with or without
 * modification, are permitted provided that the following conditions are met:
 *   * Redistributions of source code must retain the above copyright
 *     notice, this list of conditions and the following disclaimer.
 *   * Redistributions in binary form must reproduce the above copyright
 *     notice, this list of conditions and the following disclaimer in the
 *     documentation and/or other materials provided with the distribution.
 *   * Neither the name of the organization nor the names of its contributors
 *     may be used to endorse or promote products derived from this software
 *     without specific prior written permission.
 *
 *   See <http://www.opensource.org/licenses/bsd-license>
 */

#include "opencv2/core/core.hpp"
#include "opencv2/contrib/contrib.hpp"
#include "opencv2/highgui/highgui.hpp"

#include <iostream>
#include <fstream>
#include <sstream>

using namespace cv;
using namespace std;

static Mat norm_0_255(InputArray _src) {
    Mat src = _src.getMat();
    // Create and return normalized image:
    Mat dst;
    switch(src.channels()) {
    case 1:
        cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC1);
        break;
    case 3:
        cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC3);
        break;
    default:
        src.copyTo(dst);
```

```
43          break;
44      }
45      return dst;
46  }
47
48  static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
49      std::ifstream file(filename.c_str(), ifstream::in);
50      if (!file) {
51          string error_message = "No valid input file was given, please check the given filename.";
52          CV_Error(CV_StsBadArg, error_message);
53      }
54      string line, path, classlabel;
55      while (getline(file, line)) {
56          stringstream liness(line);
57          getline(liness, path, separator);
58          getline(liness, classlabel);
59          if(!path.empty() && !classlabel.empty()) {
60              images.push_back(imread(path, 0));
61              labels.push_back(atoi(classlabel.c_str()));
62          }
63      }
64  }
65
66  int main(int argc, const char *argv[]) {
67      // Check for valid command line arguments, print usage
68      // if no arguments were given.
69      if (argc < 2) {
70          cout << "usage: " << argv[0] << " <csv.ext> <output_folder> " << endl;
71          exit(1);
72      }
73      string output_folder;
74      if (argc == 3) {
75          output_folder = string(argv[2]);
76      }
77      // Get the path to your CSV.
78      string fn_csv = string(argv[1]);
79      // These vectors hold the images and corresponding labels.
80      vector<Mat> images;
81      vector<int> labels;
82      // Read in the data. This can fail if no valid
83      // input filename is given.
84      try {
85          read_csv(fn_csv, images, labels);
86      } catch (cv::Exception& e) {
87          cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
88          // nothing more we can do
89          exit(1);
90      }
91      // Quit if there are not enough images for this demo.
92      if(images.size() <= 1) {
93          string error_message = "This demo needs at least 2 images to work. Please add more images to your data set!";
94          CV_Error(CV_StsError, error_message);
95      }
96      // Get the height from the first image. We'll need this
97      // later in code to reshape the images to their original
98      // size:
99      int height = images[0].rows;
100     // The following lines simply get the last images from
```

```
101        // your dataset and remove it from the vector. This is
102        // done, so that the training data (which we learn the
103        // cv::FaceRecognizer on) and the test data we test
104        // the model with, do not overlap.
105        Mat testSample = images[images.size() - 1];
106        int testLabel = labels[labels.size() - 1];
107        images.pop_back();
108        labels.pop_back();
109        // The following lines create an Fisherfaces model for
110        // face recognition and train it with the images and
111        // labels read from the given CSV file.
112        // If you just want to keep 10 Fisherfaces, then call
113        // the factory method like this:
114        //
115        //      cv::createFisherFaceRecognizer(10);
116        //
117        // However it is not useful to discard Fisherfaces! Please
118        // always try to use _all_ available Fisherfaces for
119        // classification.
120        //
121        // If you want to create a FaceRecognizer with a
122        // confidence threshold (e.g. 123.0) and use _all_
123        // Fisherfaces, then call it with:
124        //
125        //      cv::createFisherFaceRecognizer(0, 123.0);
126        //
127        Ptr<FaceRecognizer> model = createFisherFaceRecognizer();
128        model->train(images, labels);
129        // The following line predicts the label of a given
130        // test image:
131        int predictedLabel = model->predict(testSample);
132        //
133        // To get the confidence of a prediction call the model with:
134        //
135        //      int predictedLabel = -1;
136        //      double confidence = 0.0;
137        //      model->predict(testSample, predictedLabel, confidence);
138        //
139        string result_message = format("Predicted class = %d / Actual class = %d.", predictedLabel, testLabel);
140        cout << result_message << endl;
141        // Here is how to get the eigenvalues of this Eigenfaces model:
142        Mat eigenvalues = model->getMat("eigenvalues");
143        // And we can do the same to display the Eigenvectors (read Eigenfaces):
144        Mat W = model->getMat("eigenvectors");
145        // Get the sample mean from the training data
146        Mat mean = model->getMat("mean");
147        // Display or save:
148        if(argc == 2) {
149            imshow("mean", norm_0_255(mean.reshape(1, images[0].rows)));
150        } else {
151            imwrite(format("%s/mean.png", output_folder.c_str()), norm_0_255(mean.reshape(1, images[0].rows)));
152        }
153        // Display or save the first, at most 16 Fisherfaces:
154        for (int i = 0; i < min(16, W.cols); i++) {
155            string msg = format("Eigenvalue #%d = %.5f", i, eigenvalues.at<double>(i));
156            cout << msg << endl;
157            // get eigenvector #i
158            Mat ev = W.col(i).clone();
```

```
159              // Reshape to original size & normalize to [0...255] for imshow.
160              Mat grayscale = norm_0_255(ev.reshape(1, height));
161              // Show the image & apply a Bone colormap for better sensing.
162              Mat cgrayscale;
163              applyColorMap(grayscale, cgrayscale, COLORMAP_BONE);
164              // Display or save:
165              if(argc == 2) {
166                  imshow(format("fisherface_%d", i), cgrayscale);
167              } else {
168                  imwrite(format("%s/fisherface_%d.png", output_folder.c_str(), i), norm_0_255(cgrayscale));
169              }
170          }
171          // Display or save the image reconstruction at some predefined steps:
172          for(int num_component = 0; num_component < min(16, W.cols); num_component++) {
173              // Slice the Fisherface from the model:
174              Mat ev = W.col(num_component);
175              Mat projection = subspaceProject(ev, mean, images[0].reshape(1,1));
176              Mat reconstruction = subspaceReconstruct(ev, mean, projection);
177              // Normalize the result:
178              reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
179              // Display or save:
180              if(argc == 2) {
181                  imshow(format("fisherface_reconstruction_%d", num_component), reconstruction);
182              } else {
183                  imwrite(format("%s/fisherface_reconstruction_%d.png", output_folder.c_str(), num_component), reconstructio
184              }
185          }
186          // Display if we are not writing to an output folder:
187          if(argc == 2) {
188              waitKey(0);
189          }
190          return 0;
191      }
```

The source code for this demo application is also available in the `src` folder coming with this documentation:

- `src/facerec_fisherfaces.cpp`

For this example I am going to use the Yale Facedatabase A, just because the plots are nicer. Each Fisherface has the same length as an original image, thus it can be displayed as an image. The demo shows (or saves) the first, at most 16 Fisherfaces:

The Fisherfaces method learns a class-specific transformation matrix, so the they do not capture illumination as obviously as the Eigenfaces method. The Discriminant Analysis instead finds the facial features to discriminate between the persons. It's important to mention, that the performance of the Fisherfaces heavily depends on the input data as well. Practically said: if you learn the Fisherfaces for well-illuminated pictures only and you try to recognize faces in bad-illuminated scenes, then method is likely to find the wrong components (just because those features may not be predominant on bad illuminated images). This is somewhat logical, since the method had no chance to learn the illumination.

The Fisherfaces allow a reconstruction of the projected image, just like the Eigenfaces did. But since we only identified the features to distinguish between subjects, you can't expect a nice reconstruction of the original image. For the Fisherfaces method we'll project the sample image onto each of the Fisherfaces instead. So you'll have a nice visualization, which feature each of the Fisherfaces describes:

```
// Display or save the image reconstruction at some predefined steps:
for(int num_component = 0; num_component < min(16, W.cols); num_component++) {
    // Slice the Fisherface from the model:
    Mat ev = W.col(num_component);
    Mat projection = subspaceProject(ev, mean, images[0].reshape(1,1));
    Mat reconstruction = subspaceReconstruct(ev, mean, projection);
    // Normalize the result:
    reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
    // Display or save:
    if(argc == 2) {
        imshow(format("fisherface_reconstruction_%d", num_component), reconstruction);
```

```
    } else {
        imwrite(format("%s/fisherface_reconstruction_%d.png", output_folder.c_str(), num_component), reconstruction);
    }
}
```

The differences may be subtle for the human eyes, but you should be able to see some differences:
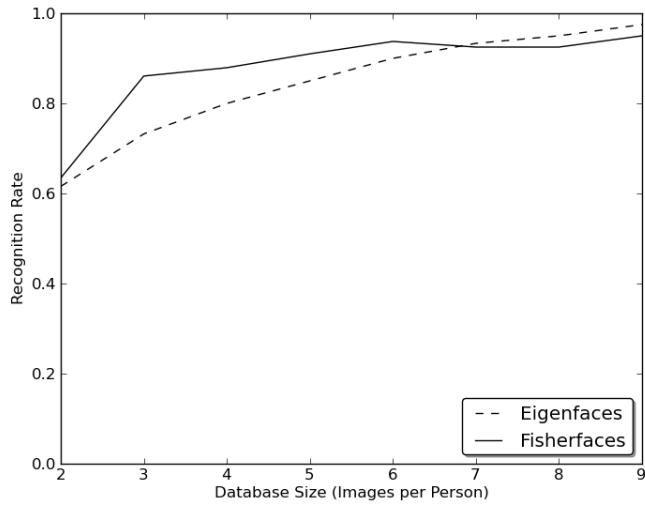


### Local Binary Patterns Histograms

Eigenfaces and Fisherfaces take a somewhat holistic approach to face recognition. You treat your data as a vector somewhere in a high-dimensional image space. We all know high-dimensionality is bad, so a lower-dimensional subspace is identified, where (probably) useful information is preserved. The Eigenfaces approach maximizes the total scatter, which can lead to problems if the variance is generated by an external source, because components with a maximum variance over all classes aren't necessarily useful for classification (see http://www.bytefish.de/wiki/pca_lda_with_gnu_octave). So to preserve some discriminative information we applied a Linear Discriminant Analysis and optimized as described in the Fisherfaces method. The Fisherfaces method worked great... at least for the constrained scenario we've assumed in our model.
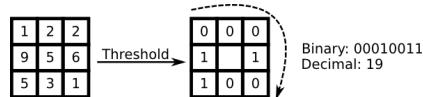
Now real life isn't perfect. You simply can't guarantee perfect light settings in your images or 10 different images of a person. So what if there's only one image for each person? Our covariance estimates for the subspace *may* be horribly wrong, so will the recognition. Remember the Eigenfaces method had a 96% recognition rate on the AT&T Facedatabase? How many images do we actually need to get such useful estimates? Here are the Rank-1 recognition

rates of the Eigenfaces and Fisherfaces method on the AT&T Facedatabase, which is a fairly easy image database:



So in order to get good recognition rates you'll need at least 8(+-1) images for each person and the Fisherfaces method doesn't really help here. The above experiment is a 10-fold cross validated result carried out with the facerec framework at: https://github.com/bytefish/facerec. This is not a publication, so I won't back these figures with a deep mathematical analysis. Please have a look into [KM01] for a detailed analysis of both methods, when it comes to small training datasets.

So some research concentrated on extracting local features from images. The idea is to not look at the whole image as a high-dimensional vector, but describe only local features of an object. The features you extract this way will have a low-dimensionality implicitly. A fine idea! But you'll soon observe the image representation we are given doesn't only suffer from illumination variations. Think of things like scale, translation or rotation in images - your local description has to be at least a bit robust against those things. Just like `SIFT`, the Local Binary Patterns methodology has its roots in 2D texture analysis. The basic idea of Local Binary Patterns is to summarize the local structure in an image by comparing each pixel with its neighborhood. Take a pixel as center and threshold its neighbors against. If the intensity of the center pixel is greater-equal its neighbor, then denote it with 1 and 0 if not. You'll end up with a binary number for each pixel, just like 11001111. So with 8 surrounding pixels you'll end up with 2^8 possible combinations, called *Local Binary Patterns* or sometimes referred to as *LBP codes*. The first LBP operator described in literature actually used a fixed 3 x 3 neighborhood just like this:



**Algorithmic Description**    A more formal description of the LBP operator can be given as:
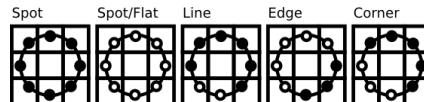
$$LBP(x_c, y_c) = \sum_{p=0}^{P-1} 2^p s(i_p - i_c)$$

, with $(x_c, y_c)$ as central pixel with intensity $i_c$; and $i_n$ being the intensity of the the neighbor pixel. $s$ is the sign function defined as:

$$s(x) = \begin{cases} 1 & \text{if } x \geq 0 \\ 0 & \text{else} \end{cases} \tag{15.1}$$

This description enables you to capture very fine grained details in images. In fact the authors were able to compete with state of the art results for texture classification. Soon after the operator was published it was noted, that a fixed

neighborhood fails to encode details differing in scale. So the operator was extended to use a variable neighborhood in [AHP04]. The idea is to align an arbitrary number of neighbors on a circle with a variable radius, which enables to capture the following neighborhoods:



For a given Point $(x_c, y_c)$ the position of the neighbor $(x_p, y_p), p \in P$ can be calculated by:
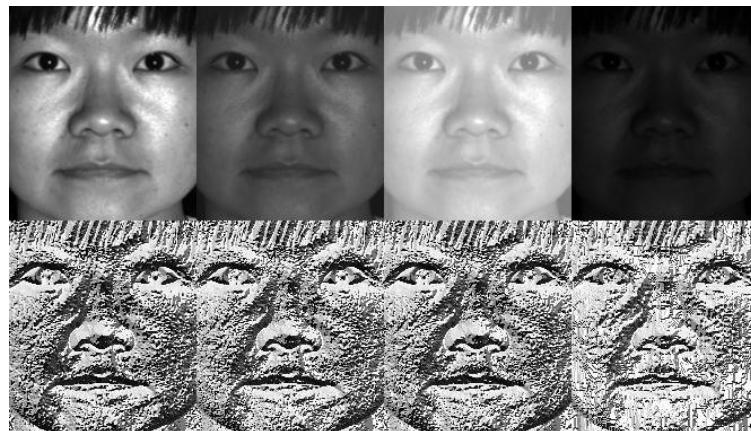
$$x_p = x_c + R\cos(\frac{2\pi p}{P})$$

$$y_p = y_c - R\sin(\frac{2\pi p}{P})$$

Where R is the radius of the circle and P is the number of sample points.

The operator is an extension to the original LBP codes, so it's sometimes called *Extended LBP* (also referred to as *Circular LBP*) . If a points coordinate on the circle doesn't correspond to image coordinates, the point get's interpolated. Computer science has a bunch of clever interpolation schemes, the OpenCV implementation does a bilinear interpolation:

$$f(x, y) \approx \begin{bmatrix} 1 - x & x \end{bmatrix} \begin{bmatrix} f(0, 0) & f(0, 1) \\ f(1, 0) & f(1, 1) \end{bmatrix} \begin{bmatrix} 1 - y \\ y \end{bmatrix}.$$

By definition the LBP operator is robust against monotonic gray scale transformations. We can easily verify this by looking at the LBP image of an artificially modified image (so you see what an LBP image looks like!):



So what's left to do is how to incorporate the spatial information in the face recognition model. The representation proposed by Ahonen et. al [AHP04] is to divide the LBP image into $m$ local regions and extract a histogram from each. The spatially enhanced feature vector is then obtained by concatenating the local histograms (**not merging them**). These histograms are called *Local Binary Patterns Histograms*.

**Local Binary Patterns Histograms in OpenCV**

```
1  /*
2   * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
3   * Released to public domain under terms of the BSD Simplified license.
4   *
5   * Redistribution and use in source and binary forms, with or without
6   * modification, are permitted provided that the following conditions are met:
7   *     * Redistributions of source code must retain the above copyright
8   *       notice, this list of conditions and the following disclaimer.
```

```
9   *   * Redistributions in binary form must reproduce the above copyright
10  *     notice, this list of conditions and the following disclaimer in the
11  *     documentation and/or other materials provided with the distribution.
12  *   * Neither the name of the organization nor the names of its contributors
13  *     may be used to endorse or promote products derived from this software
14  *     without specific prior written permission.
15  *
16  *   See <http://www.opensource.org/licenses/bsd-license>
17  */

18
19  #include "opencv2/core/core.hpp"
20  #include "opencv2/contrib/contrib.hpp"
21  #include "opencv2/highgui/highgui.hpp"
22
23  #include <iostream>
24  #include <fstream>
25  #include <sstream>
26
27  using namespace cv;
28  using namespace std;
29
30  static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
31      std::ifstream file(filename.c_str(), ifstream::in);
32      if (!file) {
33          string error_message = "No valid input file was given, please check the given filename.";
34          CV_Error(CV_StsBadArg, error_message);
35      }
36      string line, path, classlabel;
37      while (getline(file, line)) {
38          stringstream liness(line);
39          getline(liness, path, separator);
40          getline(liness, classlabel);
41          if(!path.empty() && !classlabel.empty()) {
42              images.push_back(imread(path, 0));
43              labels.push_back(atoi(classlabel.c_str()));
44          }
45      }
46  }
47
48  int main(int argc, const char *argv[]) {
49      // Check for valid command line arguments, print usage
50      // if no arguments were given.
51      if (argc != 2) {
52          cout << "usage: " << argv[0] << " <csv.ext>" << endl;
53          exit(1);
54      }
55      // Get the path to your CSV.
56      string fn_csv = string(argv[1]);
57      // These vectors hold the images and corresponding labels.
58      vector<Mat> images;
59      vector<int> labels;
60      // Read in the data. This can fail if no valid
61      // input filename is given.
62      try {
63          read_csv(fn_csv, images, labels);
64      } catch (cv::Exception& e) {
65          cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
66          // nothing more we can do
```

```
67          exit(1);
68      }
69      // Quit if there are not enough images for this demo.
70      if(images.size() <= 1) {
71          string error_message = "This demo needs at least 2 images to work. Please add more images to your data set!";
72          CV_Error(CV_StsError, error_message);
73      }
74      // Get the height from the first image. We'll need this
75      // later in code to reshape the images to their original
76      // size:
77      int height = images[0].rows;
78      // The following lines simply get the last images from
79      // your dataset and remove it from the vector. This is
80      // done, so that the training data (which we learn the
81      // cv::FaceRecognizer on) and the test data we test
82      // the model with, do not overlap.
83      Mat testSample = images[images.size() - 1];
84      int testLabel = labels[labels.size() - 1];
85      images.pop_back();
86      labels.pop_back();
87      // The following lines create an LBPH model for
88      // face recognition and train it with the images and
89      // labels read from the given CSV file.
90      //
91      // The LBPHFaceRecognizer uses Extended Local Binary Patterns
92      // (it's probably configurable with other operators at a later
93      // point), and has the following default values
94      //
95      //      radius = 1
96      //      neighbors = 8
97      //      grid_x = 8
98      //      grid_y = 8
99      //
100     // So if you want a LBPH FaceRecognizer using a radius of
101     // 2 and 16 neighbors, call the factory method with:
102     //
103     //      cv::createLBPHFaceRecognizer(2, 16);
104     //
105     // And if you want a threshold (e.g. 123.0) call it with its default values:
106     //
107     //      cv::createLBPHFaceRecognizer(1,8,8,8,123.0)
108     //
109     Ptr<FaceRecognizer> model = createLBPHFaceRecognizer();
110     model->train(images, labels);
111     // The following line predicts the label of a given
112     // test image:
113     int predictedLabel = model->predict(testSample);
114     //
115     // To get the confidence of a prediction call the model with:
116     //
117     //      int predictedLabel = -1;
118     //      double confidence = 0.0;
119     //      model->predict(testSample, predictedLabel, confidence);
120     //
121     string result_message = format("Predicted class = %d / Actual class = %d.", predictedLabel, testLabel);
122     cout << result_message << endl;
123     // Sometimes you'll need to get/set internal model data,
124     // which isn't exposed by the public cv::FaceRecognizer.
```

---

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```
125      // Since each cv::FaceRecognizer is derived from a
126      // cv::Algorithm, you can query the data.
127      //
128      // First we'll use it to set the threshold of the FaceRecognizer
129      // to 0.0 without retraining the model. This can be useful if
130      // you are evaluating the model:
131      //
132      model->set("threshold", 0.0);
133      // Now the threshold of this model is set to 0.0. A prediction
134      // now returns -1, as it's impossible to have a distance below
135      // it
136      predictedLabel = model->predict(testSample);
137      cout << "Predicted class = " << predictedLabel << endl;
138      // Show some informations about the model, as there's no cool
139      // Model data to display as in Eigenfaces/Fisherfaces.
140      // Due to efficiency reasons the LBP images are not stored
141      // within the model:
142      cout << "Model Information:" << endl;
143      string model_info = format("\tLBPH(radius=%i, neighbors=%i, grid_x=%i, grid_y=%i, threshold=%.2f)",
144              model->getInt("radius"),
145              model->getInt("neighbors"),
146              model->getInt("grid_x"),
147              model->getInt("grid_y"),
148              model->getDouble("threshold"));
149      cout << model_info << endl;
150      // We could get the histograms for example:
151      vector<Mat> histograms = model->getMatVector("histograms");
152      // But should I really visualize it? Probably the length is interesting:
153      cout << "Size of the histograms: " << histograms[0].total() << endl;
154      return 0;
155  }
```

The source code for this demo application is also available in the `src` folder coming with this documentation:

- `src/facerec_lbph.cpp`

**Conclusion**

You've learned how to use the new `FaceRecognizer` in real applications. After reading the document you also know how the algorithms work, so now it's time for you to experiment with the available algorithms. Use them, improve them and let the OpenCV community participate!

**Credits**

This document wouldn't be possible without the kind permission to use the face images of the *AT&T Database of Faces* and the *Yale Facedatabase A/B*.

