

## Camera Calibration and 3D Reconstruction

### Modules

[Fisheye camera model](#)

[C API](#)

### Classes

`struct cv::CirclesGridFinderParameters`

`struct cv::CirclesGridFinderParameters2`

`class cv::StereoBM`

Class for computing stereo correspondence using the block matching algorithm, introduced and contributed to OpenCV by K. Konolige. More...

`class cv::StereoMatcher`

The base class for stereo correspondence algorithms. More...

`class cv::StereoSGBM`

The class implements the modified H. Hirschmuller algorithm [86] that differs from the original one as follows: More...

### Enumerations

```
enum {
    cv::LMEDS = 4,
    cv::RANSAC = 8,
    cv::RHO = 16
}
```

type of the robust estimation algorithm More...

```
enum {
    cv::SOLVEPNP_ITERATIVE = 0,
    cv::SOLVEPNP_EPNP = 1,
    cv::SOLVEPNP_P3P = 2,
    cv::SOLVEPNP_DLS = 3,
    cv::SOLVEPNP_UPNP = 4,
    cv::SOLVEPNP_AP3P = 5,
    cv::SOLVEPNP_MAX_COUNT
}
```

```
enum {
    cv::CALIB_CB_ADAPTIVE_THRESH = 1,
    cv::CALIB_CB_NORMALIZE_IMAGE = 2,
    cv::CALIB_CB_FILTER_QUADS = 4,
    cv::CALIB_CB_FAST_CHECK = 8
}
```

```
enum {
    cv::CALIB_CB_SYMMETRIC_GRID = 1,
    cv::CALIB_CB_ASYMMETRIC_GRID = 2,
    cv::CALIB_CB_CLUSTERING = 4
}
```

```
enum {
    cv::CALIB_USE_INTRINSIC_GUESS = 0x00001,
    cv::CALIB_FIX_ASPECT_RATIO = 0x00002,
    cv::CALIB_FIX_PRINCIPAL_POINT = 0x00004,
    cv::CALIB_ZERO_TANGENT_DIST = 0x00008,
    cv::CALIB_FIX_FOCAL_LENGTH = 0x00010,
    cv::CALIB_FIX_K1 = 0x00020,
    cv::CALIB_FIX_K2 = 0x00040,
    cv::CALIB_FIX_K3 = 0x00080,
    cv::CALIB_FIX_K4 = 0x00800,
    cv::CALIB_FIX_K5 = 0x01000,
    cv::CALIB_FIX_K6 = 0x02000,
    cv::CALIB_RATIONAL_MODEL = 0x04000,
    cv::CALIB_THIN_PRISM_MODEL = 0x08000,
    cv::CALIB_FIX_S1_S2_S3_S4 = 0x10000,
    cv::CALIB_TILTED_MODEL = 0x40000,
    cv::CALIB_FIX_TAUX_TAUY = 0x80000,
    cv::CALIB_USE_QR = 0x100000,
    cv::CALIB_FIX_TANGENT_DIST = 0x200000,
    cv::CALIB_FIX_INTRINSIC = 0x00100,
    cv::CALIB SAME_FOCAL_LENGTH = 0x00200,
    cv::CALIB_ZERO_DISPARITY = 0x00400,
```

```

    cv::CALIB_USE_LU = (1 << 17),
    cv::CALIB_USE_EXTRINSIC_GUESS = (1 << 22)
}

enum {
    cv::FM_7POINT = 1,
    cv::FM_8POINT = 2,
    cv::FM_LMEDS = 4,
    cv::FM_RANSAC = 8
}
the algorithm for finding fundamental matrix More...

```

## Functions

double	<a href="#">cv::calibrateCamera (InputArrayOfArrays objectPoints, InputArrayOfArrays imagePoints, Size imageSize, InputOutputArray cameraMatrix, InputOutputArray distCoeffs, OutputArrayOfArrays rvecs, OutputArrayOfArrays tvecs, OutputArray stdDeviationsIntrinsics, OutputArray stdDeviationsExtrinsics, OutputArray perViewErrors, int flags=0, TermCriteria criteria=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, DBL_EPSILON))</a>	Finds the camera intrinsic and extrinsic parameters from several views of a calibration pattern. <a href="#">More...</a>
double	<a href="#">cv::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))</a>	
void	<a href="#">cv::calibrationMatrixValues (InputArray cameraMatrix, Size imageSize, double apertureWidth, double apertureHeight, double &amp;fovX, double &amp;fovY, double &amp;focalLength, Point2d &amp;principalPoint, double &amp;aspectRatio)</a>	Computes useful camera characteristics from the camera matrix. <a href="#">More...</a>
void	<a href="#">cv::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())</a>	Combines two rotation-and-shift transformations. <a href="#">More...</a>
void	<a href="#">cv::computeCorrespondEpilines (InputArray points, int whichImage, InputArray F, OutputArray lines)</a>	For points in an image of a stereo pair, computes the corresponding epilines in the other image. <a href="#">More...</a>
void	<a href="#">cv::convertPointsFromHomogeneous (InputArray src, OutputArray dst)</a>	Converts points from homogeneous to Euclidean space. <a href="#">More...</a>
void	<a href="#">cv::convertPointsHomogeneous (InputArray src, OutputArray dst)</a>	Converts points to/from homogeneous coordinates. <a href="#">More...</a>
void	<a href="#">cv::convertPointsToHomogeneous (InputArray src, OutputArray dst)</a>	Converts points from Euclidean to homogeneous space. <a href="#">More...</a>
void	<a href="#">cv::correctMatches (InputArray F, InputArray points1, InputArray points2, OutputArray newPoints1, OutputArray newPoints2)</a>	Refines coordinates of corresponding points. <a href="#">More...</a>
void	<a href="#">cv::decomposeEssentialMat (InputArray E, OutputArray R1, OutputArray R2, OutputArray t)</a>	Decompose an essential matrix to possible rotations and translation. <a href="#">More...</a>
int	<a href="#">cv::decomposeHomographyMat (InputArray H, InputArray K, OutputArrayOfArrays rotations, OutputArrayOfArrays translations, OutputArrayOfArrays normals)</a>	Decompose a homography matrix to rotation(s), translation(s) and plane normal(s). <a href="#">More...</a>
void	<a href="#">cv::decomposeProjectionMatrix (InputArray projMatrix, OutputArray cameraMatrix, OutputArray rotMatrix, OutputArray transVect, OutputArray rotMatrixX=noArray(), OutputArray rotMatrixY=noArray(), OutputArray rotMatrixZ=noArray(), OutputArray eulerAngles=noArray())</a>	Decomposes a projection matrix into a rotation matrix and a camera matrix. <a href="#">More...</a>
void	<a href="#">cv::drawChessboardCorners (InputOutputArray image, Size patternSize, InputArray corners, bool patternWasFound)</a>	Renders the detected chessboard corners. <a href="#">More...</a>
cv::Mat	<a href="#">cv::estimateAffine2D (InputArray from, InputArray to, OutputArray inliers=noArray(), int method=RANSAC, double ransacReprojThreshold=3, size_t maxIters=2000, double confidence=0.99, size_t refineIters=10)</a>	Computes an optimal affine transformation between two 2D point sets. <a href="#">More...</a>
int	<a href="#">cv::estimateAffine3D (InputArray src, InputArray dst, OutputArray out, OutputArray inliers, double ransacThreshold=3, double confidence=0.99)</a>	Computes an optimal affine transformation between two 3D point sets. <a href="#">More...</a>
cv::Mat	<a href="#">cv::estimateAffinePartial2D (InputArray from, InputArray to, OutputArray inliers=noArray(), int method=RANSAC, double ransacReprojThreshold=3, size_t maxIters=2000, double confidence=0.99, size_t refineIters=10)</a>	Computes an optimal limited affine transformation with 4 degrees of freedom between two 2D point sets. <a href="#">More...</a>
void	<a href="#">cv::filterHomographyDecompByVisibleRefpoints (InputArrayOfArrays rotations, InputArrayOfArrays normals, InputArray beforePoints, InputArray afterPoints, OutputArray possibleSolutions, InputArray pointsMask=noArray())</a>	Filters homography decompositions based on additional information. <a href="#">More...</a>
void	<a href="#">cv::filterSpeckles (InputOutputArray img, double newVal, int maxSpeckleSize, double maxDiff, InputOutputArray buf=noArray())</a>	Filters off small noise blobs (speckles) in the disparity map. <a href="#">More...</a>
bool	<a href="#">cv::find4QuadCornerSubpix (InputArray img, InputOutputArray corners, Size region_size)</a>	finds subpixel-accurate positions of the chessboard corners <a href="#">More...</a>