**The Database of Faces** ** Important: when using these images, please give credit to "AT&T Laboratories, Cambridge." **

The Database of Faces, formerly *The ORL Database of Faces*, contains a set of face images taken between April 1992 and April 1994. The database was used in the context of a face recognition project carried out in collaboration with the Speech, Vision and Robotics Group of the Cambridge University Engineering Department.

There are ten different images of each of 40 distinct subjects. For some subjects, the images were taken at different times, varying the lighting, facial expressions (open / closed eyes, smiling / not smiling) and facial details (glasses / no glasses). All the images were taken against a dark homogeneous background with the subjects in an upright, frontal position (with tolerance for some side movement).

The files are in PGM format. The size of each image is 92x112 pixels, with 256 grey levels per pixel. The images are organised in 40 directories (one for each subject), which have names of the form sX, where X indicates the subject number (between 1 and 40). In each of these directories, there are ten different images of that subject, which have names of the form Y.pgm, where Y is the image number for that subject (between 1 and 10).

A copy of the database can be retrieved from: http://www.cl.cam.ac.uk/research/dtg/attarchive/pub/data/att_faces.zip.

**Yale Facedatabase A**    *With the permission of the authors I am allowed to show a small number of images (say subject 1 and all the variations) and all images such as Fisherfaces and Eigenfaces from either Yale Facedatabase A or the Yale Facedatabase B.*

The Yale Face Database A (size 6.4MB) contains 165 grayscale images in GIF format of 15 individuals. There are 11 images per subject, one per different facial expression or configuration: center-light, w/glasses, happy, left-light, w/no glasses, normal, right-light, sad, sleepy, surprised, and wink. (Source: http://cvc.yale.edu/projects/yalefaces/yalefaces.html)

**Yale Facedatabase B**    *With the permission of the authors I am allowed to show a small number of images (say subject 1 and all the variations) and all images such as Fisherfaces and Eigenfaces from either Yale Facedatabase A or the Yale Facedatabase B.*

The extended Yale Face Database B contains 16128 images of 28 human subjects under 9 poses and 64 illumination conditions. The data format of this database is the same as the Yale Face Database B. Please refer to the homepage of the Yale Face Database B (or one copy of this page) for more detailed information of the data format.

You are free to use the extended Yale Face Database B for research purposes. All publications which use this database should acknowledge the use of "the Exteded Yale Face Database B" and reference Athinodoros Georghiades, Peter Belhumeur, and David Kriegman's paper, "From Few to Many: Illumination Cone Models for Face Recognition under Variable Lighting and Pose", PAMI, 2001, [bibtex].

The extended database as opposed to the original Yale Face Database B with 10 subjects was first reported by Kuang-Chih Lee, Jeffrey Ho, and David Kriegman in "Acquiring Linear Subspaces for Face Recognition under Variable Lighting, PAMI, May, 2005 [pdf]." All test image data used in the experiments are manually aligned, cropped, and then re-sized to 168x192 images. If you publish your experimental results with the cropped images, please reference the PAMI2005 paper as well. (Source: http://vision.ucsd.edu/~leekc/ExtYaleDatabase/ExtYaleB.html)

**Literature**

**Appendix**

**Creating the CSV File**    You don't really want to create the CSV file by hand. I have prepared you a little Python script `create_csv.py` (you find it at `/src/create_csv.py` coming with this tutorial) that automatically creates you a CSV file. If you have your images in hierarchie like this (`/basepath/<subject>/<image.ext>`):

```
philipp@mango:~/facerec/data/at$ tree
.
|-- s1
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
|-- s2
```

```
|    |-- 1.pgm
|    |-- ...
|    |-- 10.pgm
...
|-- s40
|    |-- 1.pgm
|    |-- ...
|    |-- 10.pgm
```

Then simply call `create_csv.py` with the path to the folder, just like this and you could save the output:

```
philipp@mango:~/facerec/data$ python create_csv.py
at/s13/2.pgm;0
at/s13/7.pgm;0
at/s13/6.pgm;0
at/s13/9.pgm;0
at/s13/5.pgm;0
at/s13/3.pgm;0
at/s13/4.pgm;0
at/s13/10.pgm;0
at/s13/8.pgm;0
at/s13/1.pgm;0
at/s17/2.pgm;1
at/s17/7.pgm;1
at/s17/6.pgm;1
at/s17/9.pgm;1
at/s17/5.pgm;1
at/s17/3.pgm;1
[...]
```

Here is the script, if you can't find it:

```python
1   import sys
2   import os.path
3
4   # This is a tiny script to help you creating a CSV file from a face
5   # database with a similar hierarchie:
6   #
7   #   philipp@mango:~/facerec/data/at$ tree
8   #   .
9   #   |-- README
10  #   |-- s1
11  #   |    |-- 1.pgm
12  #   |    |-- ...
13  #   |    |-- 10.pgm
14  #   |-- s2
15  #   |    |-- 1.pgm
16  #   |    |-- ...
17  #   |    |-- 10.pgm
18  #   ...
19  #   |-- s40
20  #   |    |-- 1.pgm
21  #   |    |-- ...
22  #   |    |-- 10.pgm
23  #
24
25  if __name__ == "__main__":
26
27      if len(sys.argv) != 2:
```

```
28          print "usage: create_csv <base_path>"
29          sys.exit(1)
30
31      BASE_PATH=sys.argv[1]
32      SEPARATOR=";"
33
34      label = 0
35      for dirname, dirnames, filenames in os.walk(BASE_PATH):
36          for subdirname in dirnames:
37              subject_path = os.path.join(dirname, subdirname)
38              for filename in os.listdir(subject_path):
39                  abs_path = "%s/%s" % (subject_path, filename)
40                  print "%s%s%d" % (abs_path, SEPARATOR, label)
41              label = label + 1
```

**Aligning Face Images** An accurate alignment of your image data is especially important in tasks like emotion detection, were you need as much detail as possible. Believe me... You don't want to do this by hand. So I've prepared you a tiny Python script. The code is really easy to use. To scale, rotate and crop the face image you just need to call *CropFace(image, eye_left, eye_right, offset_pct, dest_sz)*, where:

- *eye_left* is the position of the left eye

- *eye_right* is the position of the right eye

- *offset_pct* is the percent of the image you want to keep next to the eyes (horizontal, vertical direction)

- *dest_sz* is the size of the output image

If you are using the same *offset_pct* and *dest_sz* for your images, they are all aligned at the eyes.

```
1   #!/usr/bin/env python
2   # Software License Agreement (BSD License)
3   #
4   # Copyright (c) 2012, Philipp Wagner
5   # All rights reserved.
6   #
7   # Redistribution and use in source and binary forms, with or without
8   # modification, are permitted provided that the following conditions
9   # are met:
10  #
11  #  * Redistributions of source code must retain the above copyright
12  #    notice, this list of conditions and the following disclaimer.
13  #  * Redistributions in binary form must reproduce the above
14  #    copyright notice, this list of conditions and the following
15  #    disclaimer in the documentation and/or other materials provided
16  #    with the distribution.
17  #  * Neither the name of the author nor the names of its
18  #    contributors may be used to endorse or promote products derived
19  #    from this software without specific prior written permission.
20  #
21  # THIS SOFTWARE IS PROVIDED BY THE COPYRIGHT HOLDERS AND CONTRIBUTORS
22  # "AS IS" AND ANY EXPRESS OR IMPLIED WARRANTIES, INCLUDING, BUT NOT
23  # LIMITED TO, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS
24  # FOR A PARTICULAR PURPOSE ARE DISCLAIMED. IN NO EVENT SHALL THE
25  # COPYRIGHT OWNER OR CONTRIBUTORS BE LIABLE FOR ANY DIRECT, INDIRECT,
26  # INCIDENTAL, SPECIAL, EXEMPLARY, OR CONSEQUENTIAL DAMAGES (INCLUDING,
27  # BUT NOT LIMITED TO, PROCUREMENT OF SUBSTITUTE GOODS OR SERVICES;
28  # LOSS OF USE, DATA, OR PROFITS; OR BUSINESS INTERRUPTION) HOWEVER
29  # CAUSED AND ON ANY THEORY OF LIABILITY, WHETHER IN CONTRACT, STRICT
```

```python
30    # LIABILITY, OR TORT (INCLUDING NEGLIGENCE OR OTHERWISE) ARISING IN
31    # ANY WAY OUT OF THE USE OF THIS SOFTWARE, EVEN IF ADVISED OF THE
32    # POSSIBILITY OF SUCH DAMAGE.
33
34    import sys, math, Image
35
36    def Distance(p1,p2):
37      dx = p2[0] - p1[0]
38      dy = p2[1] - p1[1]
39      return math.sqrt(dx*dx+dy*dy)
40
41    def ScaleRotateTranslate(image, angle, center = None, new_center = None, scale = None, resample=Image.BICUBIC):
42      if (scale is None) and (center is None):
43        return image.rotate(angle=angle, resample=resample)
44      nx,ny = x,y = center
45      sx=sy=1.0
46      if new_center:
47        (nx,ny) = new_center
48      if scale:
49        (sx,sy) = (scale, scale)
50      cosine = math.cos(angle)
51      sine = math.sin(angle)
52      a = cosine/sx
53      b = sine/sx
54      c = x-nx*a-ny*b
55      d = -sine/sy
56      e = cosine/sy
57      f = y-nx*d-ny*e
58      return image.transform(image.size, Image.AFFINE, (a,b,c,d,e,f), resample=resample)
59
60    def CropFace(image, eye_left=(0,0), eye_right=(0,0), offset_pct=(0.2,0.2), dest_sz = (70,70)):
61      # calculate offsets in original image
62      offset_h = math.floor(float(offset_pct[0])*dest_sz[0])
63      offset_v = math.floor(float(offset_pct[1])*dest_sz[1])
64      # get the direction
65      eye_direction = (eye_right[0] - eye_left[0], eye_right[1] - eye_left[1])
66      # calc rotation angle in radians
67      rotation = -math.atan2(float(eye_direction[1]),float(eye_direction[0]))
68      # distance between them
69      dist = Distance(eye_left, eye_right)
70      # calculate the reference eye-width
71      reference = dest_sz[0] - 2.0*offset_h
72      # scale factor
73      scale = float(dist)/float(reference)
74      # rotate original around the left eye
75      image = ScaleRotateTranslate(image, center=eye_left, angle=rotation)
76      # crop the rotated image
77      crop_xy = (eye_left[0] - scale*offset_h, eye_left[1] - scale*offset_v)
78      crop_size = (dest_sz[0]*scale, dest_sz[1]*scale)
79      image = image.crop((int(crop_xy[0]), int(crop_xy[1]), int(crop_xy[0]+crop_size[0]), int(crop_xy[1]+crop_size[1])))
80      # resize it
81      image = image.resize(dest_sz, Image.ANTIALIAS)
82      return image
83
84    if __name__ == "__main__":
85      image =  Image.open("arnie.jpg")
86      CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.1,0.1), dest_sz=(200,200)).save("arnie_10_10_
87      CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2), dest_sz=(200,200)).save("arnie_20_20_
```

```
88    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.3,0.3), dest_sz=(200,200)).save("arnie_30_30_
89    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2)).save("arnie_20_20_70_70.jpg")
```

Imagine we are given this photo of Arnold Schwarzenegger, which is under a Public Domain license. The (x,y)-position of the eyes is approximately *(252,364)* for the left and *(420,366)* for the right eye. Now you only need to define the horizontal offset, vertical offset and the size your scaled, rotated & cropped face should have.

Here are some examples:

| Configuration | Cropped, Scaled, Rotated Face |
|---|---|
| 0.1 (10%), 0.1 (10%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (200,200) |  |
| 0.3 (30%), 0.3 (30%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (70,70) |  |

**CSV for the AT&T Facedatabase**

```
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2   /home/philipp/facerec/data/at/s13/7.pgm;12
3   /home/philipp/facerec/data/at/s13/6.pgm;12
4   /home/philipp/facerec/data/at/s13/9.pgm;12
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233  /home/philipp/facerec/data/at/s14/6.pgm;13
234  /home/philipp/facerec/data/at/s14/9.pgm;13
235  /home/philipp/facerec/data/at/s14/5.pgm;13
236  /home/philipp/facerec/data/at/s14/3.pgm;13
237  /home/philipp/facerec/data/at/s14/4.pgm;13
238  /home/philipp/facerec/data/at/s14/10.pgm;13
239  /home/philipp/facerec/data/at/s14/8.pgm;13
240  /home/philipp/facerec/data/at/s14/1.pgm;13
241  /home/philipp/facerec/data/at/s34/2.pgm;33
242  /home/philipp/facerec/data/at/s34/7.pgm;33
243  /home/philipp/facerec/data/at/s34/6.pgm;33
244  /home/philipp/facerec/data/at/s34/9.pgm;33
245  /home/philipp/facerec/data/at/s34/5.pgm;33
246  /home/philipp/facerec/data/at/s34/3.pgm;33
247  /home/philipp/facerec/data/at/s34/4.pgm;33
248  /home/philipp/facerec/data/at/s34/10.pgm;33
249  /home/philipp/facerec/data/at/s34/8.pgm;33
250  /home/philipp/facerec/data/at/s34/1.pgm;33
251  /home/philipp/facerec/data/at/s11/2.pgm;10
252  /home/philipp/facerec/data/at/s11/7.pgm;10
253  /home/philipp/facerec/data/at/s11/6.pgm;10
254  /home/philipp/facerec/data/at/s11/9.pgm;10
255  /home/philipp/facerec/data/at/s11/5.pgm;10
256  /home/philipp/facerec/data/at/s11/3.pgm;10
257  /home/philipp/facerec/data/at/s11/4.pgm;10
258  /home/philipp/facerec/data/at/s11/10.pgm;10
259  /home/philipp/facerec/data/at/s11/8.pgm;10
260  /home/philipp/facerec/data/at/s11/1.pgm;10
261  /home/philipp/facerec/data/at/s26/2.pgm;25
262  /home/philipp/facerec/data/at/s26/7.pgm;25
263  /home/philipp/facerec/data/at/s26/6.pgm;25
264  /home/philipp/facerec/data/at/s26/9.pgm;25
265  /home/philipp/facerec/data/at/s26/5.pgm;25
266  /home/philipp/facerec/data/at/s26/3.pgm;25
267  /home/philipp/facerec/data/at/s26/4.pgm;25
268  /home/philipp/facerec/data/at/s26/10.pgm;25
269  /home/philipp/facerec/data/at/s26/8.pgm;25
270  /home/philipp/facerec/data/at/s26/1.pgm;25
271  /home/philipp/facerec/data/at/s18/2.pgm;17
272  /home/philipp/facerec/data/at/s18/7.pgm;17
273  /home/philipp/facerec/data/at/s18/6.pgm;17
274  /home/philipp/facerec/data/at/s18/9.pgm;17
275  /home/philipp/facerec/data/at/s18/5.pgm;17
276  /home/philipp/facerec/data/at/s18/3.pgm;17
277  /home/philipp/facerec/data/at/s18/4.pgm;17
278  /home/philipp/facerec/data/at/s18/10.pgm;17
279  /home/philipp/facerec/data/at/s18/8.pgm;17
280  /home/philipp/facerec/data/at/s18/1.pgm;17
281  /home/philipp/facerec/data/at/s29/2.pgm;28
282  /home/philipp/facerec/data/at/s29/7.pgm;28
283  /home/philipp/facerec/data/at/s29/6.pgm;28
284  /home/philipp/facerec/data/at/s29/9.pgm;28
285  /home/philipp/facerec/data/at/s29/5.pgm;28
286  /home/philipp/facerec/data/at/s29/3.pgm;28
287  /home/philipp/facerec/data/at/s29/4.pgm;28
288  /home/philipp/facerec/data/at/s29/10.pgm;28
289  /home/philipp/facerec/data/at/s29/8.pgm;28
```

```
290  /home/philipp/facerec/data/at/s29/1.pgm;28
291  /home/philipp/facerec/data/at/s33/2.pgm;32
292  /home/philipp/facerec/data/at/s33/7.pgm;32
293  /home/philipp/facerec/data/at/s33/6.pgm;32
294  /home/philipp/facerec/data/at/s33/9.pgm;32
295  /home/philipp/facerec/data/at/s33/5.pgm;32
296  /home/philipp/facerec/data/at/s33/3.pgm;32
297  /home/philipp/facerec/data/at/s33/4.pgm;32
298  /home/philipp/facerec/data/at/s33/10.pgm;32
299  /home/philipp/facerec/data/at/s33/8.pgm;32
300  /home/philipp/facerec/data/at/s33/1.pgm;32
301  /home/philipp/facerec/data/at/s12/2.pgm;11
302  /home/philipp/facerec/data/at/s12/7.pgm;11
303  /home/philipp/facerec/data/at/s12/6.pgm;11
304  /home/philipp/facerec/data/at/s12/9.pgm;11
305  /home/philipp/facerec/data/at/s12/5.pgm;11
306  /home/philipp/facerec/data/at/s12/3.pgm;11
307  /home/philipp/facerec/data/at/s12/4.pgm;11
308  /home/philipp/facerec/data/at/s12/10.pgm;11
309  /home/philipp/facerec/data/at/s12/8.pgm;11
310  /home/philipp/facerec/data/at/s12/1.pgm;11
311  /home/philipp/facerec/data/at/s6/2.pgm;5
312  /home/philipp/facerec/data/at/s6/7.pgm;5
313  /home/philipp/facerec/data/at/s6/6.pgm;5
314  /home/philipp/facerec/data/at/s6/9.pgm;5
315  /home/philipp/facerec/data/at/s6/5.pgm;5
316  /home/philipp/facerec/data/at/s6/3.pgm;5
317  /home/philipp/facerec/data/at/s6/4.pgm;5
318  /home/philipp/facerec/data/at/s6/10.pgm;5
319  /home/philipp/facerec/data/at/s6/8.pgm;5
320  /home/philipp/facerec/data/at/s6/1.pgm;5
321  /home/philipp/facerec/data/at/s22/2.pgm;21
322  /home/philipp/facerec/data/at/s22/7.pgm;21
323  /home/philipp/facerec/data/at/s22/6.pgm;21
324  /home/philipp/facerec/data/at/s22/9.pgm;21
325  /home/philipp/facerec/data/at/s22/5.pgm;21
326  /home/philipp/facerec/data/at/s22/3.pgm;21
327  /home/philipp/facerec/data/at/s22/4.pgm;21
328  /home/philipp/facerec/data/at/s22/10.pgm;21
329  /home/philipp/facerec/data/at/s22/8.pgm;21
330  /home/philipp/facerec/data/at/s22/1.pgm;21
331  /home/philipp/facerec/data/at/s15/2.pgm;14
332  /home/philipp/facerec/data/at/s15/7.pgm;14
333  /home/philipp/facerec/data/at/s15/6.pgm;14
334  /home/philipp/facerec/data/at/s15/9.pgm;14
335  /home/philipp/facerec/data/at/s15/5.pgm;14
336  /home/philipp/facerec/data/at/s15/3.pgm;14
337  /home/philipp/facerec/data/at/s15/4.pgm;14
338  /home/philipp/facerec/data/at/s15/10.pgm;14
339  /home/philipp/facerec/data/at/s15/8.pgm;14
340  /home/philipp/facerec/data/at/s15/1.pgm;14
341  /home/philipp/facerec/data/at/s2/2.pgm;1
342  /home/philipp/facerec/data/at/s2/7.pgm;1
343  /home/philipp/facerec/data/at/s2/6.pgm;1
344  /home/philipp/facerec/data/at/s2/9.pgm;1
345  /home/philipp/facerec/data/at/s2/5.pgm;1
346  /home/philipp/facerec/data/at/s2/3.pgm;1
347  /home/philipp/facerec/data/at/s2/4.pgm;1
```

```
348   /home/philipp/facerec/data/at/s2/10.pgm;1
349   /home/philipp/facerec/data/at/s2/8.pgm;1
350   /home/philipp/facerec/data/at/s2/1.pgm;1
351   /home/philipp/facerec/data/at/s31/2.pgm;30
352   /home/philipp/facerec/data/at/s31/7.pgm;30
353   /home/philipp/facerec/data/at/s31/6.pgm;30
354   /home/philipp/facerec/data/at/s31/9.pgm;30
355   /home/philipp/facerec/data/at/s31/5.pgm;30
356   /home/philipp/facerec/data/at/s31/3.pgm;30
357   /home/philipp/facerec/data/at/s31/4.pgm;30
358   /home/philipp/facerec/data/at/s31/10.pgm;30
359   /home/philipp/facerec/data/at/s31/8.pgm;30
360   /home/philipp/facerec/data/at/s31/1.pgm;30
361   /home/philipp/facerec/data/at/s28/2.pgm;27
362   /home/philipp/facerec/data/at/s28/7.pgm;27
363   /home/philipp/facerec/data/at/s28/6.pgm;27
364   /home/philipp/facerec/data/at/s28/9.pgm;27
365   /home/philipp/facerec/data/at/s28/5.pgm;27
366   /home/philipp/facerec/data/at/s28/3.pgm;27
367   /home/philipp/facerec/data/at/s28/4.pgm;27
368   /home/philipp/facerec/data/at/s28/10.pgm;27
369   /home/philipp/facerec/data/at/s28/8.pgm;27
370   /home/philipp/facerec/data/at/s28/1.pgm;27
371   /home/philipp/facerec/data/at/s40/2.pgm;39
372   /home/philipp/facerec/data/at/s40/7.pgm;39
373   /home/philipp/facerec/data/at/s40/6.pgm;39
374   /home/philipp/facerec/data/at/s40/9.pgm;39
375   /home/philipp/facerec/data/at/s40/5.pgm;39
376   /home/philipp/facerec/data/at/s40/3.pgm;39
377   /home/philipp/facerec/data/at/s40/4.pgm;39
378   /home/philipp/facerec/data/at/s40/10.pgm;39
379   /home/philipp/facerec/data/at/s40/8.pgm;39
380   /home/philipp/facerec/data/at/s40/1.pgm;39
381   /home/philipp/facerec/data/at/s3/2.pgm;2
382   /home/philipp/facerec/data/at/s3/7.pgm;2
383   /home/philipp/facerec/data/at/s3/6.pgm;2
384   /home/philipp/facerec/data/at/s3/9.pgm;2
385   /home/philipp/facerec/data/at/s3/5.pgm;2
386   /home/philipp/facerec/data/at/s3/3.pgm;2
387   /home/philipp/facerec/data/at/s3/4.pgm;2
388   /home/philipp/facerec/data/at/s3/10.pgm;2
389   /home/philipp/facerec/data/at/s3/8.pgm;2
390   /home/philipp/facerec/data/at/s3/1.pgm;2
391   /home/philipp/facerec/data/at/s38/2.pgm;37
392   /home/philipp/facerec/data/at/s38/7.pgm;37
393   /home/philipp/facerec/data/at/s38/6.pgm;37
394   /home/philipp/facerec/data/at/s38/9.pgm;37
395   /home/philipp/facerec/data/at/s38/5.pgm;37
396   /home/philipp/facerec/data/at/s38/3.pgm;37
397   /home/philipp/facerec/data/at/s38/4.pgm;37
398   /home/philipp/facerec/data/at/s38/10.pgm;37
399   /home/philipp/facerec/data/at/s38/8.pgm;37
400   /home/philipp/facerec/data/at/s38/1.pgm;37
```

## Gender Classification with OpenCV

## Introduction

A lot of people interested in face recognition, also want to know how to perform image classification tasks like:

- Gender Classification (Gender Detection)

- Emotion Classification (Emotion Detection)

- Glasses Classification (Glasses Detection)

- ...

This is has become very, very easy with the new `FaceRecognizer` class. In this tutorial I'll show you how to perform gender classification with OpenCV on a set of face images. You'll also learn how to align your images to enhance the recognition results. If you want to do emotion classification instead of gender classification, all you need to do is to update is your training data and the configuration you pass to the demo.

## Prerequisites

For gender classification of faces, you'll need some images of male and female faces first. I've decided to search faces of celebrities using Google Images with the faces filter turned on (my god, they have great algorithms at Google!). My database has 8 male and 5 female subjects, each with 10 images. Here are the names, if you don't know who to search:

- Angelina Jolie

- Arnold Schwarzenegger

- Brad Pitt

- Emma Watson

- George Clooney

- Jennifer Lopez

- Johnny Depp

- Justin Timberlake

- Katy Perry

- Keanu Reeves

- Naomi Watts

- Patrick Stewart

- Tom Cruise

Once you have acquired some images, you'll need to read them. In the demo application I have decided to read the images from a very simple CSV file. Why? Because it's the simplest platform-independent approach I can think of. However, if you know a simpler solution please ping me about it. Basically all the CSV file needs to contain are lines composed of a `filename` followed by a `;` followed by the `label` (as *integer number*), making up a line like this:

```
/path/to/image.ext;0
```

Let's dissect the line. `/path/to/image.ext` is the path to an image, probably something like this if you are in Windows: `C:/faces/person0/image0.jpg`. Then there is the separator `;` and finally we assign a label `0` to the image. Think of the label as the subject (the person, the gender or whatever comes to your mind). In the gender classification scenario, the label is the gender the person has. I'll give the label `0` to *male* persons and the label `1` is for *female* subjects. So my CSV file looks like this:

```
/home/philipp/facerec/data/gender/male/keanu_reeves/keanu_reeves_01.jpg;0
/home/philipp/facerec/data/gender/male/keanu_reeves/keanu_reeves_02.jpg;0
/home/philipp/facerec/data/gender/male/keanu_reeves/keanu_reeves_03.jpg;0
...
/home/philipp/facerec/data/gender/female/katy_perry/katy_perry_01.jpg;1
/home/philipp/facerec/data/gender/female/katy_perry/katy_perry_02.jpg;1
/home/philipp/facerec/data/gender/female/katy_perry/katy_perry_03.jpg;1
...
/home/philipp/facerec/data/gender/male/brad_pitt/brad_pitt_01.jpg;0
/home/philipp/facerec/data/gender/male/brad_pitt/brad_pitt_02.jpg;0
/home/philipp/facerec/data/gender/male/brad_pitt/brad_pitt_03.jpg;0
...
/home/philipp/facerec/data/gender/female/emma_watson/emma_watson_08.jpg;1
/home/philipp/facerec/data/gender/female/emma_watson/emma_watson_02.jpg;1
/home/philipp/facerec/data/gender/female/emma_watson/emma_watson_03.jpg;1
```

All images for this example were chosen to have a frontal face perspective. They have been cropped, scaled and rotated to be aligned at the eyes, just like this set of George Clooney images:



You really don't want to create the CSV file by hand. And you really don't want scale, rotate & translate the images manually. I have prepared you two Python scripts `create_csv.py` and `crop_face.py`, you can find them in the `src` folder coming with this documentation. You'll see how to use them in the *Appendix*.

### Fisherfaces for Gender Classification

If you want to decide wether a person is *male* or *female*, you have to learn the discriminative features of both classes. The Eigenfaces method is based on the Principal Component Analysis, which is an unsupervised statistical model and not suitable for this task. Please see the Face Recognition tutorial for insights into the algorithms. The Fisherfaces instead yields a class-specific linear projection, so it is much better suited for the gender classification task.

http://www.bytefish.de/blog/gender_classification shows the recognition rate of the Fisherfaces method for gender classification.

The Fisherfaces method achieves a 98% recognition rate in a subject-independent cross-validation. A subject-independent cross-validation means *images of the person under test are never used for learning the model*. And could you believe it: you can simply use the facerec_fisherfaces demo, that's inlcuded in OpenCV.

### Fisherfaces in OpenCV

The source code for this demo application is also available in the `src` folder coming with this documentation:

- `src/facerec_fisherfaces.cpp`

```cpp
/*
 * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
 * Released to public domain under terms of the BSD Simplified license.
 *
 * Redistribution and use in source and binary forms, with or without
 * modification, are permitted provided that the following conditions are met:
 *   * Redistributions of source code must retain the above copyright
 *     notice, this list of conditions and the following disclaimer.
 *   * Redistributions in binary form must reproduce the above copyright
 *     notice, this list of conditions and the following disclaimer in the
 *     documentation and/or other materials provided with the distribution.
 *   * Neither the name of the organization nor the names of its contributors
 *     may be used to endorse or promote products derived from this software
 *     without specific prior written permission.
 *
 *   See <http://www.opensource.org/licenses/bsd-license>
 */

#include "opencv2/core/core.hpp"
#include "opencv2/contrib/contrib.hpp"
#include "opencv2/highgui/highgui.hpp"

#include <iostream>
#include <fstream>
#include <sstream>

using namespace cv;
using namespace std;

static Mat norm_0_255(InputArray _src) {
    Mat src = _src.getMat();
    // Create and return normalized image:
    Mat dst;
    switch(src.channels()) {
    case 1:
        cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC1);
        break;
    case 3:
        cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC3);
        break;
    default:
        src.copyTo(dst);
        break;
    }
    return dst;
```

```
46    }
47
48    static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
49        std::ifstream file(filename.c_str(), ifstream::in);
50        if (!file) {
51            string error_message = "No valid input file was given, please check the given filename.";
52            CV_Error(CV_StsBadArg, error_message);
53        }
54        string line, path, classlabel;
55        while (getline(file, line)) {
56            stringstream liness(line);
57            getline(liness, path, separator);
58            getline(liness, classlabel);
59            if(!path.empty() && !classlabel.empty()) {
60                images.push_back(imread(path, 0));
61                labels.push_back(atoi(classlabel.c_str()));
62            }
63        }
64    }
65
66    int main(int argc, const char *argv[]) {
67        // Check for valid command line arguments, print usage
68        // if no arguments were given.
69        if (argc < 2) {
70            cout << "usage: " << argv[0] << " <csv.ext> <output_folder> " << endl;
71            exit(1);
72        }
73        string output_folder;
74        if (argc == 3) {
75            output_folder = string(argv[2]);
76        }
77        // Get the path to your CSV.
78        string fn_csv = string(argv[1]);
79        // These vectors hold the images and corresponding labels.
80        vector<Mat> images;
81        vector<int> labels;
82        // Read in the data. This can fail if no valid
83        // input filename is given.
84        try {
85            read_csv(fn_csv, images, labels);
86        } catch (cv::Exception& e) {
87            cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
88            // nothing more we can do
89            exit(1);
90        }
91        // Quit if there are not enough images for this demo.
92        if(images.size() <= 1) {
93            string error_message = "This demo needs at least 2 images to work. Please add more images to your data set!";
94            CV_Error(CV_StsError, error_message);
95        }
96        // Get the height from the first image. We'll need this
97        // later in code to reshape the images to their original
98        // size:
99        int height = images[0].rows;
100       // The following lines simply get the last images from
101       // your dataset and remove it from the vector. This is
102       // done, so that the training data (which we learn the
103       // cv::FaceRecognizer on) and the test data we test
```

```
104        // the model with, do not overlap.
105        Mat testSample = images[images.size() - 1];
106        int testLabel = labels[labels.size() - 1];
107        images.pop_back();
108        labels.pop_back();
109        // The following lines create an Fisherfaces model for
110        // face recognition and train it with the images and
111        // labels read from the given CSV file.
112        // If you just want to keep 10 Fisherfaces, then call
113        // the factory method like this:
114        //
115        //      cv::createFisherFaceRecognizer(10);
116        //
117        // However it is not useful to discard Fisherfaces! Please
118        // always try to use _all_ available Fisherfaces for
119        // classification.
120        //
121        // If you want to create a FaceRecognizer with a
122        // confidence threshold (e.g. 123.0) and use _all_
123        // Fisherfaces, then call it with:
124        //
125        //      cv::createFisherFaceRecognizer(0, 123.0);
126        //
127        Ptr<FaceRecognizer> model = createFisherFaceRecognizer();
128        model->train(images, labels);
129        // The following line predicts the label of a given
130        // test image:
131        int predictedLabel = model->predict(testSample);
132        //
133        // To get the confidence of a prediction call the model with:
134        //
135        //      int predictedLabel = -1;
136        //      double confidence = 0.0;
137        //      model->predict(testSample, predictedLabel, confidence);
138        //
139        string result_message = format("Predicted class = %d / Actual class = %d.", predictedLabel, testLabel);
140        cout << result_message << endl;
141        // Here is how to get the eigenvalues of this Eigenfaces model:
142        Mat eigenvalues = model->getMat("eigenvalues");
143        // And we can do the same to display the Eigenvectors (read Eigenfaces):
144        Mat W = model->getMat("eigenvectors");
145        // Get the sample mean from the training data
146        Mat mean = model->getMat("mean");
147        // Display or save:
148        if(argc == 2) {
149            imshow("mean", norm_0_255(mean.reshape(1, images[0].rows)));
150        } else {
151            imwrite(format("%s/mean.png", output_folder.c_str()), norm_0_255(mean.reshape(1, images[0].rows)));
152        }
153        // Display or save the first, at most 16 Fisherfaces:
154        for (int i = 0; i < min(16, W.cols); i++) {
155            string msg = format("Eigenvalue #%d = %.5f", i, eigenvalues.at<double>(i));
156            cout << msg << endl;
157            // get eigenvector #i
158            Mat ev = W.col(i).clone();
159            // Reshape to original size & normalize to [0...255] for imshow.
160            Mat grayscale = norm_0_255(ev.reshape(1, height));
161            // Show the image & apply a Bone colormap for better sensing.
```

```
162         Mat cgrayscale;
163         applyColorMap(grayscale, cgrayscale, COLORMAP_BONE);
164         // Display or save:
165         if(argc == 2) {
166             imshow(format("fisherface_%d", i), cgrayscale);
167         } else {
168             imwrite(format("%s/fisherface_%d.png", output_folder.c_str(), i), norm_0_255(cgrayscale));
169         }
170     }
171     // Display or save the image reconstruction at some predefined steps:
172     for(int num_component = 0; num_component < min(16, W.cols); num_component++) {
173         // Slice the Fisherface from the model:
174         Mat ev = W.col(num_component);
175         Mat projection = subspaceProject(ev, mean, images[0].reshape(1,1));
176         Mat reconstruction = subspaceReconstruct(ev, mean, projection);
177         // Normalize the result:
178         reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
179         // Display or save:
180         if(argc == 2) {
181             imshow(format("fisherface_reconstruction_%d", num_component), reconstruction);
182         } else {
183             imwrite(format("%s/fisherface_reconstruction_%d.png", output_folder.c_str(), num_component), reconstructio
184         }
185     }
186     // Display if we are not writing to an output folder:
187     if(argc == 2) {
188         waitKey(0);
189     }
190     return 0;
191 }
```

### Running the Demo

If you are in Windows, then simply start the demo by running (from command line):

```
facerec_fisherfaces.exe C:/path/to/your/csv.ext
```

If you are in Linux, then simply start the demo by running:

```
./facerec_fisherfaces /path/to/your/csv.ext
```

If you don't want to display the images, but save them, then pass the desired path to the demo. It works like this in Windows:
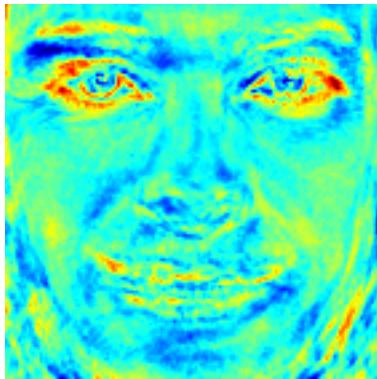
```
facerec_fisherfaces.exe C:/path/to/your/csv.ext C:/path/to/store/results/at
```

And in Linux:

```
./facerec_fisherfaces /path/to/your/csv.ext /path/to/store/results/at
```

### Results

If you run the program with your CSV file as parameter, you'll see the Fisherface that separates between male and female images. I've decided to apply a Jet colormap in this demo, so you can see which features the method identifies:

The demo also shows the average face of the male and female training images you have passed:



Moreover it the demo should yield the prediction for the correct gender:

```
Predicted class = 1 / Actual class = 1.
```

And for advanced users I have also shown the Eigenvalue for the Fisherface:

```
Eigenvalue #0 = 152.49493
```

And the Fisherfaces reconstruction:



I hope this gives you an idea how to approach gender classification and the other image classification tasks.

### Appendix

**Creating the CSV File**   You don't really want to create the CSV file by hand. I have prepared you a little Python script `create_csv.py` (you find it at `/src/create_csv.py` coming with this tutorial) that automatically creates you

a CSV file. If you have your images in hierarchie like this (`/basepath/<subject>/<image.ext>`):

```
philipp@mango:~/facerec/data/at$ tree
.
|-- s1
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
|-- s2
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
...
|-- s40
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
```

Then simply call `create_csv.py` with the path to the folder, just like this and you could save the output:

```
philipp@mango:~/facerec/data$ python create_csv.py
at/s13/2.pgm;0
at/s13/7.pgm;0
at/s13/6.pgm;0
at/s13/9.pgm;0
at/s13/5.pgm;0
at/s13/3.pgm;0
at/s13/4.pgm;0
at/s13/10.pgm;0
at/s13/8.pgm;0
at/s13/1.pgm;0
at/s17/2.pgm;1
at/s17/7.pgm;1
at/s17/6.pgm;1
at/s17/9.pgm;1
at/s17/5.pgm;1
at/s17/3.pgm;1
[...]
```

Here is the script, if you can't find it:

```python
1   import sys
2   import os.path
3
4   # This is a tiny script to help you creating a CSV file from a face
5   # database with a similar hierarchie:
6   #
7   #  philipp@mango:~/facerec/data/at$ tree
8   #  .
9   #  |-- README
10  #  |-- s1
11  #  |   |-- 1.pgm
12  #  |   |-- ...
13  #  |   |-- 10.pgm
14  #  |-- s2
15  #  |   |-- 1.pgm
16  #  |   |-- ...
17  #  |   |-- 10.pgm
18  #  ...
```

```
19   #   |-- s40
20   #   |   |-- 1.pgm
21   #   |   |-- ...
22   #   |   |-- 10.pgm
23   #

25   if __name__ == "__main__":

27       if len(sys.argv) != 2:
28           print "usage: create_csv <base_path>"
29           sys.exit(1)

31       BASE_PATH=sys.argv[1]
32       SEPARATOR=";"

34       label = 0
35       for dirname, dirnames, filenames in os.walk(BASE_PATH):
36           for subdirname in dirnames:
37               subject_path = os.path.join(dirname, subdirname)
38               for filename in os.listdir(subject_path):
39                   abs_path = "%s/%s" % (subject_path, filename)
40                   print "%s%s%d" % (abs_path, SEPARATOR, label)
41               label = label + 1
```

**Aligning Face Images** An accurate alignment of your image data is especially important in tasks like emotion detection, were you need as much detail as possible. Believe me... You don't want to do this by hand. So I've prepared you a tiny Python script. The code is really easy to use. To scale, rotate and crop the face image you just need to call *CropFace(image, eye_left, eye_right, offset_pct, dest_sz)*, where:

- *eye_left* is the position of the left eye
- *eye_right* is the position of the right eye
- *offset_pct* is the percent of the image you want to keep next to the eyes (horizontal, vertical direction)
- *dest_sz* is the size of the output image

If you are using the same *offset_pct* and *dest_sz* for your images, they are all aligned at the eyes.

```
1    #!/usr/bin/env python
2    # Software License Agreement (BSD License)
3    #
4    # Copyright (c) 2012, Philipp Wagner
5    # All rights reserved.
6    #
7    # Redistribution and use in source and binary forms, with or without
8    # modification, are permitted provided that the following conditions
9    # are met:
10   #
11   #  * Redistributions of source code must retain the above copyright
12   #    notice, this list of conditions and the following disclaimer.
13   #  * Redistributions in binary form must reproduce the above
14   #    copyright notice, this list of conditions and the following
15   #    disclaimer in the documentation and/or other materials provided
16   #    with the distribution.
17   #  * Neither the name of the author nor the names of its
18   #    contributors may be used to endorse or promote products derived
19   #    from this software without specific prior written permission.
20   #
```

```
21    # THIS SOFTWARE IS PROVIDED BY THE COPYRIGHT HOLDERS AND CONTRIBUTORS
22    # "AS IS" AND ANY EXPRESS OR IMPLIED WARRANTIES, INCLUDING, BUT NOT
23    # LIMITED TO, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS
24    # FOR A PARTICULAR PURPOSE ARE DISCLAIMED. IN NO EVENT SHALL THE
25    # COPYRIGHT OWNER OR CONTRIBUTORS BE LIABLE FOR ANY DIRECT, INDIRECT,
26    # INCIDENTAL, SPECIAL, EXEMPLARY, OR CONSEQUENTIAL DAMAGES (INCLUDING,
27    # BUT NOT LIMITED TO, PROCUREMENT OF SUBSTITUTE GOODS OR SERVICES;
28    # LOSS OF USE, DATA, OR PROFITS; OR BUSINESS INTERRUPTION) HOWEVER
29    # CAUSED AND ON ANY THEORY OF LIABILITY, WHETHER IN CONTRACT, STRICT
30    # LIABILITY, OR TORT (INCLUDING NEGLIGENCE OR OTHERWISE) ARISING IN
31    # ANY WAY OUT OF THE USE OF THIS SOFTWARE, EVEN IF ADVISED OF THE
32    # POSSIBILITY OF SUCH DAMAGE.
33
34    import sys, math, Image
35
36    def Distance(p1,p2):
37      dx = p2[0] - p1[0]
38      dy = p2[1] - p1[1]
39      return math.sqrt(dx*dx+dy*dy)
40
41    def ScaleRotateTranslate(image, angle, center = None, new_center = None, scale = None, resample=Image.BICUBIC):
42      if (scale is None) and (center is None):
43        return image.rotate(angle=angle, resample=resample)
44      nx,ny = x,y = center
45      sx=sy=1.0
46      if new_center:
47        (nx,ny) = new_center
48      if scale:
49        (sx,sy) = (scale, scale)
50      cosine = math.cos(angle)
51      sine = math.sin(angle)
52      a = cosine/sx
53      b = sine/sx
54      c = x-nx*a-ny*b
55      d = -sine/sy
56      e = cosine/sy
57      f = y-nx*d-ny*e
58      return image.transform(image.size, Image.AFFINE, (a,b,c,d,e,f), resample=resample)
59
60    def CropFace(image, eye_left=(0,0), eye_right=(0,0), offset_pct=(0.2,0.2), dest_sz = (70,70)):
61      # calculate offsets in original image
62      offset_h = math.floor(float(offset_pct[0])*dest_sz[0])
63      offset_v = math.floor(float(offset_pct[1])*dest_sz[1])
64      # get the direction
65      eye_direction = (eye_right[0] - eye_left[0], eye_right[1] - eye_left[1])
66      # calc rotation angle in radians
67      rotation = -math.atan2(float(eye_direction[1]),float(eye_direction[0]))
68      # distance between them
69      dist = Distance(eye_left, eye_right)
70      # calculate the reference eye-width
71      reference = dest_sz[0] - 2.0*offset_h
72      # scale factor
73      scale = float(dist)/float(reference)
74      # rotate original around the left eye
75      image = ScaleRotateTranslate(image, center=eye_left, angle=rotation)
76      # crop the rotated image
77      crop_xy = (eye_left[0] - scale*offset_h, eye_left[1] - scale*offset_v)
78      crop_size = (dest_sz[0]*scale, dest_sz[1]*scale)
```

```
79    image = image.crop((int(crop_xy[0]), int(crop_xy[1]), int(crop_xy[0]+crop_size[0]), int(crop_xy[1]+crop_size[1])))
80    # resize it
81    image = image.resize(dest_sz, Image.ANTIALIAS)
82    return image
83
84 if __name__ == "__main__":
85    image =  Image.open("arnie.jpg")
86    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.1,0.1), dest_sz=(200,200)).save("arnie_10_10_
87    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2), dest_sz=(200,200)).save("arnie_20_20_
88    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.3,0.3), dest_sz=(200,200)).save("arnie_30_30_
89    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2)).save("arnie_20_20_70_70.jpg")
```

Imagine we are given this photo of Arnold Schwarzenegger, which is under a Public Domain license. The (x,y)-position of the eyes is approximately *(252,364)* for the left and *(420,366)* for the right eye. Now you only need to define the horizontal offset, vertical offset and the size your scaled, rotated & cropped face should have.