bool	<code>cv::findChessboardCorners (InputArray image, Size patternSize, OutputArray corners, int flags=CV_CALIB_CB_ADAPTIVE_THRESH+CV_CALIB_CB_NORMALIZE_IMAGE)</code>	Finds the positions of internal corners of the chessboard. More...
bool	<code>cv::findCirclesGrid (InputArray image, Size patternSize, OutputArray centers, int flags, const Ptr&lt; FeatureDetector &gt; &amp;blobDetector, CirclesGridFinderParameters parameters)</code>	Finds centers in the grid of circles. More...
bool	<code>cv::findCirclesGrid (InputArray image, Size patternSize, OutputArray centers, int flags=CV_CALIB_CB_SYMMETRIC_GRID, const Ptr&lt; FeatureDetector &gt; &amp;blobDetector=SimpleBlobDetector::create())</code>	
bool	<code>cv::findCirclesGrid2 (InputArray image, Size patternSize, OutputArray centers, int flags, const Ptr&lt; FeatureDetector &gt; &amp;blobDetector, CirclesGridFinderParameters2 parameters)</code>	
Mat	<code>cv::findEssentialMat (InputArray points1, InputArray points2, InputArray cameraMatrix, int method=RANSAC, double prob=0.999, double threshold=1.0, OutputArray mask=noArray())</code>	Calculates an essential matrix from the corresponding points in two images. More...
Mat	<code>cv::findEssentialMat (InputArray points1, InputArray points2, double focal=1.0, Point2d pp=Point2d(0, 0), int method=RANSAC, double prob=0.999, double threshold=1.0, OutputArray mask=noArray())</code>	
Mat	<code>cv::findFundamentalMat (InputArray points1, InputArray points2, int method=FM_RANSAC, double ransacReprojThreshold=3., double confidence=0.99, OutputArray mask=noArray())</code>	Calculates a fundamental matrix from the corresponding points in two images. More...
Mat	<code>cv::findFundamentalMat (InputArray points1, InputArray points2, OutputArray mask, int method=FM_RANSAC, double ransacReprojThreshold=3., double confidence=0.99)</code>	
Mat	<code>cv::findHomography (InputArray srcPoints, InputArray dstPoints, int method=0, double ransacReprojThreshold=3, OutputArray mask=noArray(), const int maxIters=2000, const double confidence=0.995)</code>	Finds a perspective transformation between two planes. More...
Mat	<code>cv::findHomography (InputArray srcPoints, InputArray dstPoints, OutputArray mask, int method=0, double ransacReprojThreshold=3)</code>	
Mat	<code>cv::getOptimalNewCameraMatrix (InputArray cameraMatrix, InputArray distCoeffs, Size imageSize, double alpha, Size newImgSize=Size(), Rect *validPixROI=0, bool centerPrincipalPoint=false)</code>	Returns the new camera matrix based on the free scaling parameter. More...
Rect	<code>cv::getValidDisparityROI (Rect roi1, Rect roi2, int minDisparity, int numberOfWorkDisparities, int SADWindowSize)</code>	computes valid disparity ROI from the valid ROIs of the rectified images (that are returned by <code>cv::stereoRectify()</code> ) More...
Mat	<code>cv::initCameraMatrix2D (InputArrayOfArrays objectPoints, InputArrayOfArrays imagePoints, Size imageSize, double aspectRatio=1.0)</code>	Finds an initial camera matrix from 3D-2D point correspondences. More...
void	<code>cv::matMulDeriv (InputArray A, InputArray B, OutputArray dABdA, OutputArray dABdB)</code>	Computes partial derivatives of the matrix product for each multiplied matrix. More...
void	<code>cv::projectPoints (InputArray objectPoints, InputArray rvec, InputArray tvec, InputArray cameraMatrix, InputArray distCoeffs, OutputArray imagePoints, OutputArray jacobian=noArray(), double aspectRatio=0)</code>	Projects 3D points to an image plane. More...
int	<code>cv::recoverPose (InputArray E, InputArray points1, InputArray points2, InputArray cameraMatrix, OutputArray R, OutputArray t, InputOutputArray mask=noArray())</code>	Recover relative camera rotation and translation from an estimated essential matrix and the corresponding points in two images, using cheirality check. Returns the number of inliers which pass the check. More...
int	<code>cv::recoverPose (InputArray E, InputArray points1, InputArray points2, OutputArray R, OutputArray t, double focal=1.0, Point2d pp=Point2d(0, 0), InputOutputArray mask=noArray())</code>	
int	<code>cv::recoverPose (InputArray E, InputArray points1, InputArray points2, InputArray cameraMatrix, OutputArray R, OutputArray t, double distanceThresh, InputOutputArray mask=noArray(), OutputArray triangulatedPoints=noArray())</code>	
float	<code>cv::rectify3Collinear (InputArray cameraMatrix1, InputArray distCoeffs1, InputArray cameraMatrix2, InputArray distCoeffs2, InputArray cameraMatrix3, InputArray distCoeffs3, InputArrayOfArrays imgpt1, InputArrayOfArrays imgpt3, Size imageSize, InputArray R12, InputArray T12, InputArray R13, InputArray T13, OutputArray R1, OutputArray R2, OutputArray R3, OutputArray P1, OutputArray P2, OutputArray P3, OutputArray Q, double alpha, Size newImgSize, Rect *roi1, Rect *roi2, int flags)</code>	computes the rectification transformations for 3-head camera, where all the heads are on the same line. More...
void	<code>cv::reprojectImageTo3D (InputArray disparity, OutputArray _3dImage, InputArray Q, bool handleMissingValues=false, int ddepth=-1)</code>	Reprojects a disparity image to 3D space. More...
void	<code>cv::Rodrigues (InputArray src, OutputArray dst, OutputArray jacobian=noArray())</code>	Converts a rotation matrix to a rotation vector or vice versa. More...
Vec3d	<code>cv::RQDecomp3x3 (InputArray src, OutputArray mtxR, OutputArray mtxQ, OutputArray Qx=noArray(), OutputArray Qy=noArray(), OutputArray Qz=noArray())</code>	Computes an RQ decomposition of 3x3 matrices. More...
double	<code>cv::sampsonDistance (InputArray pt1, InputArray pt2, InputArray F)</code>	Calculates the Sampson Distance between two points. More...
int	<code>cv::solveP3P (InputArray objectPoints, InputArray imagePoints, InputArray cameraMatrix, InputArray distCoeffs, OutputArrayOfArrays rvecs, OutputArrayOfArrays tvecs, int flags)</code>	Finds an object pose from 3 3D-2D point correspondences. More...
bool	<code>cv::solvePnP (InputArray objectPoints, InputArray imagePoints, InputArray cameraMatrix, InputArray distCoeffs, OutputArray rvec, OutputArray tvec, bool useExtrinsicGuess=false, int flags=SOLVEPNP_ITERATIVE)</code>	Finds an object pose from 3D-2D point correspondences. More...