Here are some examples:

| Configuration | Cropped, Scaled, Rotated Face |
|---|---|
| 0.1 (10%), 0.1 (10%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (200,200) |  |
| 0.3 (30%), 0.3 (30%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (70,70) |  |

## Face Recognition in Videos with OpenCV

**Table of Contents**

### Introduction

Whenever you hear the term *face recognition*, you instantly think of surveillance in videos. So performing face recognition in videos (e.g. webcam) is one of the most requested features I have got. I have heard your cries, so here it is. An application, that shows you how to do face recognition in videos! For the face detection part we'll use the awesome `CascadeClassifier` and we'll use `FaceRecognizer` for face recognition. This example uses the Fisherfaces method for face recognition, because it is robust against large changes in illumination.

Here is what the final application looks like. As you can see I am only writing the id of the recognized person above the detected face (by the way this id is Arnold Schwarzenegger for my data set):



This demo is a basis for your research and it shows you how to implement face recognition in videos. You probably want to extend the application and make it more sophisticated: You could combine the id with the name, then show the confidence of the prediction, recognize the emotion... and and and. But before you send mails, asking what these Haar-Cascade thing is or what a CSV is: Make sure you have read the entire tutorial. It's all explained in here. If you just want to scroll down to the code, please note:

- The available Haar-Cascades for face detection are located in the `data` folder of your OpenCV installation! One of the available Haar-Cascades for face detection is for example `/path/to/opencv/data/haarcascades/haarcascade_frontalface_default.xml`.

I encourage you to experiment with the application. Play around with the available `FaceRecognizer` implementations, try the available cascades in OpenCV and see if you can improve your results!

**Prerequisites**

You want to do face recognition, so you need some face images to learn a `FaceRecognizer` on. I have decided to reuse the images from the gender classification example: *Gender Classification with OpenCV*.

I have the following celebrities in my training data set:

- Angelina Jolie

- Arnold Schwarzenegger

- Brad Pitt

- George Clooney

- Johnny Depp

- Justin Timberlake

- Katy Perry

- Keanu Reeves

- Patrick Stewart

- Tom Cruise

In the demo I have decided to read the images from a very simple CSV file. Why? Because it's the simplest platform-independent approach I can think of. However, if you know a simpler solution please ping me about it. Basically all the CSV file needs to contain are lines composed of a `filename` followed by a `;` followed by the `label` (as *integer number*), making up a line like this:

```
/path/to/image.ext;0
```

Let's dissect the line. `/path/to/image.ext` is the path to an image, probably something like this if you are in Windows: `C:/faces/person0/image0.jpg`. Then there is the separator `;` and finally we assign a label `0` to the image. Think of the label as the subject (the person, the gender or whatever comes to your mind). In the face recognition scenario, the label is the person this image belongs to. In the gender classification scenario, the label is the gender the person has. So my CSV file looks like this:

```
/home/philipp/facerec/data/c/keanu_reeves/keanu_reeves_01.jpg;0
/home/philipp/facerec/data/c/keanu_reeves/keanu_reeves_02.jpg;0
/home/philipp/facerec/data/c/keanu_reeves/keanu_reeves_03.jpg;0
...
/home/philipp/facerec/data/c/katy_perry/katy_perry_01.jpg;1
/home/philipp/facerec/data/c/katy_perry/katy_perry_02.jpg;1
/home/philipp/facerec/data/c/katy_perry/katy_perry_03.jpg;1
...
/home/philipp/facerec/data/c/brad_pitt/brad_pitt_01.jpg;2
/home/philipp/facerec/data/c/brad_pitt/brad_pitt_02.jpg;2
/home/philipp/facerec/data/c/brad_pitt/brad_pitt_03.jpg;2
...
/home/philipp/facerec/data/c1/crop_arnold_schwarzenegger/crop_08.jpg;6
/home/philipp/facerec/data/c1/crop_arnold_schwarzenegger/crop_05.jpg;6
```

```
/home/philipp/facerec/data/c1/crop_arnold_schwarzenegger/crop_02.jpg;6
/home/philipp/facerec/data/c1/crop_arnold_schwarzenegger/crop_03.jpg;6
```

All images for this example were chosen to have a frontal face perspective. They have been cropped, scaled and rotated to be aligned at the eyes, just like this set of George Clooney images:



### Face Recongition from Videos

The source code for the demo is available in the `src` folder coming with this documentation:

- `src/facerec_video.cpp`

This demo uses the `CascadeClassifier`:

```cpp
/*
 * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
 * Released to public domain under terms of the BSD Simplified license.
 *
 * Redistribution and use in source and binary forms, with or without
 * modification, are permitted provided that the following conditions are met:
 *   * Redistributions of source code must retain the above copyright
 *     notice, this list of conditions and the following disclaimer.
 *   * Redistributions in binary form must reproduce the above copyright
 *     notice, this list of conditions and the following disclaimer in the
 *     documentation and/or other materials provided with the distribution.
 *   * Neither the name of the organization nor the names of its contributors
 *     may be used to endorse or promote products derived from this software
 *     without specific prior written permission.
 *
 *   See <http://www.opensource.org/licenses/bsd-license>
 */

#include "opencv2/core/core.hpp"
#include "opencv2/contrib/contrib.hpp"
#include "opencv2/highgui/highgui.hpp"
#include "opencv2/imgproc/imgproc.hpp"
#include "opencv2/objdetect/objdetect.hpp"

#include <iostream>
#include <fstream>
#include <sstream>

using namespace cv;
using namespace std;
```

```
31
32  static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
33      std::ifstream file(filename.c_str(), ifstream::in);
34      if (!file) {
35          string error_message = "No valid input file was given, please check the given filename.";
36          CV_Error(CV_StsBadArg, error_message);
37      }
38      string line, path, classlabel;
39      while (getline(file, line)) {
40          stringstream liness(line);
41          getline(liness, path, separator);
42          getline(liness, classlabel);
43          if(!path.empty() && !classlabel.empty()) {
44              images.push_back(imread(path, 0));
45              labels.push_back(atoi(classlabel.c_str()));
46          }
47      }
48  }
49
50  int main(int argc, const char *argv[]) {
51      // Check for valid command line arguments, print usage
52      // if no arguments were given.
53      if (argc != 4) {
54          cout << "usage: " << argv[0] << " </path/to/haar_cascade> </path/to/csv.ext> </path/to/device id>" << endl;
55          cout << "\t </path/to/haar_cascade> -- Path to the Haar Cascade for face detection." << endl;
56          cout << "\t </path/to/csv.ext> -- Path to the CSV file with the face database." << endl;
57          cout << "\t <device id> -- The webcam device id to grab frames from." << endl;
58          exit(1);
59      }
60      // Get the path to your CSV:
61      string fn_haar = string(argv[1]);
62      string fn_csv = string(argv[2]);
63      int deviceId = atoi(argv[3]);
64      // These vectors hold the images and corresponding labels:
65      vector<Mat> images;
66      vector<int> labels;
67      // Read in the data (fails if no valid input filename is given, but you'll get an error message):
68      try {
69          read_csv(fn_csv, images, labels);
70      } catch (cv::Exception& e) {
71          cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
72          // nothing more we can do
73          exit(1);
74      }
75      // Get the height from the first image. We'll need this
76      // later in code to reshape the images to their original
77      // size AND we need to reshape incoming faces to this size:
78      int im_width = images[0].cols;
79      int im_height = images[0].rows;
80      // Create a FaceRecognizer and train it on the given images:
81      Ptr<FaceRecognizer> model = createFisherFaceRecognizer();
82      model->train(images, labels);
83      // That's it for learning the Face Recognition model. You now
84      // need to create the classifier for the task of Face Detection.
85      // We are going to use the haar cascade you have specified in the
86      // command line arguments:
87      //
88      CascadeClassifier haar_cascade;
```

```
89      haar_cascade.load(fn_haar);
90      // Get a handle to the Video device:
91      VideoCapture cap(deviceId);
92      // Check if we can use this device at all:
93      if(!cap.isOpened()) {
94          cerr << "Capture Device ID " << deviceId << "cannot be opened." << endl;
95          return -1;
96      }
97      // Holds the current frame from the Video device:
98      Mat frame;
99      for(;;) {
100         cap >> frame;
101         // Clone the current frame:
102         Mat original = frame.clone();
103         // Convert the current frame to grayscale:
104         Mat gray;
105         cvtColor(original, gray, CV_BGR2GRAY);
106         // Find the faces in the frame:
107         vector< Rect_<int> > faces;
108         haar_cascade.detectMultiScale(gray, faces);
109         // At this point you have the position of the faces in
110         // faces. Now we'll get the faces, make a prediction and
111         // annotate it in the video. Cool or what?
112         for(int i = 0; i < faces.size(); i++) {
113             // Process face by face:
114             Rect face_i = faces[i];
115             // Crop the face from the image. So simple with OpenCV C++:
116             Mat face = gray(face_i);
117             // Resizing the face is necessary for Eigenfaces and Fisherfaces. You can easily
118             // verify this, by reading through the face recognition tutorial coming with OpenCV.
119             // Resizing IS NOT NEEDED for Local Binary Patterns Histograms, so preparing the
120             // input data really depends on the algorithm used.
121             //
122             // I strongly encourage you to play around with the algorithms. See which work best
123             // in your scenario, LBPH should always be a contender for robust face recognition.
124             //
125             // Since I am showing the Fisherfaces algorithm here, I also show how to resize the
126             // face you have just found:
127             Mat face_resized;
128             cv::resize(face, face_resized, Size(im_width, im_height), 1.0, 1.0, INTER_CUBIC);
129             // Now perform the prediction, see how easy that is:
130             int prediction = model->predict(face_resized);
131             // And finally write all we've found out to the original image!
132             // First of all draw a green rectangle around the detected face:
133             rectangle(original, face_i, CV_RGB(0, 255,0), 1);
134             // Create the text we will annotate the box with:
135             string box_text = format("Prediction = %d", prediction);
136             // Calculate the position for annotated text (make sure we don't
137             // put illegal values in there):
138             int pos_x = std::max(face_i.tl().x - 10, 0);
139             int pos_y = std::max(face_i.tl().y - 10, 0);
140             // And now put it into the image:
141             putText(original, box_text, Point(pos_x, pos_y), FONT_HERSHEY_PLAIN, 1.0, CV_RGB(0,255,0), 2.0);
142         }
143         // Show the result:
144         imshow("face_recognizer", original);
145         // And display it:
146         char key = (char) waitKey(20);
```

```
147          // Exit this loop on escape:
148          if(key == 27)
149              break;
150      }
151      return 0;
152 }
```

**Running the Demo**

You'll need:

- The path to a valid Haar-Cascade for detecting a face with a `CascadeClassifier`.

- The path to a valid CSV File for learning a `FaceRecognizer`.

- A webcam and its device id (you don't know the device id? Simply start from 0 on and see what happens).

If you are in Windows, then simply start the demo by running (from command line):

```
facerec_video.exe <C:/path/to/your/haar_cascade.xml> <C:/path/to/your/csv.ext> <video device>
```

If you are in Linux, then simply start the demo by running:

```
./facerec_video </path/to/your/haar_cascade.xml> </path/to/your/csv.ext> <video device>
```

An example. If the haar-cascade is at `C:/opencv/data/haarcascades/haarcascade_frontalface_default.xml`, the CSV file is at `C:/facerec/data/celebrities.txt` and I have a webcam with deviceId 1, then I would call the demo with:

```
facerec_video.exe C:/opencv/data/haarcascades/haarcascade_frontalface_default.xml C:/facerec/data/celebrities.txt 1
```

That's it.

**Results**

Enjoy!

**Appendix**

**Creating the CSV File** You don't really want to create the CSV file by hand. I have prepared you a little Python script `create_csv.py` (you find it at `/src/create_csv.py` coming with this tutorial) that automatically creates you a CSV file. If you have your images in hierarchie like this (`/basepath/<subject>/<image.ext>`):

```
philipp@mango:~/facerec/data/at$ tree
.
|-- s1
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
|-- s2
|   |-- 1.pgm
|   |-- ...
|   |-- 10.pgm
...
|-- s40
|   |-- 1.pgm
```

```
|    |-- ...
|    |-- 10.pgm
```

Then simply call `create_csv.py` with the path to the folder, just like this and you could save the output:

```
philipp@mango:~/facerec/data$ python create_csv.py
at/s13/2.pgm;0
at/s13/7.pgm;0
at/s13/6.pgm;0
at/s13/9.pgm;0
at/s13/5.pgm;0
at/s13/3.pgm;0
at/s13/4.pgm;0
at/s13/10.pgm;0
at/s13/8.pgm;0
at/s13/1.pgm;0
at/s17/2.pgm;1
at/s17/7.pgm;1
at/s17/6.pgm;1
at/s17/9.pgm;1
at/s17/5.pgm;1
at/s17/3.pgm;1
[...]
```

Here is the script, if you can't find it:

```python
import sys
import os.path

# This is a tiny script to help you creating a CSV file from a face
# database with a similar hierarchie:
#
#  philipp@mango:~/facerec/data/at$ tree
#  .
#  |-- README
#  |-- s1
#  |    |-- 1.pgm
#  |    |-- ...
#  |    |-- 10.pgm
#  |-- s2
#  |    |-- 1.pgm
#  |    |-- ...
#  |    |-- 10.pgm
#  ...
#  |-- s40
#  |    |-- 1.pgm
#  |    |-- ...
#  |    |-- 10.pgm
#

if __name__ == "__main__":

    if len(sys.argv) != 2:
        print "usage: create_csv <base_path>"
        sys.exit(1)

    BASE_PATH=sys.argv[1]
    SEPARATOR=";"
```

```
34        label = 0
35        for dirname, dirnames, filenames in os.walk(BASE_PATH):
36            for subdirname in dirnames:
37                subject_path = os.path.join(dirname, subdirname)
38                for filename in os.listdir(subject_path):
39                    abs_path = "%s/%s" % (subject_path, filename)
40                    print "%s%s%d" % (abs_path, SEPARATOR, label)
41                label = label + 1
```

**Aligning Face Images** An accurate alignment of your image data is especially important in tasks like emotion detection, were you need as much detail as possible. Believe me... You don't want to do this by hand. So I've prepared you a tiny Python script. The code is really easy to use. To scale, rotate and crop the face image you just need to call *CropFace(image, eye_left, eye_right, offset_pct, dest_sz)*, where:

- *eye_left* is the position of the left eye
- *eye_right* is the position of the right eye
- *offset_pct* is the percent of the image you want to keep next to the eyes (horizontal, vertical direction)
- *dest_sz* is the size of the output image

If you are using the same *offset_pct* and *dest_sz* for your images, they are all aligned at the eyes.

```
1    #!/usr/bin/env python
2    # Software License Agreement (BSD License)
3    #
4    # Copyright (c) 2012, Philipp Wagner
5    # All rights reserved.
6    #
7    # Redistribution and use in source and binary forms, with or without
8    # modification, are permitted provided that the following conditions
9    # are met:
10   #
11   #  * Redistributions of source code must retain the above copyright
12   #    notice, this list of conditions and the following disclaimer.
13   #  * Redistributions in binary form must reproduce the above
14   #    copyright notice, this list of conditions and the following
15   #    disclaimer in the documentation and/or other materials provided
16   #    with the distribution.
17   #  * Neither the name of the author nor the names of its
18   #    contributors may be used to endorse or promote products derived
19   #    from this software without specific prior written permission.
20   #
21   # THIS SOFTWARE IS PROVIDED BY THE COPYRIGHT HOLDERS AND CONTRIBUTORS
22   # "AS IS" AND ANY EXPRESS OR IMPLIED WARRANTIES, INCLUDING, BUT NOT
23   # LIMITED TO, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS
24   # FOR A PARTICULAR PURPOSE ARE DISCLAIMED. IN NO EVENT SHALL THE
25   # COPYRIGHT OWNER OR CONTRIBUTORS BE LIABLE FOR ANY DIRECT, INDIRECT,
26   # INCIDENTAL, SPECIAL, EXEMPLARY, OR CONSEQUENTIAL DAMAGES (INCLUDING,
27   # BUT NOT LIMITED TO, PROCUREMENT OF SUBSTITUTE GOODS OR SERVICES;
28   # LOSS OF USE, DATA, OR PROFITS; OR BUSINESS INTERRUPTION) HOWEVER
29   # CAUSED AND ON ANY THEORY OF LIABILITY, WHETHER IN CONTRACT, STRICT
30   # LIABILITY, OR TORT (INCLUDING NEGLIGENCE OR OTHERWISE) ARISING IN
31   # ANY WAY OUT OF THE USE OF THIS SOFTWARE, EVEN IF ADVISED OF THE
32   # POSSIBILITY OF SUCH DAMAGE.
33
34   import sys, math, Image
35
```

```python
36  def Distance(p1,p2):
37    dx = p2[0] - p1[0]
38    dy = p2[1] - p1[1]
39    return math.sqrt(dx*dx+dy*dy)
40
41  def ScaleRotateTranslate(image, angle, center = None, new_center = None, scale = None, resample=Image.BICUBIC):
42    if (scale is None) and (center is None):
43      return image.rotate(angle=angle, resample=resample)
44    nx,ny = x,y = center
45    sx=sy=1.0
46    if new_center:
47      (nx,ny) = new_center
48    if scale:
49      (sx,sy) = (scale, scale)
50    cosine = math.cos(angle)
51    sine = math.sin(angle)
52    a = cosine/sx
53    b = sine/sx
54    c = x-nx*a-ny*b
55    d = -sine/sy
56    e = cosine/sy
57    f = y-nx*d-ny*e
58    return image.transform(image.size, Image.AFFINE, (a,b,c,d,e,f), resample=resample)
59
60  def CropFace(image, eye_left=(0,0), eye_right=(0,0), offset_pct=(0.2,0.2), dest_sz = (70,70)):
61    # calculate offsets in original image
62    offset_h = math.floor(float(offset_pct[0])*dest_sz[0])
63    offset_v = math.floor(float(offset_pct[1])*dest_sz[1])
64    # get the direction
65    eye_direction = (eye_right[0] - eye_left[0], eye_right[1] - eye_left[1])
66    # calc rotation angle in radians
67    rotation = -math.atan2(float(eye_direction[1]),float(eye_direction[0]))
68    # distance between them
69    dist = Distance(eye_left, eye_right)
70    # calculate the reference eye-width
71    reference = dest_sz[0] - 2.0*offset_h
72    # scale factor
73    scale = float(dist)/float(reference)
74    # rotate original around the left eye
75    image = ScaleRotateTranslate(image, center=eye_left, angle=rotation)
76    # crop the rotated image
77    crop_xy = (eye_left[0] - scale*offset_h, eye_left[1] - scale*offset_v)
78    crop_size = (dest_sz[0]*scale, dest_sz[1]*scale)
79    image = image.crop((int(crop_xy[0]), int(crop_xy[1]), int(crop_xy[0]+crop_size[0]), int(crop_xy[1]+crop_size[1])))
80    # resize it
81    image = image.resize(dest_sz, Image.ANTIALIAS)
82    return image
83
84  if __name__ == "__main__":
85    image =  Image.open("arnie.jpg")
86    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.1,0.1), dest_sz=(200,200)).save("arnie_10_10_
87    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2), dest_sz=(200,200)).save("arnie_20_20_
88    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.3,0.3), dest_sz=(200,200)).save("arnie_30_30_
89    CropFace(image, eye_left=(252,364), eye_right=(420,366), offset_pct=(0.2,0.2)).save("arnie_20_20_70_70.jpg")
```

Imagine we are given this photo of Arnold Schwarzenegger, which is under a Public Domain license. The (x,y)-position of the eyes is approximately *(252,364)* for the left and *(420,366)* for the right eye. Now you only need to define the horizontal offset, vertical offset and the size your scaled, rotated & cropped face should have.

Here are some examples:

| Configuration | Cropped, Scaled, Rotated Face |
|---|---|
| 0.1 (10%), 0.1 (10%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (200,200) |  |
| 0.3 (30%), 0.3 (30%), (200,200) |  |
| 0.2 (20%), 0.2 (20%), (70,70) |  |

### Saving and Loading a FaceRecognizer

#### Introduction

Saving and loading a `FaceRecognizer` is very important. Training a FaceRecognizer can be a very time-intense task, plus it's often impossible to ship the whole face database to the user of your product. The task of saving and loading a FaceRecognizer is easy with `FaceRecognizer`. You only have to call `FaceRecognizer::load()` for loading and `FaceRecognizer::save()` for saving a `FaceRecognizer`.

I'll adapt the Eigenfaces example from the *Face Recognition with OpenCV*: Imagine we want to learn the Eigenfaces of the AT&T Facedatabase, store the model to a YAML file and then load it again.

From the loaded model, we'll get a prediction, show the mean, Eigenfaces and the image reconstruction.