bool	<code>cv::solvePnP</code>	( <b>InputArray</b> objectPoints, <b>InputArray</b> imagePoints, <b>InputArray</b> cameraMatrix, <b>InputArray</b> distCoeffs, <b>OutputArray</b> rvec, <b>OutputArray</b> tvec, bool useExtrinsicGuess=false, int iterationsCount=100, float reprojectionError=8.0, double confidence=0.99, <b>OutputArray</b> inliers= <b>noArray</b> (), int flags= <b>SOLVEPNP_ITERATIVE</b> )	Finds an object pose from 3D-2D point correspondences using the RANSAC scheme. <a href="#">More...</a>
double	<code>cv::stereoCalibrate</code>	( <b>InputArrayOfArrays</b> objectPoints, <b>InputArrayOfArrays</b> imagePoints1, <b>InputArrayOfArrays</b> imagePoints2, <b>InputOutputArray</b> cameraMatrix1, <b>InputOutputArray</b> distCoeffs1, <b>InputOutputArray</b> cameraMatrix2, <b>InputOutputArray</b> distCoeffs2, <b>Size</b> imageSize, <b>InputOutputArray</b> R, <b>InputOutputArray</b> T, <b>OutputArray</b> E, <b>OutputArray</b> F, <b>OutputArray</b> perViewErrors, int flags= <b>CALIB_FIX_INTRINSIC</b> , TermCriteria criteria=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 1e-6))	Calibrates the stereo camera. <a href="#">More...</a>
double	<code>cv::stereoCalibrate</code>	( <b>InputArrayOfArrays</b> objectPoints, <b>InputArrayOfArrays</b> imagePoints1, <b>InputArrayOfArrays</b> imagePoints2, <b>InputOutputArray</b> cameraMatrix1, <b>InputOutputArray</b> distCoeffs1, <b>InputOutputArray</b> cameraMatrix2, <b>InputOutputArray</b> distCoeffs2, <b>Size</b> imageSize, <b>OutputArray</b> R, <b>OutputArray</b> T, <b>OutputArray</b> E, <b>OutputArray</b> F, int flags= <b>CALIB_FIX_INTRINSIC</b> , TermCriteria criteria=TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 1e-6))	Calibrates the stereo camera. <a href="#">More...</a>
void	<code>cv::stereoRectify</code>	( <b>InputArray</b> cameraMatrix1, <b>InputArray</b> distCoeffs1, <b>InputArray</b> cameraMatrix2, <b>InputArray</b> distCoeffs2, <b>Size</b> imageSize, <b>InputArray</b> R, <b>InputArray</b> T, <b>OutputArray</b> R1, <b>OutputArray</b> R2, <b>OutputArray</b> P1, <b>OutputArray</b> P2, <b>OutputArray</b> Q, int flags= <b>CALIB_ZERO_DISPARITY</b> , double alpha=-1, <b>Size</b> newImageSize= <b>Size</b> (), <b>Rect</b> *validPixROI1=0, <b>Rect</b> *validPixROI2=0)	Computes rectification transforms for each head of a calibrated stereo camera. <a href="#">More...</a>
bool	<code>cv::stereoRectifyUncalibrated</code>	( <b>InputArray</b> points1, <b>InputArray</b> points2, <b>InputArray</b> F, <b>Size</b> imgSize, <b>OutputArray</b> H1, <b>OutputArray</b> H2, double threshold=5)	Computes a rectification transform for an uncalibrated stereo camera. <a href="#">More...</a>
void	<code>cv::triangulatePoints</code>	( <b>InputArray</b> projMatr1, <b>InputArray</b> projMatr2, <b>InputArray</b> projPoints1, <b>InputArray</b> projPoints2, <b>OutputArray</b> points4D)	Reconstructs points by triangulation. <a href="#">More...</a>
void	<code>cv::validateDisparity</code>	( <b>InputOutputArray</b> disparity, <b>InputArray</b> cost, int minDisparity, int numberOfDisparities, int disp12MaxDisp=1)	Validates disparity using the left-right check. The matrix "cost" should be computed by the stereo correspondence algorithm <a href="#">More...</a>

## Detailed Description

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
- $f_x, f_y$  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  $z \neq 0$ ):

$$\begin{bmatrix} x \\ y \\ z \end{bmatrix} = R \begin{bmatrix} X \\ Y \\ Z \end{bmatrix} + t$$

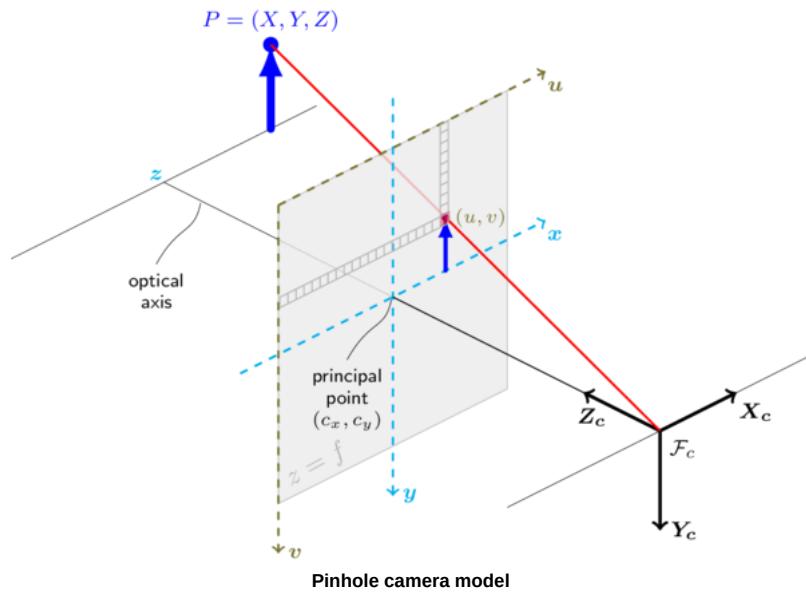
$$x' = x/z$$

$$y' = y/z$$

$$u = f_x * x' + c_x$$

$$v = f_y * y' + c_y$$

The following figure illustrates the pinhole camera model.



Real lenses usually have some distortion, mostly radial distortion and slight tangential distortion. So, the above model is extended as:

$$\begin{bmatrix} x \\ y \\ z \end{bmatrix} = R \begin{bmatrix} X \\ Y \\ Z \end{bmatrix} + t$$

$$x' = x/z$$

$$y' = y/z$$

$$x'' = x' \frac{1+k_1r^2+k_2r^4+k_3r^6}{1+k_4r^2+k_5r^4+k_6r^6} + 2p_1x'y' + p_2(r^2 + 2x'^2) + s_1r^2 + s_2r^4$$

$$y'' = y' \frac{1+k_1r^2+k_2r^4+k_3r^6}{1+k_4r^2+k_5r^4+k_6r^6} + p_1(r^2 + 2y'^2) + 2p_2x'y' + s_3r^2 + s_4r^4$$

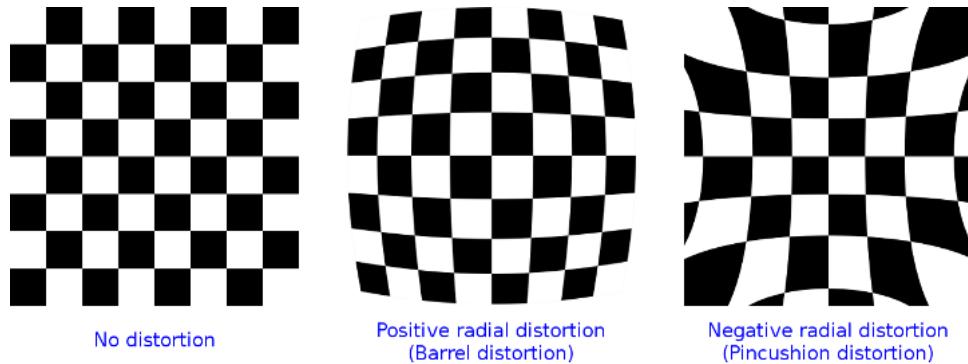
where  $r^2 = x'^2 + y'^2$

$$u = f_x * x'' + c_x$$

$$v = f_y * y'' + c_y$$

$k_1, k_2, k_3, k_4, k_5$ , and  $k_6$  are radial distortion coefficients.  $p_1$  and  $p_2$  are tangential distortion coefficients.  $s_1, s_2, s_3$ , and  $s_4$ , are the thin prism distortion coefficients. Higher-order coefficients are not considered in OpenCV.