#### Using FaceRecognizer::save and FaceRecognizer::load

The source code for this demo application is also available in the `src` folder coming with this documentation:

- `src/facerec_save_load.cpp`

```
1   /*
2    * Copyright (c) 2011. Philipp Wagner <bytefish[at]gmx[dot]de>.
3    * Released to public domain under terms of the BSD Simplified license.
4    *
5    * Redistribution and use in source and binary forms, with or without
6    * modification, are permitted provided that the following conditions are met:
7    *   * Redistributions of source code must retain the above copyright
8    *     notice, this list of conditions and the following disclaimer.
9    *   * Redistributions in binary form must reproduce the above copyright
10   *     notice, this list of conditions and the following disclaimer in the
11   *     documentation and/or other materials provided with the distribution.
12   *   * Neither the name of the organization nor the names of its contributors
13   *     may be used to endorse or promote products derived from this software
14   *     without specific prior written permission.
15   *
16   *   See <http://www.opensource.org/licenses/bsd-license>
17   */
18
19  #include "opencv2/contrib/contrib.hpp"
20  #include "opencv2/core/core.hpp"
21  #include "opencv2/highgui/highgui.hpp"
22
23  #include <iostream>
24  #include <fstream>
25  #include <sstream>
26
27  using namespace cv;
28  using namespace std;
29
30  static Mat norm_0_255(InputArray _src) {
31      Mat src = _src.getMat();
32      // Create and return normalized image:
33      Mat dst;
34      switch(src.channels()) {
35      case 1:
36          cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC1);
37          break;
```

```
38      case 3:
39          cv::normalize(_src, dst, 0, 255, NORM_MINMAX, CV_8UC3);
40          break;
41      default:
42          src.copyTo(dst);
43          break;
44      }
45      return dst;
46  }
47
48  static void read_csv(const string& filename, vector<Mat>& images, vector<int>& labels, char separator = ';') {
49      std::ifstream file(filename.c_str(), ifstream::in);
50      if (!file) {
51          string error_message = "No valid input file was given, please check the given filename.";
52          CV_Error(CV_StsBadArg, error_message);
53      }
54      string line, path, classlabel;
55      while (getline(file, line)) {
56          stringstream liness(line);
57          getline(liness, path, separator);
58          getline(liness, classlabel);
59          if(!path.empty() && !classlabel.empty()) {
60              images.push_back(imread(path, 0));
61              labels.push_back(atoi(classlabel.c_str()));
62          }
63      }
64  }
65
66  int main(int argc, const char *argv[]) {
67      // Check for valid command line arguments, print usage
68      // if no arguments were given.
69      if (argc < 2) {
70          cout << "usage: " << argv[0] << " <csv.ext> <output_folder> " << endl;
71          exit(1);
72      }
73      string output_folder;
74      if (argc == 3) {
75          output_folder = string(argv[2]);
76      }
77      // Get the path to your CSV.
78      string fn_csv = string(argv[1]);
79      // These vectors hold the images and corresponding labels.
80      vector<Mat> images;
81      vector<int> labels;
82      // Read in the data. This can fail if no valid
83      // input filename is given.
84      try {
85          read_csv(fn_csv, images, labels);
86      } catch (cv::Exception& e) {
87          cerr << "Error opening file \"" << fn_csv << "\". Reason: " << e.msg << endl;
88          // nothing more we can do
89          exit(1);
90      }
91      // Quit if there are not enough images for this demo.
92      if(images.size() <= 1) {
93          string error_message = "This demo needs at least 2 images to work. Please add more images to your data set!";
94          CV_Error(CV_StsError, error_message);
95      }
```

```
96    // Get the height from the first image. We'll need this
97    // later in code to reshape the images to their original
98    // size:
99    int height = images[0].rows;
100   // The following lines simply get the last images from
101   // your dataset and remove it from the vector. This is
102   // done, so that the training data (which we learn the
103   // cv::FaceRecognizer on) and the test data we test
104   // the model with, do not overlap.
105   Mat testSample = images[images.size() - 1];
106   int testLabel = labels[labels.size() - 1];
107   images.pop_back();
108   labels.pop_back();
109   // The following lines create an Eigenfaces model for
110   // face recognition and train it with the images and
111   // labels read from the given CSV file.
112   // This here is a full PCA, if you just want to keep
113   // 10 principal components (read Eigenfaces), then call
114   // the factory method like this:
115   //
116   //      cv::createEigenFaceRecognizer(10);
117   //
118   // If you want to create a FaceRecognizer with a
119   // confidence threshold (e.g. 123.0), call it with:
120   //
121   //      cv::createEigenFaceRecognizer(10, 123.0);
122   //
123   // If you want to use _all_ Eigenfaces and have a threshold,
124   // then call the method like this:
125   //
126   //      cv::createEigenFaceRecognizer(0, 123.0);
127   //
128   Ptr<FaceRecognizer> model0 = createEigenFaceRecognizer();
129   model0->train(images, labels);
130   // save the model to eigenfaces_at.yaml
131   model0->save("eigenfaces_at.yml");
132   //
133   //
134   // Now create a new Eigenfaces Recognizer
135   //
136   Ptr<FaceRecognizer> model1 = createEigenFaceRecognizer();
137   model1->load("eigenfaces_at.yml");
138   // The following line predicts the label of a given
139   // test image:
140   int predictedLabel = model1->predict(testSample);
141   //
142   // To get the confidence of a prediction call the model with:
143   //
144   //      int predictedLabel = -1;
145   //      double confidence = 0.0;
146   //      model->predict(testSample, predictedLabel, confidence);
147   //
148   string result_message = format("Predicted class = %d / Actual class = %d.", predictedLabel, testLabel);
149   cout << result_message << endl;
150   // Here is how to get the eigenvalues of this Eigenfaces model:
151   Mat eigenvalues = model1->getMat("eigenvalues");
152   // And we can do the same to display the Eigenvectors (read Eigenfaces):
153   Mat W = model1->getMat("eigenvectors");
```

**15.2. FaceRecognizer - Face Recognition with OpenCV**

```
154        // Get the sample mean from the training data
155        Mat mean = model1->getMat("mean");
156        // Display or save:
157        if(argc == 2) {
158            imshow("mean", norm_0_255(mean.reshape(1, images[0].rows)));
159        } else {
160            imwrite(format("%s/mean.png", output_folder.c_str()), norm_0_255(mean.reshape(1, images[0].rows)));
161        }
162        // Display or save the Eigenfaces:
163        for (int i = 0; i < min(10, W.cols); i++) {
164            string msg = format("Eigenvalue #%d = %.5f", i, eigenvalues.at<double>(i));
165            cout << msg << endl;
166            // get eigenvector #i
167            Mat ev = W.col(i).clone();
168            // Reshape to original size & normalize to [0...255] for imshow.
169            Mat grayscale = norm_0_255(ev.reshape(1, height));
170            // Show the image & apply a Jet colormap for better sensing.
171            Mat cgrayscale;
172            applyColorMap(grayscale, cgrayscale, COLORMAP_JET);
173            // Display or save:
174            if(argc == 2) {
175                imshow(format("eigenface_%d", i), cgrayscale);
176            } else {
177                imwrite(format("%s/eigenface_%d.png", output_folder.c_str(), i), norm_0_255(cgrayscale));
178            }
179        }
180        // Display or save the image reconstruction at some predefined steps:
181        for(int num_components = 10; num_components < 300; num_components+=15) {
182            // slice the eigenvectors from the model
183            Mat evs = Mat(W, Range::all(), Range(0, num_components));
184            Mat projection = subspaceProject(evs, mean, images[0].reshape(1,1));
185            Mat reconstruction = subspaceReconstruct(evs, mean, projection);
186            // Normalize the result:
187            reconstruction = norm_0_255(reconstruction.reshape(1, images[0].rows));
188            // Display or save:
189            if(argc == 2) {
190                imshow(format("eigenface_reconstruction_%d", num_components), reconstruction);
191            } else {
192                imwrite(format("%s/eigenface_reconstruction_%d.png", output_folder.c_str(), num_components), reconstructio
193            }
194        }
195        // Display if we are not writing to an output folder:
196        if(argc == 2) {
197            waitKey(0);
198        }
199        return 0;
200    }
```

## Results

eigenfaces_at.yml then contains the model state, we'll simply look at the first 10 lines with head eigenfaces_at.yml:

```
philipp@mango:~/github/libfacerec-build$ head eigenfaces_at.yml
%YAML:1.0
num_components: 399
mean: !!opencv-matrix
```

```
rows: 1
cols: 10304
dt: d
data: [ 8.5558897243107765e+01, 8.5511278195488714e+01,
    8.5854636591478695e+01, 8.5796992481203006e+01,
    8.5952380952380949e+01, 8.6162907268170414e+01,
    8.6082706766917283e+01, 8.5776942355889716e+01,
```

And here is the Reconstruction, which is the same as the original:



## ColorMaps in OpenCV

### applyColorMap

Applies a GNU Octave/MATLAB equivalent colormap on a given image.

**C++:** void **applyColorMap**(InputArray **src**, OutputArray **dst**, int **colormap**)

>> **Parameters**

>>>> **src** – The source image, grayscale or colored does not matter.

>>>> **dst** – The result is the colormapped source image. Note: `Mat::create()` is called on dst.

>>>> **colormap** – The colormap to apply, see the list of available colormaps below.

Currently the following GNU Octave/MATLAB equivalent colormaps are implemented:

```
enum
{
    COLORMAP_AUTUMN = 0,
    COLORMAP_BONE = 1,
    COLORMAP_JET = 2,
    COLORMAP_WINTER = 3,
    COLORMAP_RAINBOW = 4,
    COLORMAP_OCEAN = 5,
    COLORMAP_SUMMER = 6,
    COLORMAP_SPRING = 7,
    COLORMAP_COOL = 8,
    COLORMAP_HSV = 9,
    COLORMAP_PINK = 10,
    COLORMAP_HOT = 11
}
```

**Description**

The human perception isn't built for observing fine changes in grayscale images. Human eyes are more sensitive to observing changes between colors, so you often need to recolor your grayscale images to get a clue about them. OpenCV now comes with various colormaps to enhance the visualization in your computer vision application.

In OpenCV 2.4 you only need `applyColorMap()` to apply a colormap on a given image. The following sample code reads the path to an image from command line, applies a Jet colormap on it and shows the result:

```cpp
#include <opencv2/contrib/contrib.hpp>
#include <opencv2/core/core.hpp>
#include <opencv2/highgui/highgui.hpp>

using namespace cv;

int main(int argc, const char *argv[]) {
    // Get the path to the image, if it was given
    // if no arguments were given.
    string filename;
    if (argc > 1) {
        filename = string(argv[1]);
    }
    // The following lines show how to apply a colormap on a given image
    // and show it with cv::imshow example with an image. An exception is
    // thrown if the path to the image is invalid.
    if(!filename.empty()) {
        Mat img0 = imread(filename);
        // Throw an exception, if the image can't be read:
        if(img0.empty()) {
            CV_Error(CV_StsBadArg, "Sample image is empty. Please adjust your path, so it points to a valid input imag
        }
```

```
    // Holds the colormap version of the image:
    Mat cm_img0;
    // Apply the colormap:
    applyColorMap(img0, cm_img0, COLORMAP_JET);
    // Show the result:
    imshow("cm_img0", cm_img0);
    waitKey(0);
  }

  return 0;
}
```

And here are the color scales for each of the available colormaps:

| Class | Scale |
|---|---|
| COLORMAP_AUTUMN | |
| COLORMAP_BONE | |
| COLORMAP_COOL | |
| COLORMAP_HOT | |
| COLORMAP_HSV | |
| COLORMAP_JET | |
| COLORMAP_OCEAN | |
| COLORMAP_PINK | |
| COLORMAP_RAINBOW | |
| COLORMAP_SPRING | |
| COLORMAP_SUMMER | |
| COLORMAP_WINTER | |

## Changelog

### Release 0.05

This library is now included in the official OpenCV distribution (from 2.4 on). The :ocv:class'FaceRecognizer' is now an `Algorithm`, which better fits into the overall OpenCV API.

To reduce the confusion on user side and minimize my work, libfacerec and OpenCV have been synchronized and are now based on the same interfaces and implementation.

The library now has an extensive documentation:

- The API is explained in detail and with a lot of code examples.
- The face recognition guide I had written for Python and GNU Octave/MATLAB has been adapted to the new OpenCV C++ `cv::FaceRecognizer`.
- A tutorial for gender classification with Fisherfaces.
- A tutorial for face recognition in videos (e.g. webcam).

#### Release highlights

- There are no single highlights to pick from, this release is a highlight itself.

### Release 0.04

This version is fully Windows-compatible and works with OpenCV 2.3.1. Several bugfixes, but none influenced the recognition rate.

#### Release highlights

- A whole lot of exceptions with meaningful error messages.
- A tutorial for Windows users: http://bytefish.de/blog/opencv_visual_studio_and_libfacerec

### Release 0.03

Reworked the library to provide separate implementations in cpp files, because it's the preferred way of contributing OpenCV libraries. This means the library is not header-only anymore. Slight API changes were done, please see the documentation for details.

#### Release highlights

- New Unit Tests (for LBP Histograms) make the library more robust.
- Added more documentation.

### Release 0.02

Reworked the library to provide separate implementations in cpp files, because it's the preferred way of contributing OpenCV libraries. This means the library is not header-only anymore. Slight API changes were done, please see the documentation for details.

---

**Release highlights**

- New Unit Tests (for LBP Histograms) make the library more robust.

- Added a documentation and changelog in reStructuredText.

### Release 0.01

Initial release as header-only library.

**Release highlights**

- Colormaps for OpenCV to enhance the visualization.

- Face Recognition algorithms implemented:

    - Eigenfaces [TP91]

    - Fisherfaces [BHK97]

    - Local Binary Patterns Histograms [AHP04]

- Added persistence facilities to store the models with a common API.

- Unit Tests (using gtest).

- Providing a CMakeLists.txt to enable easy cross-platform building.

## Indices and tables

- *genindex*

- *modindex*

- *search*

# 15.3 Retina : a Bio mimetic human retina model

## Retina

**class Retina**

Class which provides the main controls to the Gipsa/Listic labs human retina model. Spatio-temporal filtering modelling the two main retina information channels :

- foveal vision for detailled color vision : the parvocellular pathway).

- periphearal vision for sensitive transient signals detection (motion and events) : the magnocellular pathway.

The retina can be settled up with various parameters, by default, the retina cancels mean luminance and enforces all details of the visual scene. In order to use your own parameters, you can use at least one time the *write(std::string fs)* method which will write a proper XML file with all default parameters. Then, tweak it on your own and reload them at any time using method *setup(std::string fs)*. These methods update a *Retina::RetinaParameters* member structure that is described hereafter.

```
class Retina
{
public:
  // parameters setup instance
  struct RetinaParameters; // this class is detailled later

  // constructors
  Retina (Size inputSize);
  Retina (Size inputSize, const bool colorMode, RETINA_COLORSAMPLINGMETHOD colorSamplingMethod=RETINA_COLOR_BAYER, con

  // main method for input frame processing
  void run (const Mat &inputImage);

  // output buffers retreival methods
  // -> foveal color vision details channel with luminance and noise correction
  void getParvo (Mat &retinaOutput_parvo);
  void getParvo (std::valarray< float > &retinaOutput_parvo);
  const std::valarray< float > & getParvo () const;
  // -> peripheral monochrome motion and events (transient information) channel
  void getMagno (Mat &retinaOutput_magno);
  void getMagno (std::valarray< float > &retinaOutput_magno);
  const std::valarray< float > & getMagno () const;

  // reset retina buffers... equivalent to closing your eyes for some seconds
  void clearBuffers ();

  // retreive input and output buffers sizes
  Size inputSize ();
  Size outputSize ();

  // setup methods with specific parameters specification of global xml config file loading/write
  void setup (std::string retinaParameterFile="", const bool applyDefaultSetupOnFailure=true);
  void setup (FileStorage &fs, const bool applyDefaultSetupOnFailure=true);
  void setup (RetinaParameters newParameters);
  struct Retina::RetinaParameters getParameters ();
  const std::string printSetup ();
  virtual void write (std::string fs) const;
  virtual void write (FileStorage &fs) const;
  void setupOPLandIPLParvoChannel (const bool colorMode=true, const bool normaliseOutput=true, const float photorecept
  void setupIPLMagnoChannel (const bool normaliseOutput=true, const float parasolCells_beta=0, const float parasolCell
  void setColorSaturation (const bool saturateColors=true, const float colorSaturationValue=4.0);
  void activateMovingContoursProcessing (const bool activate);
  void activateContoursProcessing (const bool activate);
};
```

### Description

Class which allows the Gipsa (preliminary work) / Listic (code maintainer) labs retina model to be used. This class allows human retina spatio-temporal image processing to be applied on still images, images sequences and video sequences. Briefly, here are the main human retina model properties:

- spectral whithening (mid-frequency details enhancement)

- high frequency spatio-temporal noise reduction (temporal noise and high frequency spatial noise are minimized)

- low frequency luminance reduction (luminance range compression) : high luminance regions do not hide details in darker regions anymore

• local logarithmic luminance compression allows details to be enhanced even in low light conditions

Use : this model can be used basically for spatio-temporal video effects but also in the aim of :

• performing texture analysis with enhanced signal to noise ratio and enhanced details robust against input images luminance ranges (check out the parvocellular retina channel output, by using the provided **getParvo** methods)

• performing motion analysis also taking benefit of the previously cited properties (check out the magnocellular retina channel output, by using the provided **getMagno** methods)

For more information, refer to the following papers :

• Benoit A., Caplier A., Durette B., Herault, J., "Using Human Visual System Modeling For Bio-Inspired Low Level Image Processing", Elsevier, Computer Vision and Image Understanding 114 (2010), pp. 758-773. DOI <http://dx.doi.org/10.1016/j.cviu.2010.01.011>

• Please have a look at the reference work of Jeanny Herault that you can read in his book :

Vision: Images, Signals and Neural Networks: Models of Neural Processing in Visual Perception (Progress in Neural Processing),By: Jeanny Herault, ISBN: 9814273686. WAPI (Tower ID): 113266891.

This retina filter code includes the research contributions of phd/research collegues from which code has been redrawn by the author :

• take a look at the *retinacolor.hpp* module to discover Brice Chaix de Lavarene phD color mosaicing/demosaicing and his reference paper: B. Chaix de Lavarene, D. Alleysson, B. Durette, J. Herault (2007). "Efficient demosaicing through recursive filtering", IEEE International Conference on Image Processing ICIP 2007

• take a look at *imagelogpolprojection.hpp* to discover retina spatial log sampling which originates from Barthelemy Durette phd with Jeanny Herault. A Retina / V1 cortex projection is also proposed and originates from Jeanny's discussions. ====> more informations in the above cited Jeanny Heraults's book.

## Demos and experiments !

Take a look at the C++ examples provided with OpenCV :

• **samples/cpp/retinademo.cpp shows how to use the retina module for details enhancement (Parvo channel output) and tran**

Typical uses are (provided your OpenCV installation is situated in folder *OpenCVReleaseFolder*)

– image processing : **OpenCVReleaseFolder/bin/retinademo -image myPicture.jpg**

– video processing : **OpenCVReleaseFolder/bin/retinademo -video myMovie.avi**

– webcam processing: **OpenCVReleaseFolder/bin/retinademo -video**

**Note :** This demo generates the file *RetinaDefaultParameters.xml* which contains the default parameters of the retina. Then, rename this as *RetinaSpecificParameters.xml*, adjust the parameters the way you want and reload the program to check the effect.

• **samples/cpp/OpenEXRimages_HighDynamicRange_Retina_toneMapping.cpp** shows how to use the retina to perform High Dynamic Range (HDR) luminance compression

Then, take a HDR image using bracketing with your camera and generate an OpenEXR image and then process it using the demo.

Typical use, supposing that you have the OpenEXR image *memorial.exr* (present in the samples/cpp/ folder)

**OpenCVReleaseFolder/bin/OpenEXRimages_HighDynamicRange_Retina_toneMapping memorial.exr**

Note that some sliders are made available to allow you to play with luminance compression.

# Methods description

Here are detailed the main methods to control the retina model

## Retina::Retina

**C++:** Retina::**Retina**(Size **inputSize**)

**C++:** Retina::**Retina**(Size **inputSize**, const bool **colorMode**, RETINA_COLORSAMPLINGMETHOD **colorSamplingMethod**=RETINA_COLOR_BAYER, const bool **useRetinaLogSampling**=false, const double **reductionFactor**=1.0, const double **samplingStrenght**=10.0 )

Constructors

### Parameters

**inputSize** – the input frame size

**colorMode** – the chosen processing mode : with or without color processing

**colorSamplingMethod** – specifies which kind of color sampling will be used * RETINA_COLOR_RANDOM: each pixel position is either R, G or B in a random choice * RETINA_COLOR_DIAGONAL: color sampling is RGBRGBRGB..., line 2 BRGBRG-BRG..., line 3, GBRGBRGBR... * RETINA_COLOR_BAYER: standard bayer sampling

**useRetinaLogSampling** – activate retina log sampling, if true, the 2 following parameters can be used

**reductionFactor** – only usefull if param useRetinaLogSampling=true, specifies the reduction factor of the output frame (as the center (fovea) is high resolution and corners can be underscaled, then a reduction of the output is allowed without precision leak

**samplingStrenght** – only usefull if param useRetinaLogSampling=true, specifies the strenght of the log scale that is applied

## Retina::activateContoursProcessing

**C++:** void Retina::**activateContoursProcessing**(const bool **activate**)

Activate/desactivate the Parvocellular pathway processing (contours information extraction), by default, it is activated

### Parameters

**activate** – true if Parvocellular (contours information extraction) output should be activated, false if not... if activated, the Parvocellular output can be retrieved using the **getParvo** methods

## Retina::activateMovingContoursProcessing

**C++:** void Retina::**activateMovingContoursProcessing**(const bool **activate**)

Activate/desactivate the Magnocellular pathway processing (motion information extraction), by default, it is activated

### Parameters

**activate** – true if Magnocellular output should be activated, false if not... if activated, the Magnocellular output can be retrieved using the **getMagno** methods

### Retina::clearBuffers

**C++:** void Retina::**clearBuffers**()

Clears all retina buffers (equivalent to opening the eyes after a long period of eye close ;o) whatchout the temporal transition occuring just after this method call.

### Retina::getParvo

**C++:** void Retina::**getParvo**(Mat& **retinaOutput_parvo**)

**C++:** void Retina::**getParvo**(std::valarray<float>& **retinaOutput_parvo**)

**C++:** const std::valarray<float>& Retina::**getParvo**() const

Accessor of the details channel of the retina (models foveal vision)

> **Parameters**
>
>> **retinaOutput_parvo** – the output buffer (reallocated if necessary), format can be :
>>
>> – a Mat, this output is rescaled for standard 8bits image processing use in OpenCV
>>
>> – a 1D std::valarray Buffer (encoding is R1, R2, ... Rn), this output is the original retina filter model output, without any quantification or rescaling

### Retina::getMagno

**C++:** void Retina::**getMagno**(Mat& **retinaOutput_magno**)

**C++:** void Retina::**getMagno**(std::valarray<float>& **retinaOutput_magno**)

**C++:** const std::valarray<float>& Retina::**getMagno**() const

Accessor of the motion channel of the retina (models peripheral vision)

> **Parameters**
>
>> **retinaOutput_magno** – the output buffer (reallocated if necessary), format can be :
>>
>> – a Mat, this output is rescaled for standard 8bits image processing use in OpenCV
>>
>> – a 1D std::valarray Buffer (encoding is R1, R2, ... Rn), this output is the original retina filter model output, without any quantification or rescaling

### Retina::getParameters

**struct Retina::RetinaParameters Retina::getParameters()**

Retrieve the current parameters values in a *Retina::RetinaParameters* structure

> **Returns** the current parameters setup

### Retina::inputSize

**C++:** Size Retina::**inputSize**()

Retreive retina input buffer size

> **Returns** the retina input buffer size

---

### Retina::outputSize

**C++:** Size Retina::**outputSize**()

> Retreive retina output buffer size that can be different from the input if a spatial log transformation is applied

>> **Returns** the retina output buffer size

### Retina::printSetup

**C++:** const std::string Retina::**printSetup**()

> Outputs a string showing the used parameters setup

>> **Returns** a string which contains formatted parameters information

### Retina::run

**C++:** void Retina::**run**(const Mat& **inputImage**)

> Method which allows retina to be applied on an input image, after run, encapsulated retina module is ready to deliver its outputs using dedicated acccessors, see getParvo and getMagno methods

>> **Parameters**

>>> **inputImage** – the input Mat image to be processed, can be gray level or BGR coded in any format (from 8bit to 16bits)

### Retina::setColorSaturation

**C++:** void Retina::**setColorSaturation**(const bool **saturateColors**=true, const float **colorSaturationValue**=4.0 )

> Activate color saturation as the final step of the color demultiplexing process -> this saturation is a sigmoide function applied to each channel of the demultiplexed image.