The next figure shows two common types of radial distortion: barrel distortion (typically  $k_1 > 0$ ) and pincushion distortion (typically  $k_1 < 0$ ).



In some cases the image sensor may be tilted in order to focus an oblique plane in front of the camera (Scheimpfug condition). This can be useful for particle image velocimetry (PIV) or triangulation with a laser fan. The tilt causes a perspective distortion of  $x''$  and  $y''$ . This distortion can be modelled in the following way, see e.g. [117].

$$s \begin{bmatrix} x''' \\ y''' \\ 1 \end{bmatrix} = \begin{bmatrix} R_{33}(\tau_x, \tau_y) & 0 & -R_{13}(\tau_x, \tau_y) \\ 0 & R_{33}(\tau_x, \tau_y) & -R_{23}(\tau_x, \tau_y) \\ 0 & 0 & 1 \end{bmatrix} R(\tau_x, \tau_y) \begin{bmatrix} x'' \\ y'' \\ 1 \end{bmatrix}$$

$$u = f_x * x''' + c_x$$

$$v = f_y * y''' + c_y$$

where the matrix  $R(\tau_x, \tau_y)$  is defined by two rotations with angular parameter  $\tau_x$  and  $\tau_y$ , respectively,

$$R(\tau_x, \tau_y) = \begin{bmatrix} \cos(\tau_y) & 0 & -\sin(\tau_y) \\ 0 & 1 & 0 \\ \sin(\tau_y) & 0 & \cos(\tau_y) \end{bmatrix} \begin{bmatrix} 1 & 0 & 0 \\ 0 & \cos(\tau_x) & \sin(\tau_x) \\ 0 & -\sin(\tau_x) & \cos(\tau_x) \end{bmatrix} = \begin{bmatrix} \cos(\tau_y) & \sin(\tau_y)\sin(\tau_x) & -\sin(\tau_y)\cos(\tau_x) \\ 0 & \cos(\tau_x) & \sin(\tau_x) \\ \sin(\tau_y) & -\cos(\tau_y)\sin(\tau_x) & \cos(\tau_y)\cos(\tau_x) \end{bmatrix}.$$

In the functions below the coefficients are passed or returned as

$$(k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6[, s_1, s_2, s_3, s_4[, \tau_x, \tau_y]]]])$$

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  $320 \times 240$  resolution, absolutely the same distortion coefficients can be used for  $640 \times 480$  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.

#### Note

- A calibration sample for 3 cameras in horizontal position can be found at `opencv_source_code/samples/cpp/3calibration.cpp`
  - A calibration sample based on a sequence of images can be found at `opencv_source_code/samples/cpp/calibration.cpp`
  - A calibration sample in order to do 3D reconstruction can be found at `opencv_source_code/samples/cpp/build3dmodel.cpp`
  - A calibration sample of an artificially generated camera and chessboard patterns can be found at `opencv_source_code/samples/cpp/calibration_artificial.cpp`
  - A calibration example on stereo calibration can be found at `opencv_source_code/samples/cpp/stereo_calib.cpp`
  - A calibration example on stereo matching can be found at `opencv_source_code/samples/cpp/stereo_match.cpp`
  - (Python) A camera calibration sample can be found at `opencv_source_code/samples/python/calibrate.py`

## Enumeration Type Documentation

### § anonymous enum

anonymous enum

type of the robust estimation algorithm

Enumerator	
LMEDS Python: cv.LMEDS	least-median of squares algorithm
RANSAC Python: cv.RANSAC	RANSAC algorithm.
RHO Python: cv.RHO	RHO algorithm.

### § anonymous enum

anonymous enum

Enumerator	
SOLVEPNP_ITERATIVE Python: cv.SOLVEPNP_ITERATIVE	
SOLVEPNP_EPNP Python: cv.SOLVEPNP_EPNP	EPnP: Efficient Perspective-n-Point Camera Pose Estimation [107].
SOLVEPNP_P3P Python: cv.SOLVEPNP_P3P	Complete Solution Classification for the Perspective-Three-Point Problem [64].
SOLVEPNP_DLS Python: cv.SOLVEPNP_DLS	A Direct Least-Squares (DLS) Method for PnP [84].
SOLVEPNP_UPNP Python: cv.SOLVEPNP_UPNP	Exhaustive Linearization for Robust Camera Pose and Focal Length Estimation [148].
SOLVEPNP_AP3P Python: cv.SOLVEPNP_AP3P	An Efficient Algebraic Solution to the Perspective-Three-Point Problem [96].
SOLVEPNP_MAX_COUNT Python: cv.SOLVEPNP_MAX_COUNT	Used for count.

### § anonymous enum

## anonymous enum

## Enumerator

CALIB_CB_ADAPTIVE_THRESH	
Python: cv.CALIB_CB_ADAPTIVE_THRESH	
CALIB_CB_NORMALIZE_IMAGE	
Python: cv.CALIB_CB_NORMALIZE_IMAGE	
CALIB_CB_FILTER_QUADS	
Python: cv.CALIB_CB_FILTER_QUADS	
CALIB_CB_FAST_CHECK	
Python: cv.CALIB_CB_FAST_CHECK	

## § anonymous enum

## anonymous enum

## Enumerator

CALIB_CB_SYMMETRIC_GRID	
Python: cv.CALIB_CB_SYMMETRIC_GRID	
CALIB_CB_ASYMMETRIC_GRID	
Python: cv.CALIB_CB_ASYMMETRIC_GRID	

## § anonymous enum

## anonymous enum

Enumerator	
CALIB_USE_INTRINSIC_GUESS Python: cv.CALIB_USE_INTRINSIC_GUESS	
CALIB_FIX_ASPECT_RATIO Python: cv.CALIB_FIX_ASPECT_RATIO	
CALIB_FIX_PRINCIPAL_POINT Python: cv.CALIB_FIX_PRINCIPAL_POINT	
CALIB_ZERO_TANGENT_DIST Python: cv.CALIB_ZERO_TANGENT_DIST	
CALIB_FIX_FOCAL_LENGTH Python: cv.CALIB_FIX_FOCAL_LENGTH	
CALIB_FIX_K1 Python: cv.CALIB_FIX_K1	
CALIB_FIX_K2 Python: cv.CALIB_FIX_K2	
CALIB_FIX_K3 Python: cv.CALIB_FIX_K3	
CALIB_FIX_K4 Python: cv.CALIB_FIX_K4	
CALIB_FIX_K5 Python: cv.CALIB_FIX_K5	
CALIB_FIX_K6 Python: cv.CALIB_FIX_K6	
CALIB_RATIONAL_MODEL Python: cv.CALIB_RATIONAL_MODEL	
CALIB_THIN_PRISM_MODEL Python: cv.CALIB_THIN_PRISM_MODEL	
CALIB_FIX_S1_S2_S3_S4 Python: cv.CALIB_FIX_S1_S2_S3_S4	
CALIB_TILTED_MODEL Python: cv.CALIB_TILTED_MODEL	
CALIB_FIX_TAUX_TAUY Python: cv.CALIB_FIX_TAUX_TAUY	
CALIB_USE_QR Python: cv.CALIB_USE_QR	use QR instead of <b>SVD</b> decomposition for solving. Faster but potentially less precise
CALIB_FIX_TANGENT_DIST Python: cv.CALIB_FIX_TANGENT_DIST	
CALIB_FIX_INTRINSIC Python: cv.CALIB_FIX_INTRINSIC	
CALIB_SAME_FOCAL_LENGTH Python: cv.CALIB_SAME_FOCAL_LENGTH	
CALIB_ZERO_DISPARITY Python: cv.CALIB_ZERO_DISPARITY	
CALIB_USE_LU Python: cv.CALIB_USE_LU	use LU instead of <b>SVD</b> decomposition for solving. much faster but potentially less precise
CALIB_USE_EXTRINSIC_GUESS Python: cv.CALIB_USE_EXTRINSIC_GUESS	for stereoCalibrate

## § anonymous enum

anonymous enum

the algorithm for finding fundamental matrix

Enumerator	
FM_7POINT Python: cv.FM_7POINT	7-point algorithm
FM_8POINT Python: cv.FM_8POINT	8-point algorithm
FM_LMEDS Python: cv.FM_LMEDS	least-median algorithm. 7-point algorithm is used.
FM_RANSAC Python: cv.FM_RANSAC	RANSAC algorithm. It needs at least 15 points. 7-point algorithm is used.