>> **Parameters**

>>> **saturateColors** – boolean that activates color saturation (if true) or desactivate (if false)

>>> **colorSaturationValue** – the saturation factor : a simple factor applied on the chrominance buffers

### Retina::setup

**C++:** void Retina::**setup**(std::string **retinaParameterFile**="", const bool **applyDefaultSetupOnFailure**=true )

**C++:** void Retina::**setup**(FileStorage& **fs**, const bool **applyDefaultSetupOnFailure**=true )

**C++:** void Retina::**setup**(RetinaParameters **newParameters**)

> Try to open an XML retina parameters file to adjust current retina instance setup => if the xml file does not exist, then default setup is applied => warning, Exceptions are thrown if read XML file is not valid

>> **Parameters**

>>> **retinaParameterFile** – the parameters filename

>>> **applyDefaultSetupOnFailure** – set to true if an error must be thrown on error

>>> **fs** – the open Filestorage which contains retina parameters

>>> **newParameters** – a parameters structures updated with the new target configuration

### Retina::write

**C++:** void Retina::**write**(std::string **fs**) const

**C++:** void Retina::**write**(FileStorage& **fs**) const
    Write xml/yml formated parameters information

>       **Parameters**
>
>>              **fs** – the filename of the xml file that will be open and writen with formatted parameters
>>              information

### Retina::setupIPLMagnoChannel

**C++:** void Retina::**setupIPLMagnoChannel**(const bool **normaliseOutput**=true, const float **parasolCells_beta**=0, const float **parasolCells_tau**=0, const float **parasolCells_k**=7, const float **amacrinCellsTemporalCutFrequency**=1.2, const float **V0CompressionParameter**=0.95, const float **localAdaptintegration_tau**=0, const float **localAdaptintegration_k**=7 )
    Set parameters values for the Inner Plexiform Layer (IPL) magnocellular channel this channel processes signals output from OPL processing stage in peripheral vision, it allows motion information enhancement. It is decorrelated from the details channel. See reference papers for more details.

>       **Parameters**
>
>>              **normaliseOutput** – specifies if (true) output is rescaled between 0 and 255 of not (false)
>>
>>              **parasolCells_beta** – the low pass filter gain used for local contrast adaptation at the IPL
>>              level of the retina (for ganglion cells local adaptation), typical value is 0
>>
>>              **parasolCells_tau** – the low pass filter time constant used for local contrast adaptation at the
>>              IPL level of the retina (for ganglion cells local adaptation), unit is frame, typical value is 0
>>              (immediate response)
>>
>>              **parasolCells_k** – the low pass filter spatial constant used for local contrast adaptation at the
>>              IPL level of the retina (for ganglion cells local adaptation), unit is pixels, typical value is 5
>>
>>              **amacrinCellsTemporalCutFrequency** – the time constant of the first order high pass fiter
>>              of the magnocellular way (motion information channel), unit is frames, typical value is 1.2
>>
>>              **V0CompressionParameter** – the compression strengh of the ganglion cells local adaptation
>>              output, set a value between 0.6 and 1 for best results, a high value increases more the low
>>              value sensitivity... and the output saturates faster, recommended value: 0.95
>>
>>              **localAdaptintegration_tau** – specifies the temporal constant of the low pas filter involved
>>              in the computation of the local "motion mean" for the local adaptation computation
>>
>>              **localAdaptintegration_k** – specifies the spatial constant of the low pas filter involved in
>>              the computation of the local "motion mean" for the local adaptation computation

### Retina::setupOPLandIPLParvoChannel

**C++:** void Retina::**setupOPLandIPLParvoChannel**(const bool **colorMode**=true, const bool **normalise-Output**=true, const float **photoreceptorsLocal-AdaptationSensitivity**=0.7, const float **photore-ceptorsTemporalConstant**=0.5, const float **pho-toreceptorsSpatialConstant**=0.53, const float **horizontalCellsGain**=0, const float **HcellsTemporal-Constant**=1, const float **HcellsSpatialConstant**=7, const float **ganglionCellsSensitivity**=0.7 )

Setup the OPL and IPL parvo channels (see biologocal model) OPL is referred as Outer Plexiform Layer of the retina, it allows the spatio-temporal filtering which withens the spectrum and reduces spatio-temporal noise while attenuating global luminance (low frequency energy) IPL parvo is the OPL next processing stage, it refers to a part of the Inner Plexiform layer of the retina, it allows high contours sensitivity in foveal vision. See reference papers for more informations.

> **Parameters**
>
> > **colorMode** – specifies if (true) color is processed of not (false) to then processing gray level image
> >
> > **normaliseOutput** – specifies if (true) output is rescaled between 0 and 255 of not (false)
> >
> > **photoreceptorsLocalAdaptationSensitivity** – the photoreceptors sensitivity renage is 0-1 (more log compression effect when value increases)
> >
> > **photoreceptorsTemporalConstant** – the time constant of the first order low pass filter of the photoreceptors, use it to cut high temporal frequencies (noise or fast motion), unit is frames, typical value is 1 frame
> >
> > **photoreceptorsSpatialConstant** – the spatial constant of the first order low pass filter of the photoreceptors, use it to cut high spatial frequencies (noise or thick contours), unit is pixels, typical value is 1 pixel
> >
> > **horizontalCellsGain** – gain of the horizontal cells network, if 0, then the mean value of the output is zero, if the parameter is near 1, then, the luminance is not filtered and is still reachable at the output, typicall value is 0
> >
> > **HcellsTemporalConstant** – the time constant of the first order low pass filter of the horizontal cells, use it to cut low temporal frequencies (local luminance variations), unit is frames, typical value is 1 frame, as the photoreceptors
> >
> > **HcellsSpatialConstant** – the spatial constant of the first order low pass filter of the horizontal cells, use it to cut low spatial frequencies (local luminance), unit is pixels, typical value is 5 pixel, this value is also used for local contrast computing when computing the local contrast adaptation at the ganglion cells level (Inner Plexiform Layer parvocellular channel model)
> >
> > **ganglionCellsSensitivity** – the compression strengh of the ganglion cells local adaptation output, set a value between 0.6 and 1 for best results, a high value increases more the low value sensitivity... and the output saturates faster, recommended value: 0.7

## Retina::RetinaParameters

**struct** Retina::**RetinaParameters**

This structure merges all the parameters that can be adjusted threw the **Retina::setup**(), **Retina::setupOPLandIPLParvoChannel** and **Retina::setupIPLMagnoChannel** setup methods Parameters

structure for better clarity, check explenations on the comments of methods : setupOPLandIPLParvoChannel and setupIPLMagnoChannel.

```cpp
class RetinaParameters{
    struct OPLandIplParvoParameters{ // Outer Plexiform Layer (OPL) and Inner Plexiform Layer Parvocellular (IplParvo)
            OPLandIplParvoParameters():colorMode(true),
                normaliseOutput(true), // specifies if (true) output is rescaled between 0 and 255 of not (false)
                photoreceptorsLocalAdaptationSensitivity(0.7f), // the photoreceptors sensitivity renage is 0-1 (more lo
                photoreceptorsTemporalConstant(0.5f),// the time constant of the first order low pass filter of the phot
                photoreceptorsSpatialConstant(0.53f),// the spatial constant of the first order low pass filter of the p
                horizontalCellsGain(0.0f),//gain of the horizontal cells network, if 0, then the mean value of the outpu
                hcellsTemporalConstant(1.f),// the time constant of the first order low pass filter of the horizontal ce
                hcellsSpatialConstant(7.f),//the spatial constant of the first order low pass filter of the horizontal c
                ganglionCellsSensitivity(0.7f)//the compression strengh of the ganglion cells local adaptation output, s
                {};// default setup
            bool colorMode, normaliseOutput;
            float photoreceptorsLocalAdaptationSensitivity, photoreceptorsTemporalConstant, photoreceptorsSpatialConsta
    };
    struct IplMagnoParameters{ // Inner Plexiform Layer Magnocellular channel (IplMagno)
            IplMagnoParameters():
                normaliseOutput(true), //specifies if (true) output is rescaled between 0 and 255 of not (false)
                parasolCells_beta(0.f), // the low pass filter gain used for local contrast adaptation at the IPL level
                parasolCells_tau(0.f), //the low pass filter time constant used for local contrast adaptation at the IPL
                parasolCells_k(7.f), //the low pass filter spatial constant used for local contrast adaptation at the IP
                amacrinCellsTemporalCutFrequency(1.2f), //the time constant of the first order high pass fiter of the ma
                V0CompressionParameter(0.95f), the compression strengh of the ganglion cells local adaptation output, se
                localAdaptintegration_tau(0.f), // specifies the temporal constant of the low pas filter involved in the
                localAdaptintegration_k(7.f) // specifies the spatial constant of the low pas filter involved in the com
                {};// default setup
            bool normaliseOutput;
            float parasolCells_beta, parasolCells_tau, parasolCells_k, amacrinCellsTemporalCutFrequency, V0CompressionP
    };
     struct OPLandIplParvoParameters OPLandIplParvo;
     struct IplMagnoParameters IplMagno;
};
```

# LEGACY. DEPRECATED STUFF

## 16.1 Motion Analysis

### CalcOpticalFlowBM

Calculates the optical flow for two images by using the block matching method.

**C:** void **cvCalcOpticalFlowBM**(const CvArr* **prev**, const CvArr* **curr**, CvSize **block_size**, CvSize **shift_size**, CvSize **max_range**, int **use_previous**, CvArr* **velx**, CvArr* **vely**)

**Python:** cv.**CalcOpticalFlowBM**(prev, curr, blockSize, shiftSize, max_range, usePrevious, velx, vely) → None

> **Parameters**
>
>> **prev** – First image, 8-bit, single-channel
>>
>> **curr** – Second image, 8-bit, single-channel
>>
>> **block_size** – Size of basic blocks that are compared
>>
>> **shift_size** – Block coordinate increments
>>
>> **max_range** – Size of the scanned neighborhood in pixels around the block
>>
>> **use_previous** – Flag that specifies whether to use the input velocity as initial approximations or not.
>>
>> **velx** – Horizontal component of the optical flow of
>>
>> $$\left\lfloor \frac{\texttt{prev->width} - \texttt{block\_size.width}}{\texttt{shift\_size.width}} \right\rfloor \times \left\lfloor \frac{\texttt{prev->height} - \texttt{block\_size.height}}{\texttt{shift\_size.height}} \right\rfloor$$
>>
>> size, 32-bit floating-point, single-channel
>>
>> **vely** – Vertical component of the optical flow of the same size velx , 32-bit floating-point, single-channel

The function calculates the optical flow for overlapped blocks block_size.width x block_size.height pixels each, thus the velocity fields are smaller than the original images. For every block in prev the functions tries to find a similar block in curr in some neighborhood of the original block or shifted by (velx(x0,y0), vely(x0,y0)) block as has been calculated by previous function call (if use_previous=1)

## CalcOpticalFlowHS

Calculates the optical flow for two images using Horn-Schunck algorithm.

**C:** void **cvCalcOpticalFlowHS**(const CvArr* **prev**, const CvArr* **curr**, int **use_previous**, CvArr* **velx**, CvArr* **vely**, double **lambda**, CvTermCriteria **criteria**)

**Python:** cv.**CalcOpticalFlowHS**(prev, curr, usePrevious, velx, vely, lambda, criteria) → None

> **Parameters**

>> **prev** – First image, 8-bit, single-channel

>> **curr** – Second image, 8-bit, single-channel

>> **use_previous** – Flag that specifies whether to use the input velocity as initial approximations or not.

>> **velx** – Horizontal component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

>> **vely** – Vertical component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

>> **lambda** – Smoothness weight. The larger it is, the smoother optical flow map you get.

>> **criteria** – Criteria of termination of velocity computing

The function computes the flow for every pixel of the first input image using the Horn and Schunck algorithm [Horn81]. The function is obsolete. To track sparse features, use `calcOpticalFlowPyrLK()`. To track all the pixels, use `calcOpticalFlowFarneback()`.

## CalcOpticalFlowLK

Calculates the optical flow for two images using Lucas-Kanade algorithm.

**C:** void **cvCalcOpticalFlowLK**(const CvArr* **prev**, const CvArr* **curr**, CvSize **win_size**, CvArr* **velx**, CvArr* **vely**)

**Python:** cv.**CalcOpticalFlowLK**(prev, curr, winSize, velx, vely) → None

> **Parameters**

>> **prev** – First image, 8-bit, single-channel

>> **curr** – Second image, 8-bit, single-channel

>> **win_size** – Size of the averaging window used for grouping pixels

>> **velx** – Horizontal component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

>> **vely** – Vertical component of the optical flow of the same size as input images, 32-bit floating-point, single-channel

The function computes the flow for every pixel of the first input image using the Lucas and Kanade algorithm [Lucas81]. The function is obsolete. To track sparse features, use `calcOpticalFlowPyrLK()`. To track all the pixels, use `calcOpticalFlowFarneback()`.

## 16.2 Expectation Maximization

This section describes obsolete `C` interface of EM algorithm. Details of the algorithm and its `C++` interface can be found in the other section *Expectation Maximization*.

### CvEMParams

**struct `CvEMParams`**

Parameters of the EM algorithm. All parameters are public. You can initialize them by a constructor and then override some of them directly if you want.

### CvEMParams::CvEMParams

The constructors

**C++:** `CvEMParams::`**`CvEMParams`**`()`

**C++:** `CvEMParams::`**`CvEMParams`**`(`int **nclusters**, int **cov_mat_type**=EM::COV_MAT_DIAGONAL, int **start_step**=EM::START_AUTO_STEP, CvTermCriteria **term_crit**=cvTermCriteria(CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 100, FLT_EPSILON), const CvMat* **probs**=0, const CvMat* **weights**=0, const CvMat* **means**=0, const CvMat** **covs**=0 )

> **Parameters**
>
> > **nclusters** – The number of mixture components in the Gaussian mixture model. Some of EM implementation could determine the optimal number of mixtures within a specified value range, but that is not the case in ML yet.
> >
> > **cov_mat_type** – Constraint on covariance matrices which defines type of matrices. Possible values are:
> >
> > – **CvEM::COV_MAT_SPHERICAL** A scaled identity matrix $\mu_k * I$. There is the only parameter $\mu_k$ to be estimated for each matrix. The option may be used in special cases, when the constraint is relevant, or as a first step in the optimization (for example in case when the data is preprocessed with PCA). The results of such preliminary estimation may be passed again to the optimization procedure, this time with `cov_mat_type=CvEM::COV_MAT_DIAGONAL`.
> >
> > – **CvEM::COV_MAT_DIAGONAL** A diagonal matrix with positive diagonal elements. The number of free parameters is `d` for each matrix. This is most commonly used option yielding good estimation results.
> >
> > – **CvEM::COV_MAT_GENERIC** A symmetric positively defined matrix. The number of free parameters in each matrix is about $d^2/2$. It is not recommended to use this option, unless there is pretty accurate initial estimation of the parameters and/or a huge number of training samples.
> >
> > **start_step** – The start step of the EM algorithm:
> >
> > – **CvEM::START_E_STEP** Start with Expectation step. You need to provide means $a_k$ of mixture components to use this option. Optionally you can pass weights $\pi_k$ and covariance matrices $S_k$ of mixture components.
> >
> > – **CvEM::START_M_STEP** Start with Maximization step. You need to provide initial probabilities $p_{i,k}$ to use this option.

– **CvEM::START_AUTO_STEP** Start with Expectation step. You need not provide any parameters because they will be estimated by the kmeans algorithm.

**term_crit** – The termination criteria of the EM algorithm. The EM algorithm can be terminated by the number of iterations `term_crit.max_iter` (number of M-steps) or when relative change of likelihood logarithm is less than `term_crit.epsilon`.

**probs** – Initial probabilities $p_{i,k}$ of sample $i$ to belong to mixture component $k$. It is a floating-point matrix of $nsamples \times nclusters$ size. It is used and must be not NULL only when `start_step=CvEM::START_M_STEP`.

**weights** – Initial weights $\pi_k$ of mixture components. It is a floating-point vector with $nclusters$ elements. It is used (if not NULL) only when `start_step=CvEM::START_E_STEP`.

**means** – Initial means $a_k$ of mixture components. It is a floating-point matrix of $nclusters \times dims$ size. It is used used and must be not NULL only when `start_step=CvEM::START_E_STEP`.

**covs** – Initial covariance matrices $S_k$ of mixture components. Each of covariance matrices is a valid square floating-point matrix of $dims \times dims$ size. It is used (if not NULL) only when `start_step=CvEM::START_E_STEP`.

The default constructor represents a rough rule-of-the-thumb:

```
CvEMParams() : nclusters(10), cov_mat_type(1/*CvEM::COV_MAT_DIAGONAL*/),
    start_step(0/*CvEM::START_AUTO_STEP*/), probs(0), weights(0), means(0), covs(0)
{
    term_crit=cvTermCriteria( CV_TERMCRIT_ITER+CV_TERMCRIT_EPS, 100, FLT_EPSILON );
}
```

With another constructor it is possible to override a variety of parameters from a single number of mixtures (the only essential problem-dependent parameter) to initial values for the mixture parameters.

## CvEM

**class CvEM** : **public CvStatModel**
The class implements the EM algorithm as described in the beginning of the section *Expectation Maximization*.

## CvEM::train

Estimates the Gaussian mixture parameters from a sample set.

**C++:** bool CvEM::**train**(const Mat& **samples**, const Mat& **sampleIdx**=Mat(), CvEMParams **params**=CvEMParams(), Mat* **labels**=0 )

**C++:** bool CvEM::**train**(const CvMat* **samples**, const CvMat* **sampleIdx**=0, CvEMParams **params**=CvEMParams(), CvMat* **labels**=0 )

> Parameters

>> **samples** – Samples from which the Gaussian mixture model will be estimated.

>> **sampleIdx** – Mask of samples to use. All samples are used by default.

>> **params** – Parameters of the EM algorithm.

>> **labels** – The optional output "class label" for each sample: $labels_i = arg\ max_k(p_{i,k}), i = 1..N$ (indices of the most probable mixture component for each sample).

Unlike many of the ML models, EM is an unsupervised learning algorithm and it does not take responses (class labels or function values) as input. Instead, it computes the *Maximum Likelihood Estimate* of the Gaussian mixture parameters from an input sample set, stores all the parameters inside the structure: $p_{i,k}$ in probs, $a_k$ in means , $S_k$ in covs[k], $\pi_k$ in weights , and optionally computes the output "class label" for each sample: $labels_i = \arg \max_k(p_{i,k}), i = 1..N$ (indices of the most probable mixture component for each sample).

The trained model can be used further for prediction, just like any other classifier. The trained model is similar to the `CvNormalBayesClassifier`.

For an example of clustering random samples of the multi-Gaussian distribution using EM, see `em.cpp` sample in the OpenCV distribution.

## CvEM::predict

Returns a mixture component index of a sample.

**C++:** float CvEM::**predict**(const Mat& **sample**, Mat* **probs**=0 ) const

**C++:** float CvEM::**predict**(const CvMat* **sample**, CvMat* **probs**) const

> **Parameters**

>> **sample** – A sample for classification.

>> **probs** – If it is not null then the method will write posterior probabilities of each component given the sample data to this parameter.

## CvEM::getNClusters

Returns the number of mixture components $M$ in the Gaussian mixture model.

**C++:** int CvEM::**getNClusters**() const

**C++:** int CvEM::**get_nclusters**() const

## CvEM::getMeans

Returns mixture means $a_k$.

**C++:** Mat CvEM::**getMeans**() const

**C++:** const CvMat* CvEM::**get_means**() const

## CvEM::getCovs

Returns mixture covariance matrices $S_k$.

**C++:** void CvEM::**getCovs**(std::vector<cv::Mat>& **covs**) const

**C++:** const CvMat** CvEM::**get_covs**() const

## CvEM::getWeights

Returns mixture weights $\pi_k$.

**C++:** Mat CvEM::**getWeights**() const

**C++:** const CvMat* CvEM::`get_weights`() const

## CvEM::getProbs

Returns vectors of probabilities for each training sample.

**C++:** Mat CvEM::`getProbs`() const

**C++:** const CvMat* CvEM::`get_probs`() const

For each training sample i (that have been passed to the constructor or to `CvEM::train()`) returns probabilities $p_{i,k}$ to belong to a mixture component k.

## CvEM::getLikelihood

Returns logarithm of likelihood.

**C++:** double CvEM::`getLikelihood`() const

**C++:** double CvEM::`get_log_likelihood`() const

## CvEM::write

Writes the trained Gaussian mixture model to the file storage.

**C++:** void CvEM::`write`(CvFileStorage* **fs**, const char* **name**) const

>    **Parameters**

>    >    **fs** – A file storage where the model will be written.

>    >    **name** – A name of the file node where the model data will be written.

## CvEM::read

Reads the trained Gaussian mixture model from the file storage.

**C++:** void CvEM::`read`(CvFileStorage* **fs**, CvFileNode* **node**)

>    **Parameters**

>    >    **fs** – A file storage with the trained model.