## Function Documentation

[§ calibrateCamera\(\)](#) [1/2]

```

double cv::calibrateCamera ( InputArrayOfArrays objectPoints,
                           InputArrayOfArrays imagePoints,
                           Size imageSize,
                           InputOutputArray cameraMatrix,
                           InputOutputArray distCoeffs,
                           OutputArrayOfArrays rvecs,
                           OutputArrayOfArrays tvecs,
                           OutputArray stdDeviationsIntrinsics,
                           OutputArray stdDeviationsExtrinsics,
                           OutputArray perViewErrors,
                           int flags = 0,
                           TermCriteria criteria = TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, DBL_EPSILON)
)

```

**Python:**

```

retval, cameraMatrix, distCoeffs, rvecs, tvecs = cv.calibrateCamera( objectPoints, imagePoints, imageSize, cameraMatrix, distCoeffs[, rvecs[, tvecs[, flags[, criteria]]]]]

retval, cameraMatrix, distCoeffs, rvecs, tvecs, stdDeviationsIntrinsics, stdDeviationsExtrinsics, perViewErrors = cv.calibrateCameraExtended( objectPoints, imagePoints, imageSize, cameraMatrix, distCoeffs[, rvecs[, tvecs[, stdDeviationsIntrinsics[, stdDeviationsExtrinsics[, perViewErrors]]]]]

```

Finds the camera intrinsic and extrinsic parameters from several views of a calibration pattern.

**Parameters**

<b>objectPoints</b>	In the new interface it is a vector of vectors of calibration pattern points in the calibration pattern coordinate space (e.g. std::vector< 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 partially occluded patterns, or even different patterns in different views. Then, the vectors will be different. The points are 3D, but since the pattern 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 different views are concatenated together.
<b>imagePoints</b>	In the new interface it is a vector of vectors of the projections of calibration pattern points (e.g. std::vector<std::vector<cv::Vec2f>>). 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.
<b>imageSize</b>	Size of the image used only to initialize the intrinsic camera matrix.
<b>cameraMatrix</b>	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 CALIB_FIX_ASPECT_RATIO are set, the matrix must be initialized before calling the function.
<b>distCoeffs</b>	Output vector of distortion coefficients ( $k_1, k_2, p_1, p_2, [k_3, k_4, k_5, k_6, s_1, s_2, s_3, s_4, [\tau_x, \tau_y]]$ ) of 4, 5, 8, 12 or 14 elements.
<b>rvecs</b>	Output vector of rotation vectors (see Rodrigues ) estimated for each pattern view (e.g. std::vector<cv::Mat>>). That is, each k-th rotation vector (see the next output parameter description) brings the calibration pattern from the model coordinate space (in which the pattern is located, that is, a real position of the calibration pattern in the k-th pattern view (k=0.. M-1).
<b>tvecs</b>	Output vector of translation vectors estimated for each pattern view.
<b>stdDeviationsIntrinsics</b>	Output vector of standard deviations estimated for intrinsic parameters. Order of deviations values: ( $f_x, f_y, c_x, c_y, k_1, k_2, p_1, p_2, [s_1, s_2, s_3, s_4]$ ). If the parameter is not estimated, its deviation is equals to zero.
<b>stdDeviationsExtrinsics</b>	Output vector of standard deviations estimated for extrinsic parameters. Order of deviations values: ( $R_1, T_1, \dots, R_M, T_M$ ) where concatenated 1x3 vectors.
<b>perViewErrors</b>	Output vector of the RMS re-projection error estimated for each pattern view.
<b>flags</b>	Different flags that may be zero or a combination of the following values: <ul style="list-style-type: none"> <li>• <b>CALIB_USE_INTRINSIC_GUESS</b> cameraMatrix contains valid initial values of fx, fy, cx, cy that are optimized further. Otherwise, the imageSize is used, and focal distances are computed in a least-squares fashion. Note, that if intrinsic parameters are known, set the flag to zero.</li> <li>• <b>CALIB_FIX_PRINCIPAL_POINT</b> The principal point is not changed during the global optimization. It stays at the center or at the center of the image if CALIB_USE_INTRINSIC_GUESS is set too.</li> <li>• <b>CALIB_FIX_ASPECT_RATIO</b> The functions considers only fy as a free parameter. The ratio fx/fy stays the same as in the intrinsic matrix. If CALIB_USE_INTRINSIC_GUESS is not set, the actual input values of fx and fy are ignored, only their ratio is computed and used.</li> <li>• <b>CALIB_ZERO_TANGENT_DIST</b> Tangential distortion coefficients (<math>p_1, p_2</math>) are set to zeros and stay zero.</li> <li>• <b>CALIB_FIX_K1,...,CALIB_FIX_K6</b> The corresponding radial distortion coefficient is not changed during the optimization. If CALIB_FIX_K1 is set, the supplied distCoeffs matrix is used. Otherwise, it is set to 0.</li> <li>• <b>CALIB_RATIONAL_MODEL</b> Coefficients k4, k5, and k6 are enabled. To provide the backward compatibility, this extra flag is used. If the flag is not set, the function use the rational model and return 8 coefficients. If the flag is set, the function computes and returns only 5 distortion coefficients.</li> <li>• <b>CALIB_THIN_PRISM_MODEL</b> Coefficients s1, s2, s3 and s4 are enabled. To provide the backward compatibility, this extra flag is used. If the flag is not set, the function use the thin prism model and return 12 coefficients. If the flag is set, the function computes and returns only 5 distortion coefficients.</li> <li>• <b>CALIB_FIX_S1_S2_S3_S4</b> The thin prism distortion coefficients are not changed during the optimization. If CALIB_USE_INTRINSIC_GUESS is set, the supplied distCoeffs matrix is used. Otherwise, it is set to 0.</li> </ul>

- **CALIB\_TILTED\_MODEL** Coefficients tauX and tauY are enabled. To provide the backward compatibility, this extra flag should use the tilted sensor model and return 14 coefficients. If the flag is not set, the function computes and returns only 5 distortion coefficients.
- **CALIB\_FIX\_TAUX\_TAUY** The coefficients of the tilted sensor model are not changed during the optimization. If CALIB\_USE\_INTRINSIC\_GUESS is supplied distCoeffs matrix is used. Otherwise, it is set to 0.

**criteria**

Termination criteria for the iterative optimization algorithm.

**Returns**

the overall RMS re-projection error.

The function estimates the intrinsic camera parameters and extrinsic parameters for each of the views. The algorithm is based on [223] and [21]. The coordinates of object points 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 (see findChessboardCorners). Currently, initialization of intrinsic parameters (when CALIB\_USE\_INTRINSIC\_GUESS is set) is supported only 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:

- Compute the initial intrinsic parameters (the option only available for planar calibration patterns) or read them from the input parameters. The distortion coefficients  $\kappa_1, \kappa_2, p_1, p_2$  and  $\text{CALIB_FIX_K}_1, \text{CALIB_FIX_K}_2$  are specified.
- Estimate the initial camera pose as if the intrinsic parameters have been already known. This is done using solvePnP.
- Run the global Levenberg-Marquardt optimization algorithm to minimize the reprojection error, that is, the total sum of squared distances between the observed object points and the current estimates for camera parameters and the poses) object points objectPoints. See projectPoints for details.

**Note**

If you use a non-square (=non-NxN) grid and findChessboardCorners for calibration, and calibrateCamera returns bad values (zero distortion coefficients, an invalid value for  $\text{CALIB_OK}$ , 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).