>    >    **node** – The parent map. If it is NULL, the function searches a node with parameters in all the top-level nodes (streams), starting with the first one.

# 16.3 Histograms

## CalcPGH

Calculates a pair-wise geometrical histogram for a contour.

**C:** void **cvCalcPGH**(const CvSeq* **contour**, CvHistogram* **hist**)

>    **Parameters**

>    >    **contour** – Input contour. Currently, only integer point coordinates are allowed.

**hist** – Calculated histogram. It must be two-dimensional.

The function calculates a 2D pair-wise geometrical histogram (PGH), described in [Iivarinen97] for the contour. The algorithm considers every pair of contour edges. The angle between the edges and the minimum/maximum distances are determined for every pair. To do this, each of the edges in turn is taken as the base, while the function loops through all the other edges. When the base edge and any other edge are considered, the minimum and maximum distances from the points on the non-base edge and line of the base edge are selected. The angle between the edges defines the row of the histogram in which all the bins that correspond to the distance between the calculated minimum and maximum distances are incremented (that is, the histogram is transposed relatively to the definition in the original paper). The histogram can be used for contour matching.

## QueryHistValue*D

Queries the value of the histogram bin.

**C:** float **cvQueryHistValue_1D**(CvHistogram **hist**, int **idx0**)

**C:** float **cvQueryHistValue_2D**(CvHistogram **hist**, int **idx0**, int **idx1**)

**C:** float **cvQueryHistValue_3D**(CvHistogram **hist**, int **idx0**, int **idx1**, int **idx2**)

**C:** float **cvQueryHistValue_nD**(CvHistogram **hist**, const int* **idx**)

**Python:** cv.**QueryHistValue_1D**(hist, idx0) → float

**Python:** cv.**QueryHistValue_2D**(hist, idx0, idx1) → float

**Python:** cv.**QueryHistValue_3D**(hist, idx0, idx1, idx2) → float

**Python:** cv.**QueryHistValue_nD**(hist, idx) → float

>> **Parameters**

>>> **hist** – Histogram.

>>> **idx0** – 0-th index.

>>> **idx1** – 1-st index.

>>> **idx2** – 2-nd index.

>>> **idx** – Array of indices.

The macros return the value of the specified bin of the 1D, 2D, 3D, or N-D histogram. In case of a sparse histogram, the function returns 0. If the bin is not present in the histogram, no new bin is created.

## GetHistValue_?D

Returns a pointer to the histogram bin.

**C:** float **cvGetHistValue_1D**(CvHistogram **hist**, int **idx0**)

**C:** float **cvGetHistValue_2D**(CvHistogram **hist**, int **idx0**, int **idx1**)

**C:** float **cvGetHistValue_3D**(CvHistogram **hist**, int **idx0**, int **idx1**, int **idx2**)

**C:** float **cvGetHistValue_nD**(CvHistogram **hist**, int **idx**)

>> **Parameters**

>>> **hist** – Histogram.

>>> **idx0** – 0-th index.

> **idx1** – 1-st index.

> **idx2** – 2-nd index.

> **idx** – Array of indices.

```
#define cvGetHistValue_1D( hist, idx0 )
    ((float*)(cvPtr1D( (hist)->bins, (idx0), 0 ))
#define cvGetHistValue_2D( hist, idx0, idx1 )
    ((float*)(cvPtr2D( (hist)->bins, (idx0), (idx1), 0 )))
#define cvGetHistValue_3D( hist, idx0, idx1, idx2 )
    ((float*)(cvPtr3D( (hist)->bins, (idx0), (idx1), (idx2), 0 )))
#define cvGetHistValue_nD( hist, idx )
    ((float*)(cvPtrND( (hist)->bins, (idx), 0 )))
```

The macros `GetHistValue` return a pointer to the specified bin of the 1D, 2D, 3D, or N-D histogram. In case of a sparse histogram, the function creates a new bin and sets it to 0, unless it exists already.

## 16.4 Planar Subdivisions (C API)

### CvSubdiv2D

**struct CvSubdiv2D**

Planar subdivision.
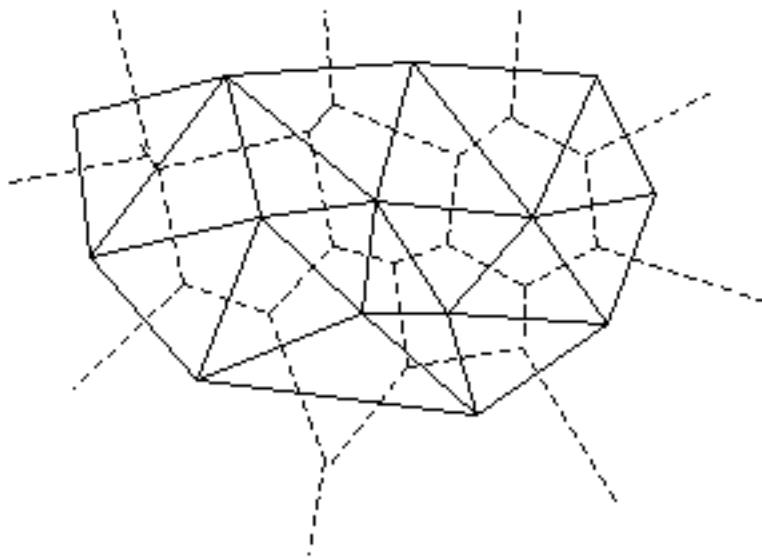
```
#define CV_SUBDIV2D_FIELDS()    \
    CV_GRAPH_FIELDS()           \
    int   quad_edges;           \
    int   is_geometry_valid;    \
    CvSubdiv2DEdge recent_edge; \
    CvPoint2D32f  topleft;      \
    CvPoint2D32f  bottomright;

typedef struct CvSubdiv2D
{
    CV_SUBDIV2D_FIELDS()
}
CvSubdiv2D;
```

Planar subdivision is the subdivision of a plane into a set of non-overlapped regions (facets) that cover the whole plane. The above structure describes a subdivision built on a 2D point set, where the points are linked together and form a planar graph, which, together with a few edges connecting the exterior subdivision points (namely, convex hull points) with infinity, subdivides a plane into facets by its edges.

For every subdivision, there is a dual subdivision in which facets and points (subdivision vertices) swap their roles. This means that a facet is treated as a vertex (called a virtual point below) of the dual subdivision and the original subdivision vertices become facets. In the figure below, the original subdivision is marked with solid lines and dual subdivision - with dotted lines.

OpenCV subdivides a plane into triangles using the Delaunay's algorithm. Subdivision is built iteratively starting from a dummy triangle that includes all the subdivision points for sure. In this case, the dual subdivision is a Voronoi diagram of the input 2D point set. The subdivisions can be used for the 3D piece-wise transformation of a plane, morphing, fast location of points on the plane, building special graphs (such as NNG,RNG), and so forth.

## CvQuadEdge2D

**struct `CvQuadEdge2D`**
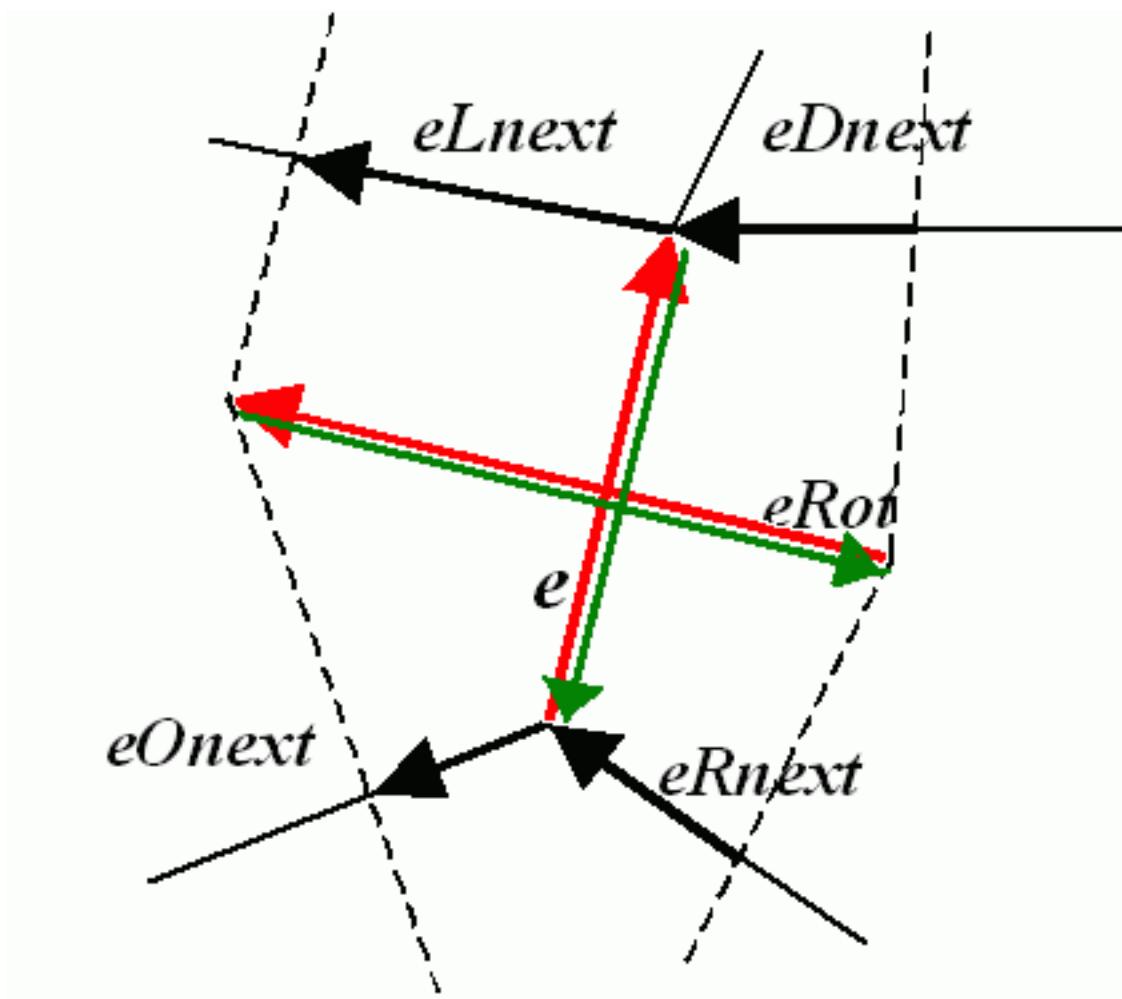
Quad-edge of a planar subdivision.

```
/* one of edges within quad-edge, lower 2 bits is index (0..3)
   and upper bits are quad-edge pointer */
typedef long CvSubdiv2DEdge;

/* quad-edge structure fields */
#define CV_QUADEDGE2D_FIELDS()      \
    int flags;                      \
    struct CvSubdiv2DPoint* pt[4]; \
    CvSubdiv2DEdge  next[4];

typedef struct CvQuadEdge2D
{
    CV_QUADEDGE2D_FIELDS()
}
CvQuadEdge2D;
```

Quad-edge is a basic element of a subdivision containing four edges (e, eRot, reversed e, and reversed eRot):

## CvSubdiv2DPoint

**struct CvSubdiv2DPoint**

Point of an original or dual subdivision.

```
#define CV_SUBDIV2D_POINT_FIELDS()\
    int            flags;      \
    CvSubdiv2DEdge first;      \
    CvPoint2D32f   pt;         \
    int id;

#define CV_SUBDIV2D_VIRTUAL_POINT_FLAG (1 << 30)

typedef struct CvSubdiv2DPoint
{
    CV_SUBDIV2D_POINT_FIELDS()
}
CvSubdiv2DPoint;
```

- **id** This integer can be used to index auxiliary data associated with each vertex of the planar subdivision.

## CalcSubdivVoronoi2D

Calculates the coordinates of the Voronoi diagram cells.

**C:** void **cvCalcSubdivVoronoi2D**(CvSubdiv2D* **subdiv**)

**Python:** cv.**CalcSubdivVoronoi2D**(subdiv) → None

>> **Parameters**

>>>> **subdiv** – Delaunay subdivision, in which all the points are already added.

The function calculates the coordinates of virtual points. All virtual points corresponding to a vertex of the original subdivision form (when connected together) a boundary of the Voronoi cell at that point.

## ClearSubdivVoronoi2D

Removes all virtual points.

**C:** void **cvClearSubdivVoronoi2D**(CvSubdiv2D* **subdiv**)

**Python:** cv.**ClearSubdivVoronoi2D**(subdiv) → None

>> **Parameters**

>>>> **subdiv** – Delaunay subdivision.

The function removes all of the virtual points. It is called internally in `CalcSubdivVoronoi2D()` if the subdivision was modified after the previous call to the function.

## CreateSubdivDelaunay2D

Creates an empty Delaunay triangulation.

**C:** CvSubdiv2D* **cvCreateSubdivDelaunay2D**(CvRect **rect**, CvMemStorage* **storage**)

**Python:** cv.**CreateSubdivDelaunay2D**(rect, storage) → CvSubdiv2D

>> **Parameters**

>>>> **rect** – Rectangle that includes all of the 2D points that are to be added to the subdivision.

>>>> **storage** – Container for the subdivision.

The function creates an empty Delaunay subdivision where 2D points can be added using the function `SubdivDelaunay2DInsert()` . All of the points to be added must be within the specified rectangle, otherwise a runtime error is raised.

Note that the triangulation is a single large triangle that covers the given rectangle. Hence the three vertices of this triangle are outside the rectangle `rect` .

## FindNearestPoint2D

Finds the subdivision vertex closest to the given point.

**C:** CvSubdiv2DPoint* **cvFindNearestPoint2D**(CvSubdiv2D* **subdiv**, CvPoint2D32f **pt**)

**Python:** cv.**FindNearestPoint2D**(subdiv, pt) → point

>> **Parameters**

>>>> **subdiv** – Delaunay or another subdivision.

> **pt** – Input point.

The function is another function that locates the input point within the subdivision. It finds the subdivision vertex that is the closest to the input point. It is not necessarily one of vertices of the facet containing the input point, though the facet (located using `Subdiv2DLocate()` ) is used as a starting point. The function returns a pointer to the found subdivision vertex.

## Subdiv2DEdgeDst

Returns the edge destination.

**C:** CvSubdiv2DPoint* **cvSubdiv2DEdgeDst**(CvSubdiv2DEdge **edge**)

**Python:** cv.**Subdiv2DEdgeDst**(edge) → point

> **Parameters**
>
> > **edge** – Subdivision edge (not a quad-edge).

The function returns the edge destination. The returned pointer may be NULL if the edge is from a dual subdivision and the virtual point coordinates are not calculated yet. The virtual points can be calculated using the function `CalcSubdivVoronoi2D()`.

## Subdiv2DGetEdge

Returns one of the edges related to the given edge.

**C:** CvSubdiv2DEdge **cvSubdiv2DGetEdge**(CvSubdiv2DEdge **edge**, CvNextEdgeType **type**)
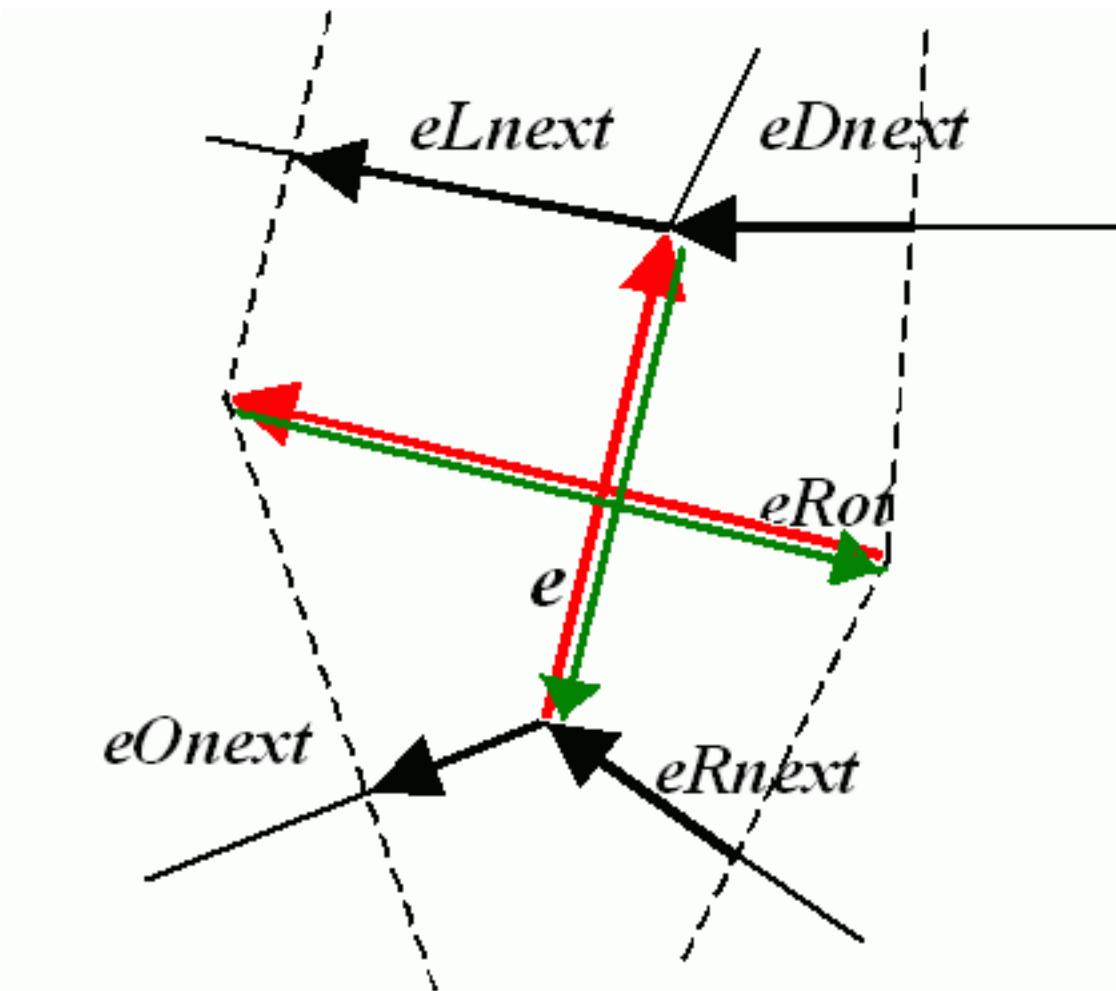
**Python:** cv.**Subdiv2DGetEdge**(edge, type) → CvSubdiv2DEdge

> **Parameters**
>
> > **edge** – Subdivision edge (not a quad-edge).
> >
> > **type** – Parameter specifying which of the related edges to return. The following values are possible:
> >
> > – **CV_NEXT_AROUND_ORG** next around the edge origin ( `eOnext` on the picture below if `e` is the input edge)
> >
> > – **CV_NEXT_AROUND_DST** next around the edge vertex ( `eDnext` )
> >
> > – **CV_PREV_AROUND_ORG** previous around the edge origin (reversed `eRnext` )
> >
> > – **CV_PREV_AROUND_DST** previous around the edge destination (reversed `eLnext` )
> >
> > – **CV_NEXT_AROUND_LEFT** next around the left facet ( `eLnext` )
> >
> > – **CV_NEXT_AROUND_RIGHT** next around the right facet ( `eRnext` )
> >
> > – **CV_PREV_AROUND_LEFT** previous around the left facet (reversed `eOnext` )
> >
> > – **CV_PREV_AROUND_RIGHT** previous around the right facet (reversed `eDnext` )

The function returns one of the edges related to the input edge.

### Subdiv2DNextEdge

Returns next edge around the edge origin.

**C:** CvSubdiv2DEdge **cvSubdiv2DNextEdge**(CvSubdiv2DEdge **edge**)

**Python:** cv.**Subdiv2DNextEdge**(edge) → CvSubdiv2DEdge

> **Parameters**
>
> > **edge** – Subdivision edge (not a quad-edge).

The function returns the next edge around the edge origin: `eOnext` on the picture above if `e` is the input edge).

### Subdiv2DLocate

Returns the location of a point within a Delaunay triangulation.

**C:** CvSubdiv2DPointLocation **cvSubdiv2DLocate**(CvSubdiv2D* **subdiv**, CvPoint2D32f **pt**, CvSubdiv2DEdge* **edge**, CvSubdiv2DPoint** **vertex**=NULL )

---

**Python:** cv.**Subdiv2DLocate**(subdiv, pt) -> (loc, where)

>> **Parameters**

>>> **subdiv** – Delaunay or another subdivision.

>>> **pt** – Point to locate.

>>> **edge** – Output edge that the point belongs to or is located to the right of it.

>>> **vertex** – Optional output vertex double pointer the input point coincides with.

The function locates the input point within the subdivision. There are five cases:

• The point falls into some facet. The function returns `CV_PTLOC_INSIDE` and `*edge` will contain one of edges of the facet.

• The point falls onto the edge. The function returns `CV_PTLOC_ON_EDGE` and `*edge` will contain this edge.

• The point coincides with one of the subdivision vertices. The function returns `CV_PTLOC_VERTEX` and `*vertex` will contain a pointer to the vertex.

• The point is outside the subdivision reference rectangle. The function returns `CV_PTLOC_OUTSIDE_RECT` and no pointers are filled.

• One of input arguments is invalid. A runtime error is raised or, if silent or "parent" error processing mode is selected, `CV_PTLOC_ERROR` is returnd.

## Subdiv2DRotateEdge

Returns another edge of the same quad-edge.

**C:** CvSubdiv2DEdge **cvSubdiv2DRotateEdge**(CvSubdiv2DEdge **edge**, int **rotate**)

**Python:** cv.**Subdiv2DRotateEdge**(edge, rotate) → CvSubdiv2DEdge

>> **Parameters**

>>> **edge** – Subdivision edge (not a quad-edge).

>>> **rotate** – Parameter specifying which of the edges of the same quad-edge as the input one to return. The following values are possible:

>>>> – **0** the input edge ( `e` on the picture below if `e` is the input edge)

>>>> – **1** the rotated edge ( `eRot` )

>>>> – **2** the reversed edge (reversed `e` (in green))

>>>> – **3** the reversed rotated edge (reversed `eRot` (in green))

The function returns one of the edges of the same quad-edge as the input edge.