**See also**

[findChessboardCorners](#), [solvePnP](#), [initCameraMatrix2D](#), [stereoCalibrate](#), [undistort](#)

**§ [calibrateCamera\(\)](#) [2/2]**

```
double cv::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:**

```
retval, cameraMatrix, distCoeffs, rvecs, tvecs = cv.calibrateCamera(objectPoints, imagePoints, imageSize, cameraMatrix, distCoeffs[, rvecs[, tvecs[, flags[, criteria]]]]]
retval, cameraMatrix, distCoeffs, rvecs, tvecs, stdDeviationsIntrinsics, stdDeviationsExtrinsics, perViewErrors = cv.calibrateCameraExtended( objectPoints, imagePoints, imageSize, cameraMatrix, distCoeffs[, rvecs[, tvecs[, stdDeviationsIntrinsics[, stdDeviationsExtrinsics[, perViewErrors]]]]])
```

**§ [calibrationMatrixValues\(\)](#)**

```
void cv::calibrationMatrixValues ( InputArray cameraMatrix,
                                  Size      imageSize,
                                  double    apertureWidth,
                                  double    apertureHeight,
                                  double &   fovx,
                                  double &   fovy,
                                  double &   focalLength,
                                  Point2d &  principalPoint,
                                  double &   aspectRatio
                                )
```

**Python:**

```
fovx, fovy, focalLength, principalPoint, aspectRatio = cv.calibrationMatrixValues( cameraMatrix, imageSize, apertureWidth, apertureHeight )
```

Computes useful camera characteristics from the camera matrix.

**Parameters**

**cameraMatrix** Input camera matrix that can be estimated by `calibrateCamera` or `stereoCalibrate` .  
**imageSize** Input image size in pixels.  
**apertureWidth** Physical width in mm of the sensor.  
**apertureHeight** Physical height in mm 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 mm.  
**aspectRatio**  $f_y/f_x$

The function computes various useful camera characteristics from the previously estimated camera matrix.

**Note**

Do keep in mind that the unity measure 'mm' stands for whatever unit of measure one chooses for the chessboard pitch (it can thus be any value).

**§ composeRT()**

```

void cv::composeRT ( InputArray rvec1,
                    InputArray tvec1,
                    InputArray rvec2,
                    InputArray tvec2,
                    OutputArray rvec3,
                    OutputArray tvec3,
                    dr3dr1 =
                    OutputArray noArray(),
                    dr3dt1 =
                    OutputArray noArray(),
                    dr3dr2 =
                    OutputArray noArray(),
                    dr3dt2 =
                    OutputArray noArray(),
                    dt3dr1 =
                    OutputArray noArray(),
                    dt3dt1 =
                    OutputArray noArray(),
                    dt3dr2 =
                    OutputArray noArray(),
                    OutputArray dt3dt2 = noArray()
)

```

**Python:**

```

rvec3, tvec3,
dr3dr1, dr3dt1,
dr3dr2, dr3dt2,     = cv.composeRT( rvec1, tvec1, rvec2, tvec2[, rvec3[, tvec3[, dr3dr1[, dr3dt1[, dr3dr2[, dr3dt2[, dt3dr1[, dt3dt1[, dt3dr2[, dt3dt2]]]]]]]]])
dt3dr1, dt3dt1,
dt3dr2, dt3dt2

```

Combines two rotation-and-shift transformations.

**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.
- dr3dr1**
- dr3dt1**
- dr3dr2**
- dr3dt2**
- dt3dr1**
- dt3dt1**
- dt3dr2**
- dt3dt2** Optional output derivatives of rvec3 or tvec3 with regard to rvec1, rvec2, tvec1 and tvec2, respectively.

The functions compute:

$$\begin{aligned} \text{rvec3} &= \text{rodrigues}^{-1}(\text{rodrigues}(\text{rvec2}) \cdot \text{rodrigues}(\text{rvec1})) \\ \text{tvec3} &= \text{rodrigues}(\text{rvec2}) \cdot \text{tvec1} + \text{tvec2} \end{aligned}$$

where `rodrigues` denotes a rotation vector to a rotation matrix transformation, and `rodrigues`<sup>-1</sup> 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()

```
void cv::computeCorrespondEpilines ( InputArray points,
                                    int whichImage,
                                    InputArray F,
                                    OutputArray lines
                                )
```

**Python:**

```
lines = cv.computeCorrespondEpilines( points, whichImage, F[, lines] )
```

For points in an image of a stereo pair, computes the corresponding epilines in the other image.

**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 which Image=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  $a_i^2 + b_i^2 = 1$  .

**§ convertPointsFromHomogeneous()**

```
void cv::convertPointsFromHomogeneous ( InputArray src,
                                         OutputArray dst
                                       )
```

**Python:**

```
dst = cv.convertPointsFromHomogeneous( src[, dst] )
```

Converts points from homogeneous to Euclidean space.

**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  $(x_1, x_2, \dots, x_{n-1}, x_n)$  is converted to  $(x_1/x_n, x_2/x_n, \dots, x_{n-1}/x_n)$ . When  $x_n=0$ , the output point coordinates will be  $(0,0,0,\dots)$ .

**§ convertPointsHomogeneous()**

```
void cv::convertPointsHomogeneous ( InputArray src,
                                    OutputArray dst
                                  )
```

Converts points to/from homogeneous coordinates.

**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.

## § convertPointsToHomogeneous()

```
void cv::convertPointsToHomogeneous ( InputArray src,
                                     OutputArray dst
                                    )
```

### Python:

```
dst = cv.convertPointsToHomogeneous( src[, dst] )
```

Converts points from Euclidean to homogeneous space.

#### 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  $(x_1, x_2, \dots, x_n)$  is converted to  $(x_1, x_2, \dots, x_n, 1)$ .

## § correctMatches()

```
void cv::correctMatches ( InputArray F,
                         InputArray points1,
                         InputArray points2,
                         OutputArray newPoints1,
                         OutputArray newPoints2
                        )
```

### Python:

```
newPoints1, newPoints2 = cv.correctMatches( F, points1, points2[, newPoints1[, newPoints2]] )
```

Refines coordinates of corresponding points.

#### 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  $\text{points1}[i] \leftrightarrow \text{points2}[i]$ , and a fundamental matrix  $F$ , it computes the corrected correspondences  $\text{newPoints1}[i] \leftrightarrow \text{newPoints2}[i]$  that minimize the geometric error  $d(\text{points1}[i], \text{newPoints1}[i])^2 + d(\text{points2}[i], \text{newPoints2}[i])^2$  (where  $d(a, b)$  is the geometric distance between points  $a$  and  $b$ ) subject to the epipolar constraint  $\text{newPoints2}^T * F * \text{newPoints1} = 0$ .

## § decomposeEssentialMat()

```
void cv::decomposeEssentialMat ( InputArray E,
                                 OutputArray R1,
                                 OutputArray R2,
                                 OutputArray t
                                )
```

### Python:

```
R1, R2, t = cv.decomposeEssentialMat( E[, R1[, R2[, t]]] )
```

Decompose an essential matrix to possible rotations and translation.

#### Parameters

**E** The input essential matrix.

**R1** One possible rotation matrix.

**R2** Another possible rotation matrix.

**t** One possible translation.

This function decomposes an essential matrix  $E$  using svd decomposition [77]. Generally 4 possible poses exist for a given  $E$ . They are  $[R_1, t]$ ,  $[R_1, -t]$ ,  $[R_2, t]$ ,  $[R_2, -t]$ . By decomposing  $E$ , you can only get the direction of the translation, so the function returns unit  $t$ .

## § decomposeHomographyMat()

```
int cv::decomposeHomographyMat ( InputArray H,  
                                InputArray K,  
                                OutputArrayOfArrays rotations,  
                                OutputArrayOfArrays translations,  
                                OutputArrayOfArrays normals  
)
```

### Python:

```
retval, rotations, translations, normals = cv.decomposeHomographyMat( H, K[, rotations[, translations[, normals]]] )
```

Decompose a homography matrix to rotation(s), translation(s) and plane normal(s).

#### Parameters

- H** The input homography matrix between two images.
- K** The input intrinsic camera calibration matrix.
- rotations** Array of rotation matrices.
- translations** Array of translation matrices.
- normals** Array of plane normal matrices.

This function extracts relative camera motion between two views observing a planar object from the homography  $H$  induced by the plane. The intrinsic camera matrix  $K$  must also be provided. The function may return up to four mathematical solution sets. At least two of the solutions may further be invalidated if point correspondences are available by applying positive depth constraint (all points must be in front of the camera). The decomposition method is described in detail in [\[124\]](#).

#### Examples:

[samples/cpp/tutorial\\_code/features2D/Homography/decompose\\_homography.cpp](#).

## § decomposeProjectionMatrix()

```
void cv::decomposeProjectionMatrix ( InputArray projMatrix,
                                    OutputArray cameraMatrix,
                                    OutputArray rotMatrix,
                                    OutputArray transVect,
                                    OutputArray rotMatrixX = noArray(),
                                    OutputArray rotMatrixY = noArray(),
                                    OutputArray rotMatrixZ = noArray(),
                                    OutputArray eulerAngles = noArray()
                                )
```

**Python:**

```
cameraMatrix,
rotMatrix,
transVect,
rotMatrixX,    = cv.decomposeProjectionMatrix( projMatrix[, cameraMatrix[, rotMatrix[, transVect[, rotMatrixX[, rotMatrixY[, rotMatrixZ[, eulerAngles]]]]]]] )
rotMatrixY,
rotMatrixZ,
eulerAngles
```

Decomposes a projection matrix into a rotation matrix and a camera matrix.

**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.
- rotMatrixX** Optional 3x3 rotation matrix around x-axis.
- rotMatrixY** Optional 3x3 rotation matrix around y-axis.
- rotMatrixZ** 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 principal axes that results in the same orientation of an object, e.g. see [173]. Returned tree rotation matrices and corresponding three Euler angles are only one of the possible solutions.

The function is based on RQDecomp3x3.

**drawChessboardCorners()**

```
void cv::drawChessboardCorners ( InputOutputArray image,
                               Size patternSize,
                               InputArray corners,
                               bool patternWasFound
                            )
```

**Python:**

```
image = cv.drawChessboardCorners( image, patternSize, corners, patternWasFound )
```

Renders the detected chessboard corners.

**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.

**Examples:**

[samples/cpp/tutorial\\_code/features2D/Homography/pose\\_from\\_homography.cpp](#).

**estimateAffine2D()**

```
cv::Mat cv::estimateAffine2D ( InputArray from,
                               InputArray to,
                               OutputArray inliers = noArray(),
                               int method = RANSAC,
                               ransacReprojThreshold =
                               3,
                               size_t maxIters = 2000,
                               double confidence = 0.99,
                               size_t refinelters = 10
                           )
```

**Python:**

```
retval, inliers = cv.estimateAffine2D( from, to[, inliers[, method[, ransacReprojThreshold[, maxIters[, confidence[, refinelters]]]]]])
```

Computes an optimal affine transformation between two 2D point sets.

It computes

$$\begin{bmatrix} x \\ y \end{bmatrix} = \begin{bmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{bmatrix} \begin{bmatrix} X \\ Y \end{bmatrix} + \begin{bmatrix} b_1 \\ b_2 \end{bmatrix}$$

**Parameters**

- from** First input 2D point set containing  $(X, Y)$ .
- to** Second input 2D point set containing  $(x, y)$ .
- inliers** Output vector indicating which points are inliers (1-inlier, 0-outlier).
- method** Robust method used to compute transformation. The following methods are possible:
  - **cv::RANSAC** - RANSAC-based robust method
  - **cv::LMEADS** - Least-Median robust method RANSAC is the default method.

**ransacReprojThreshold** Maximum reprojection error in the RANSAC algorithm to consider a point as an inlier. Applies only to RANSAC.

**maxIters** The maximum number of robust method iterations.

**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.

**refinelters** Maximum number of iterations of refining algorithm (Levenberg-Marquardt). Passing 0 will disable refining, so the output matrix will be output of robust method.

**Returns**

Output 2D affine transformation matrix  $2 \times 3$  or empty matrix if transformation could not be estimated. The returned matrix has the following form:

$$\begin{bmatrix} a_{11} & a_{12} & b_1 \\ a_{21} & a_{22} & b_2 \end{bmatrix}$$

The function estimates an optimal 2D affine transformation between two 2D point sets using the selected robust algorithm.

The computed transformation is then refined further (using only inliers) with the Levenberg-Marquardt method to reduce the re-projection error even more.

**Note**

The RANSAC method can handle practically any ratio of outliers but 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.

**See also**

[estimateAffinePartial2D](#), [getAffineTransform](#)

**§ [estimateAffine3D\(\)](#)**

```
int cv::estimateAffine3D ( InputArray src,
                           InputArray dst,
                           OutputArray out,
                           OutputArray inliers,
                           double ransacThreshold =
                           double 3,
                           double confidence = 0.99
                           )
```

**Python:**

```
retval, out, inliers = cv.estimateAffine3D( src, dst[, out[, inliers[, ransacThreshold[, confidence]]]] )
```

Computes an optimal affine transformation between two 3D point sets.

It computes

$$\begin{bmatrix} x \\ y \\ z \end{bmatrix} = \begin{bmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{33} \end{bmatrix} \begin{bmatrix} X \\ Y \\ Z \end{bmatrix} + \begin{bmatrix} b_1 \\ b_2 \\ b_3 \end{bmatrix}$$

**Parameters**

**src** First input 3D point set containing  $(X, Y, Z)$ .

**dst** Second input 3D point set containing  $(x, y, z)$ .

**out** Output 3D affine transformation matrix  $3 \times 4$  of the form

$$\begin{bmatrix} a_{11} & a_{12} & a_{13} & b_1 \\ a_{21} & a_{22} & a_{23} & b_2 \\ a_{31} & a_{32} & a_{33} & b_3 \end{bmatrix}$$

**inliers** Output vector indicating which points are inliers (1-inlier, 0-outlier).

**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.

## § estimateAffinePartial2D()

```
cv::Mat cv::estimateAffinePartial2D ( InputArray from,
                                      InputArray to,
                                      OutputArray inliers = noArray(),
                                      int method = RANSAC,
                                      ransacReprojThreshold =
                                      double 3,
                                      size_t maxIters = 2000,
                                      double confidence = 0.99,
                                      size_t refinIters = 10
) 
```

**Python:**

```
retval, inliers = cv.estimateAffinePartial2D( from, to[, inliers[, method[, ransacReprojThreshold[, maxIters[, confidence[, refinIters]]]]]]) 
```

Computes an optimal limited affine transformation with 4 degrees of freedom between two 2D point sets.

#### Parameters

<b>from</b>	First input 2D point set.
<b>to</b>	Second input 2D point set.
<b>inliers</b>	Output vector indicating which points are inliers.
<b>method</b>	Robust method used to compute transformation. The following methods are possible: <ul style="list-style-type: none"> <li>• <b>cv::RANSAC</b> - RANSAC-based robust method</li> <li>• <b>cv::LMEDE</b> - Least-Median robust method RANSAC is the default method.</li> </ul>
<b>ransacReprojThreshold</b>	Maximum reprojection error in the RANSAC algorithm to consider a point as an inlier. Applies only to RANSAC.
<b>maxIters</b>	The maximum number of robust method iterations.
<b>confidence</b>	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.
<b>refinIters</b>	Maximum number of iterations of refining algorithm (Levenberg-Marquardt). Passing 0 will disable refining, so the output matrix will be output of robust method.

#### Returns

Output 2D affine transformation (4 degrees of freedom) matrix  $2 \times 3$  or empty matrix if transformation could not be estimated.

The function estimates an optimal 2D affine transformation with 4 degrees of freedom limited to combinations of translation, rotation, and uniform scaling. Uses the selected algorithm for robust estimation.

The computed transformation is then refined further (using only inliers) with the Levenberg-Marquardt method to reduce the re-projection error even more.

Estimated transformation matrix is:

$$\begin{bmatrix} \cos(\theta) \cdot s & -\sin(\theta) \cdot s & t_x \\ \sin(\theta) \cdot s & \cos(\theta) \cdot s & t_y \end{bmatrix}$$

Where  $\theta$  is the rotation angle,  $s$  the scaling factor and  $t_x, t_y$  are translations in  $x, y$  axes respectively.

#### Note

The RANSAC method can handle practically any ratio of outliers but need 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.

#### See also

[estimateAffine2D](#), [getAffineTransform](#)

§ [filterHomographyDecompByVisibleRefpoints\(\)](#)