## SubdivDelaunay2DInsert

Inserts a single point into a Delaunay triangulation.

**C:** CvSubdiv2DPoint* **cvSubdivDelaunay2DInsert**(CvSubdiv2D* **subdiv**, CvPoint2D32f **pt**)

**Python:** cv.**SubdivDelaunay2DInsert**(subdiv, pt) → point

>> **Parameters**

>>> **subdiv** – Delaunay subdivision created by the function `CreateSubdivDelaunay2D()`.

**pt** – Inserted point.

The function inserts a single point into a subdivision and modifies the subdivision topology appropriately. If a point with the same coordinates exists already, no new point is added. The function returns a pointer to the allocated point. No virtual point coordinates are calculated at this stage.

# 16.5 Feature Detection and Description

## RandomizedTree

**class RandomizedTree**

Class containing a base structure for `RTreeClassifier`.

```cpp
class CV_EXPORTS RandomizedTree
{
public:
        friend class RTreeClassifier;

        RandomizedTree();
        ~RandomizedTree();

        void train(std::vector<BaseKeypoint> const& base_set,
                RNG &rng, int depth, int views,
                size_t reduced_num_dim, int num_quant_bits);
        void train(std::vector<BaseKeypoint> const& base_set,
                RNG &rng, PatchGenerator &make_patch, int depth,
                int views, size_t reduced_num_dim, int num_quant_bits);

        // next two functions are EXPERIMENTAL
        //(do not use unless you know exactly what you do)
        static void quantizeVector(float *vec, int dim, int N, float bnds[2],
                int clamp_mode=0);
        static void quantizeVector(float *src, int dim, int N, float bnds[2],
                uchar *dst);

        // patch_data must be a 32x32 array (no row padding)
        float* getPosterior(uchar* patch_data);
        const float* getPosterior(uchar* patch_data) const;
        uchar* getPosterior2(uchar* patch_data);

        void read(const char* file_name, int num_quant_bits);
        void read(std::istream &is, int num_quant_bits);
        void write(const char* file_name) const;
        void write(std::ostream &os) const;

        int classes() { return classes_; }
        int depth() { return depth_; }

        void discardFloatPosteriors() { freePosteriors(1); }

        inline void applyQuantization(int num_quant_bits)
                { makePosteriors2(num_quant_bits); }

private:
        int classes_;
        int depth_;
```

```
        int num_leaves_;
        std::vector<RTreeNode> nodes_;
        float **posteriors_;        // 16-byte aligned posteriors
        uchar **posteriors2_;       // 16-byte aligned posteriors
        std::vector<int> leaf_counts_;

        void createNodes(int num_nodes, RNG &rng);
        void allocPosteriorsAligned(int num_leaves, int num_classes);
        void freePosteriors(int which);
                // which: 1=posteriors_, 2=posteriors2_, 3=both
        void init(int classes, int depth, RNG &rng);
        void addExample(int class_id, uchar* patch_data);
        void finalize(size_t reduced_num_dim, int num_quant_bits);
        int getIndex(uchar* patch_data) const;
        inline float* getPosteriorByIndex(int index);
        inline uchar* getPosteriorByIndex2(int index);
        inline const float* getPosteriorByIndex(int index) const;
        void convertPosteriorsToChar();
        void makePosteriors2(int num_quant_bits);
        void compressLeaves(size_t reduced_num_dim);
        void estimateQuantPercForPosteriors(float perc[2]);
};
```

## RandomizedTree::train

Trains a randomized tree using an input set of keypoints.

**C++:** void RandomizedTree::**train**(vector<BaseKeypoint> const& **base_set**, RNG& **rng**, int **depth**, int **views**, size_t **reduced_num_dim**, int **num_quant_bits**)

**C++:** void RandomizedTree::**train**(vector<BaseKeypoint> const& **base_set**, RNG& **rng**, PatchGenerator& **make_patch**, int **depth**, int **views**, size_t **reduced_num_dim**, int **num_quant_bits**)

> **Parameters**
>
> > **base_set** – Vector of the BaseKeypoint type. It contains image keypoints used for training.
> >
> > **rng** – Random-number generator used for training.
> >
> > **make_patch** – Patch generator used for training.
> >
> > **depth** – Maximum tree depth.
> >
> > **views** – Number of random views of each keypoint neighborhood to generate.
> >
> > **reduced_num_dim** – Number of dimensions used in the compressed signature.
> >
> > **num_quant_bits** – Number of bits used for quantization.

## RandomizedTree::read

Reads a pre-saved randomized tree from a file or stream.

**C++:** RandomizedTree::**read**(const char* **file_name**, int **num_quant_bits**)

**C++:** RandomizedTree::**read**(std::istream& **is**, int **num_quant_bits**)

> **Parameters**
>
> > **file_name** – Name of the file that contains randomized tree data.

> **is** – Input stream associated with the file that contains randomized tree data.
>
> **num_quant_bits** – Number of bits used for quantization.

## RandomizedTree::write

Writes the current randomized tree to a file or stream.

**C++:** void RandomizedTree::**write**(const char* **file_name**) const

**C++:** void RandomizedTree::**write**(std::ostream& **os**) const

> **Parameters**
>
> > **file_name** – Name of the file where randomized tree data is stored.
> >
> > **os** – Output stream associated with the file where randomized tree data is stored.

## RandomizedTree::applyQuantization

**C++:** void RandomizedTree::**applyQuantization**(int **num_quant_bits**)
> Applies quantization to the current randomized tree.

> > **Parameters**
> >
> > > **num_quant_bits** – Number of bits used for quantization.

## RTreeNode

**struct RTreeNode**

Class containing a base structure for `RandomizedTree`.

```
struct RTreeNode
{
        short offset1, offset2;

        RTreeNode() {}

        RTreeNode(uchar x1, uchar y1, uchar x2, uchar y2)
                : offset1(y1*PATCH_SIZE + x1),
                offset2(y2*PATCH_SIZE + x2)
        {}

        //! Left child on 0, right child on 1
        inline bool operator() (uchar* patch_data) const
        {
                return patch_data[offset1] > patch_data[offset2];
        }
};
```

## RTreeClassifier

**class RTreeClassifier**

Class containing `RTreeClassifier`. It represents the Calonder descriptor originally introduced by Michael Calonder.

```cpp
class CV_EXPORTS RTreeClassifier
{
public:
        static const int DEFAULT_TREES = 48;
        static const size_t DEFAULT_NUM_QUANT_BITS = 4;

        RTreeClassifier();

        void train(std::vector<BaseKeypoint> const& base_set,
                RNG &rng,
                int num_trees = RTreeClassifier::DEFAULT_TREES,
                int depth = DEFAULT_DEPTH,
                int views = DEFAULT_VIEWS,
                size_t reduced_num_dim = DEFAULT_REDUCED_NUM_DIM,
                int num_quant_bits = DEFAULT_NUM_QUANT_BITS,
                        bool print_status = true);
        void train(std::vector<BaseKeypoint> const& base_set,
                RNG &rng,
                PatchGenerator &make_patch,
                int num_trees = RTreeClassifier::DEFAULT_TREES,
                int depth = DEFAULT_DEPTH,
                int views = DEFAULT_VIEWS,
                size_t reduced_num_dim = DEFAULT_REDUCED_NUM_DIM,
                int num_quant_bits = DEFAULT_NUM_QUANT_BITS,
                 bool print_status = true);

        // sig must point to a memory block of at least
        //classes()*sizeof(float|uchar) bytes
        void getSignature(IplImage *patch, uchar *sig);
        void getSignature(IplImage *patch, float *sig);
        void getSparseSignature(IplImage *patch, float *sig,
                float thresh);

        static int countNonZeroElements(float *vec, int n, double tol=1e-10);
        static inline void safeSignatureAlloc(uchar **sig, int num_sig=1,
                        int sig_len=176);
        static inline uchar* safeSignatureAlloc(int num_sig=1,
                        int sig_len=176);

        inline int classes() { return classes_; }
        inline int original_num_classes()
                { return original_num_classes_; }

        void setQuantization(int num_quant_bits);
        void discardFloatPosteriors();

        void read(const char* file_name);
        void read(std::istream &is);
        void write(const char* file_name) const;
        void write(std::ostream &os) const;

        std::vector<RandomizedTree> trees_;

private:
        int classes_;
        int num_quant_bits_;
        uchar **posteriors_;
        ushort *ptemp_;
```

```
        int original_num_classes_;
        bool keep_floats_;
};
```

## RTreeClassifier::train

Trains a randomized tree classifier using an input set of keypoints.

**C++:** void RTreeClassifier::**train**(vector<BaseKeypoint> const& **base_set**, RNG& **rng**, int **num_trees**=RTreeClassifier::DEFAULT_TREES, int **depth**=RandomizedTree::DEFAULT_DEPTH, int **views**=RandomizedTree::DEFAULT_VIEWS, size_t **reduced_num_dim**=RandomizedTree::DEFAULT_REDUCED_NUM_DIM, int **num_quant_bits**=DEFAULT_NUM_QUANT_BITS )

**C++:** void RTreeClassifier::**train**(vector<BaseKeypoint> const& **base_set**, RNG& **rng**, PatchGenerator& **make_patch**, int **num_trees**=RTreeClassifier::DEFAULT_TREES, int **depth**=RandomizedTree::DEFAULT_DEPTH, int **views**=RandomizedTree::DEFAULT_VIEWS, size_t **reduced_num_dim**=RandomizedTree::DEFAULT_REDUCED_NUM_DIM, int **num_quant_bits**=DEFAULT_NUM_QUANT_BITS )

> **Parameters**

>> **base_set** – Vector of the `BaseKeypoint` type. It contains image keypoints used for training.

>> **rng** – Random-number generator used for training.

>> **make_patch** – Patch generator used for training.

>> **num_trees** – Number of randomized trees used in `RTreeClassificator`.

>> **depth** – Maximum tree depth.

>> **views** – Number of random views of each keypoint neighborhood to generate.

>> **reduced_num_dim** – Number of dimensions used in the compressed signature.

>> **num_quant_bits** – Number of bits used for quantization.

## RTreeClassifier::getSignature

Returns a signature for an image patch.

**C++:** void RTreeClassifier::**getSignature**(IplImage* **patch**, uchar* **sig**)

**C++:** void RTreeClassifier::**getSignature**(IplImage* **patch**, float* **sig**)

> **Parameters**

>> **patch** – Image patch to calculate the signature for.

>> **sig** – Output signature (array dimension is `reduced_num_dim`) .

## RTreeClassifier::getSparseSignature

Returns a sparse signature for an image patch

**C++:** void RTreeClassifier::**getSparseSignature**(IplImage* **patch**, float* **sig**, float **thresh**)

> **Parameters**

>> **patch** – Image patch to calculate the signature for.

>> **sig** – Output signature (array dimension is `reduced_num_dim`) .

>> **thresh** – Threshold used for compressing the signature.

Returns a signature for an image patch similarly to `getSignature` but uses a threshold for removing all signature elements below the threshold so that the signature is compressed.

## RTreeClassifier::countNonZeroElements

Returns the number of non-zero elements in an input array.

**C++:** **static** int RTreeClassifier::**countNonZeroElements**(float* **vec**, int **n**, double **tol**=1e-10)

> **Parameters**

>> **vec** – Input vector containing float elements.

>> **n** – Input vector size.

>> **tol** – Threshold used for counting elements. All elements less than `tol` are considered as zero elements.

## RTreeClassifier::read

Reads a pre-saved `RTreeClassifier` from a file or stream.

**C++:** void RTreeClassifier::**read**(const char* **file_name**)

**C++:** void RTreeClassifier::**read**(std::istream& **is**)

> **Parameters**

>> **file_name** – Name of the file that contains randomized tree data.

>> **is** – Input stream associated with the file that contains randomized tree data.

## RTreeClassifier::write

Writes the current `RTreeClassifier` to a file or stream.

**C++:** void RTreeClassifier::**write**(const char* **file_name**) const

**C++:** void RTreeClassifier::**write**(std::ostream& **os**) const

> **Parameters**

>> **file_name** – Name of the file where randomized tree data is stored.

>> **os** – Output stream associated with the file where randomized tree data is stored.

## RTreeClassifier::setQuantization

Applies quantization to the current randomized tree.

**C++:** void RTreeClassifier::**setQuantization**(int **num_quant_bits**)

> **Parameters**

      **num_quant_bits** – Number of bits used for quantization.

The example below demonstrates the usage of `RTreeClassifier` for matching the features. The features are extracted from the test and train images with SURF. Output is $best\_corr$ and $best\_corr\_idx$ arrays that keep the best probabilities and corresponding features indices for every train feature.

```cpp
CvMemStorage* storage = cvCreateMemStorage(0);
CvSeq *objectKeypoints = 0, *objectDescriptors = 0;
CvSeq *imageKeypoints = 0, *imageDescriptors = 0;
CvSURFParams params = cvSURFParams(500, 1);
cvExtractSURF( test_image, 0, &imageKeypoints, &imageDescriptors,
                storage, params );
cvExtractSURF( train_image, 0, &objectKeypoints, &objectDescriptors,
                storage, params );


RTreeClassifier detector;
int patch_width = PATCH_SIZE;
iint patch_height = PATCH_SIZE;
vector<BaseKeypoint> base_set;
int i=0;
CvSURFPoint* point;
for (i=0;i<(n_points > 0 ? n_points : objectKeypoints->total);i++)
{
        point=(CvSURFPoint*)cvGetSeqElem(objectKeypoints,i);
        base_set.push_back(
                BaseKeypoint(point->pt.x,point->pt.y,train_image));
}


        //Detector training
 RNG rng( cvGetTickCount() );
PatchGenerator gen(0,255,2,false,0.7,1.3,-CV_PI/3,CV_PI/3,
                        -CV_PI/3,CV_PI/3);

printf("RTree Classifier training...n");
detector.train(base_set,rng,gen,24,DEFAULT_DEPTH,2000,
        (int)base_set.size(), detector.DEFAULT_NUM_QUANT_BITS);
printf("Donen");

float* signature = new float[detector.original_num_classes()];
float* best_corr;
int* best_corr_idx;
if (imageKeypoints->total > 0)
{
        best_corr = new float[imageKeypoints->total];
        best_corr_idx = new int[imageKeypoints->total];
}

for(i=0; i < imageKeypoints->total; i++)
{
        point=(CvSURFPoint*)cvGetSeqElem(imageKeypoints,i);
        int part_idx = -1;
        float prob = 0.0f;

        CvRect roi = cvRect((int)(point->pt.x) - patch_width/2,
                (int)(point->pt.y) - patch_height/2,
                 patch_width, patch_height);
        cvSetImageROI(test_image, roi);
        roi = cvGetImageROI(test_image);
        if(roi.width != patch_width || roi.height != patch_height)
```

```
        {
                best_corr_idx[i] = part_idx;
                best_corr[i] = prob;
        }
        else
        {
                cvSetImageROI(test_image, roi);
                IplImage* roi_image =
                        cvCreateImage(cvSize(roi.width, roi.height),
                        test_image->depth, test_image->nChannels);
                cvCopy(test_image,roi_image);

                detector.getSignature(roi_image, signature);
                for (int j = 0; j< detector.original_num_classes();j++)
                {
                        if (prob < signature[j])
                        {
                                part_idx = j;
                                prob = signature[j];
                        }
                }

                best_corr_idx[i] = part_idx;
                best_corr[i] = prob;

                if (roi_image)
                        cvReleaseImage(&roi_image);
        }
        cvResetImageROI(test_image);
}
```

## 16.6 Common Interfaces of Descriptor Extractors

Extractors of keypoint descriptors in OpenCV have wrappers with a common interface that enables you to easily switch between different algorithms solving the same problem. This section is devoted to computing descriptors represented as vectors in a multidimensional space. All objects that implement the `vector` descriptor extractors inherit the `DescriptorExtractor` interface.

### CalonderDescriptorExtractor

**class** `CalonderDescriptorExtractor` : **public** `DescriptorExtractor`

Wrapping class for computing descriptors by using the `RTreeClassifier` class.

```
template<typename T>
class CalonderDescriptorExtractor : public DescriptorExtractor
{
public:
    CalonderDescriptorExtractor( const string& classifierFile );

    virtual void read( const FileNode &fn );
    virtual void write( FileStorage &fs ) const;
    virtual int descriptorSize() const;
    virtual int descriptorType() const;
protected:
```

```
    ...
}
```

## 16.7 Common Interfaces of Generic Descriptor Matchers

### OneWayDescriptorMatcher

**class OneWayDescriptorMatcher** : **public GenericDescriptorMatcher**

Wrapping class for computing, matching, and classifying descriptors using the `OneWayDescriptorBase` class.

```cpp
class OneWayDescriptorMatcher : public GenericDescriptorMatcher
{
public:
    class Params
    {
    public:
        static const int POSE_COUNT = 500;
        static const int PATCH_WIDTH = 24;
        static const int PATCH_HEIGHT = 24;
        static float GET_MIN_SCALE() { return 0.7f; }
        static float GET_MAX_SCALE() { return 1.5f; }
        static float GET_STEP_SCALE() { return 1.2f; }

        Params( int poseCount = POSE_COUNT,
                Size patchSize = Size(PATCH_WIDTH, PATCH_HEIGHT),
                string pcaFilename = string(),
                string trainPath = string(), string trainImagesList = string(),
                float minScale = GET_MIN_SCALE(), float maxScale = GET_MAX_SCALE(),
                float stepScale = GET_STEP_SCALE() );

        int poseCount;
        Size patchSize;
        string pcaFilename;
        string trainPath;
        string trainImagesList;

        float minScale, maxScale, stepScale;
    };

    OneWayDescriptorMatcher( const Params& params=Params() );
    virtual ~OneWayDescriptorMatcher();

    void initialize( const Params& params, const Ptr<OneWayDescriptorBase>& base=Ptr<OneWayDescriptorBase>() );

    // Clears keypoints stored in collection and OneWayDescriptorBase
    virtual void clear();

    virtual void train();

    virtual bool isMaskSupported();

    virtual void read( const FileNode &fn );
    virtual void write( FileStorage& fs ) const;

    virtual Ptr<GenericDescriptorMatcher> clone( bool emptyTrainData=false ) const;
```

```
protected:
    ...
};
```

## FernDescriptorMatcher

**class FernDescriptorMatcher** : **public GenericDescriptorMatcher**

Wrapping class for computing, matching, and classifying descriptors using the FernClassifier class.

```
class FernDescriptorMatcher : public GenericDescriptorMatcher
{
public:
    class Params
    {
    public:
        Params( int nclasses=0,
                int patchSize=FernClassifier::PATCH_SIZE,
                int signatureSize=FernClassifier::DEFAULT_SIGNATURE_SIZE,
                int nstructs=FernClassifier::DEFAULT_STRUCTS,
                int structSize=FernClassifier::DEFAULT_STRUCT_SIZE,
                int nviews=FernClassifier::DEFAULT_VIEWS,
                int compressionMethod=FernClassifier::COMPRESSION_NONE,
                const PatchGenerator& patchGenerator=PatchGenerator() );

        Params( const string& filename );

        int nclasses;
        int patchSize;
        int signatureSize;
        int nstructs;
        int structSize;
        int nviews;
        int compressionMethod;
        PatchGenerator patchGenerator;

        string filename;
    };

    FernDescriptorMatcher( const Params& params=Params() );
    virtual ~FernDescriptorMatcher();

    virtual void clear();

    virtual void train();

    virtual bool isMaskSupported();

    virtual void read( const FileNode &fn );
    virtual void write( FileStorage& fs ) const;

    virtual Ptr<GenericDescriptorMatcher> clone( bool emptyTrainData=false ) const;

protected:
        ...
};
```

# OCL. OPENCL-ACCELERATED COMPUTER VISION

## 17.1 OpenCL Module Introduction

### General Information

The OpenCV OCL module is a set of classes and functions to utilize OpenCL compatible device. It should support any device compatible with OpenCL 1.1. The module includes utility functions, low-level vision primitives, and a few high-level algorithms ready to be used in the end-user applications.

The OpenCV OCL module is designed as a host-level API plus device-level kernels. The device-level kernels are converted to text string and are compiled at runtime, so it need OpenCL runtime support. To correctly run the OpenCV OCL module, make sure you have OpenCL runtime provided by your device vendor, which is device driver normally.

The OpenCV OCL module is designed for ease of use and does not require any knowledge of OpenCL. Though, such a knowledge will certainly be useful to handle non-trivial cases or achieve the highest performance. It is helpful to understand the cost of various operations, what the module does, what the preferred data formats are, and so on. Since there is data transfer between OpenCL host and OpenCL device, for better performance it's recommended to copy data once to the OpenCL host memory (i.e. copy `cv::Mat` to `cv::ocl::OclMat`), then call several `cv::ocl` functions and then copy the result back to CPU memory, rather than do forward and backward transfer for each OCL function.

To enable OCL support, configure OpenCV using CMake with the option `WITH\_OPENCL=ON`. If the option is passed and if OpenCL SDK is installed (e.g. on MacOSX it's always the case), the full-featured OpenCV OCL module will be built. Otherwise, the module will not be built.

Right now, the user should define the `cv::ocl::Info` class in the application and call `cv::ocl::getDevice` before any `cv::ocl::<func>`. This operation initialize OpenCL runtime and set the first found device as computing device. If there is more than one device and you want to use non-default device, you should call `cv::ocl::setDevice`.

In the current version, all the threads share the same context and device so the multi-devices are not supported. This is to be fixed in future releases.

## 17.2 Data Structures and Functions

### ocl::Info

**class** `ocl::`**`Info`**

this class should be maintained by the user and be passed to getDevice

## ocl::getDevice

Returns the list of devices

**C++:** int ocl::**getDevice**(std::vector<Info>& **oclinfo**, int **devicetype**=CVCL_DEVICE_TYPE_GPU )

>   **Parameters**

>>   **oclinfo** – Output vector of `ocl::Info` structures

>>   **devicetype** – One of `CVCL_DEVICE_TYPE_GPU`, `CVCL_DEVICE_TYPE_CPU` or `CVCL_DEVICE_TYPE_DEFAULT`.

the function must be called before any other `cv::ocl` functions; it initializes ocl runtime.

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