```
void cv::filterHomographyDecompByVisibleRefpoints ( InputArrayOfArrays rotations,
                                                    InputArrayOfArrays normals,
                                                    InputArray beforePoints,
                                                    InputArray afterPoints,
                                                    OutputArray possibleSolutions,
                                                    InputArray pointsMask = noArray()
)
```

**Python:**

```
possibleSolutions = cv.filterHomographyDecompByVisibleRefpoints( rotations, normals, beforePoints, afterPoints[, possibleSolutions[, pointsMask]] )
```

Filters homography decompositions based on additional information.

**Parameters**

<b>rotations</b>	Vector of rotation matrices.
<b>normals</b>	Vector of plane normal matrices.
<b>beforePoints</b>	Vector of (rectified) visible reference points before the homography is applied
<b>afterPoints</b>	Vector of (rectified) visible reference points after the homography is applied
<b>possibleSolutions</b>	Vector of int indices representing the viable solution set after filtering
<b>pointsMask</b>	optional Mat/Vector of 8u type representing the mask for the inliers as given by the findHomography function

This function is intended to filter the output of the decomposeHomographyMat based on additional information as described in [124]. The summary of the method: the decomposeHomographyMat function returns 2 unique solutions and their "opposites" for a total of 4 solutions. If we have access to the sets of points visible in the camera frame before and after the homography transformation is applied, we can determine which are the true potential solutions and which are the opposites by verifying which homographies are consistent with all visible reference points being in front of the camera. The inputs are left unchanged; the filtered solution set is returned as indices into the existing one.

**§ filterSpeckles()**

```
void cv::filterSpeckles ( InputOutputArray img,
                        double newVal,
                        int maxSpeckleSize,
                        double maxDiff,
                        InputOutputArray buf = noArray()
)
```

**Python:**

```
img, buf = cv.filterSpeckles( img, newVal, maxSpeckleSize, maxDiff[, buf] )
```

Filters off small noise blobs (speckles) in the disparity map.

**Parameters**

<b>img</b>	The input 16-bit signed disparity image
<b>newVal</b>	The disparity value used to paint-off the speckles
<b>maxSpeckleSize</b>	The maximum speckle size to consider it a speckle. Larger blobs are not affected by the algorithm
<b>maxDiff</b>	Maximum difference between neighbor disparity pixels to put them into the same blob. Note that since <b>StereoBM</b> , <b>StereoSGBM</b> 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.
<b>buf</b>	The optional temporary buffer to avoid memory allocation within the function.

**§ find4QuadCornerSubpix()**

```
bool cv::find4QuadCornerSubpix ( InputArray img,
                                 InputOutputArray corners,
                                 Size region_size
)
```

finds subpixel-accurate positions of the chessboard corners

**§ findChessboardCorners()**

```
bool cv::findChessboardCorners ( InputArray image,
                                Size patternSize,
                                OutputArray corners,
                                int flags = CALIB_CB_ADAPTIVE_THRESH+CALIB_CB_NORMALIZE_IMAGE
                            )
```

**Python:**

```
retval, corners = cv.findChessboardCorners( image, patternSize[, corners[, flags]] )
```

Finds the positions of internal corners of the chessboard.

**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\_col) = cvSize(columns,rows) ).

**corners** Output array of detected corners.

**flags** Various operation flags that can be zero or a combination of the following values:

- **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).
- **CALIB\_CB\_NORMALIZE\_IMAGE** Normalize the image gamma with equalizeHist before applying fixed or adaptive thresholding.
- **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.

**Examples:**

[samples/cpp/tutorial\\_code/features2D/Homography/decompose\\_homography.cpp](#),  
[samples/cpp/tutorial\\_code/features2D/Homography/homography\\_from\\_camera\\_displacement.cpp](#), and  
[samples/cpp/tutorial\\_code/features2D/Homography/pose\\_from\\_homography.cpp](#).

**§ [findCirclesGrid\(\)](#) [1/2]**

```
bool cv::findCirclesGrid ( InputArray image,
                           Size patternSize,
                           OutputArray centers,
                           int flags,
                           const Ptr< FeatureDetector > & blobDetector,
                           CirclesGridFinderParameters parameters
                         )
```

**Python:**

```
retval, centers = cv.findCirclesGrid( image, patternSize, flags, blobDetector, parameters[, centers] )
retval, centers = cv.findCirclesGrid( image, patternSize[, centers[, flags[, blobDetector]]] )
```

Finds centers in the grid of circles.

**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_column)` ).
- 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.

**parameters** struct for finding circles in a grid pattern.

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.

**§ findCirclesGrid() [2/2]**

```
bool cv::findCirclesGrid ( InputArray image,
                           Size patternSize,
                           OutputArray centers,
                           int flags = CALIB_CB_SYMMETRIC_GRID,
                           const Ptr< FeatureDetector > & blobDetector = SimpleBlobDetector::create()
                         )
```

**Python:**

```
retval, centers = cv.findCirclesGrid( image, patternSize, flags, blobDetector, parameters[, centers] )
retval, centers = cv.findCirclesGrid( image, patternSize[, centers[, flags[, blobDetector]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**§ findCirclesGrid2()**

```
bool cv::findCirclesGrid2 ( InputArray
                           Size
                           OutputArray
                           int
                           const Ptr< FeatureDetector > & blobDetector,
                           CirclesGridFinderParameters2 parameters
                           )
```

**Python:**

```
retval, centers = cv.findCirclesGrid2( image, patternSize, flags, blobDetector, parameters[, centers] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**§ findEssentialMat() [1/2]**

```
Mat cv::findEssentialMat ( InputArray points1,
                           InputArray points2,
                           InputArray cameraMatrix,
                           int method = RANSAC,
                           double prob = 0.999,
                           double threshold = 1.0,
                           OutputArray mask = noArray()
                           )
```

**Python:**

```
retval, mask = cv.findEssentialMat( points1, points2, cameraMatrix[, method[, prob[, threshold[, mask]]]] )
retval, mask = cv.findEssentialMat( points1, points2[, focal[, pp[, method[, prob[, threshold[, mask]]]]]] )
```

Calculates an essential matrix from the corresponding points in two images.

**Parameters**

- points1** Array of N (N >= 5) 2D 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 .
- cameraMatrix** Camera matrix  $K = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ . Note that this function assumes that points1 and points2 are feature points from cameras with the same camera matrix.
- method** Method for computing an essential matrix.
  - **RANSAC** for the RANSAC algorithm.
  - **LMEDS** for the LMedS algorithm.
- prob** Parameter used for the RANSAC or LMedS methods only. It specifies a desirable level of confidence (probability) that the estimated matrix is correct.
- threshold** 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.
- mask** 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.

This function estimates essential matrix based on the five-point algorithm solver in [144] . [174] is also a related. The epipolar geometry is described by the following equation:

$$[p_2; 1]^T K^{-T} E K^{-1} [p_1; 1] = 0$$

where  $E$  is an essential matrix,  $p_1$  and  $p_2$  are corresponding points in the first and the second images, respectively. The result of this function may be passed further to decomposeEssentialMat or recoverPose to recover the relative pose between cameras.

**§ findEssentialMat() [2/2]**

```
Mat cv::findEssentialMat ( InputArray points1,
                           InputArray points2,
                           double focal = 1.0,
                           Point2d pp =
                           Point2d(0, 0),
                           int method = RANSAC,
                           double prob = 0.999,
                           double threshold = 1.0,
                           OutputArray mask = noArray()
)
```

**Python:**

```
retval, mask = cv.findEssentialMat( points1, points2, cameraMatrix[, method[, prob[, threshold[, mask]]]] )
retval, mask = cv.findEssentialMat( points1, points2[, focal[, pp[, method[, prob[, threshold[, mask]]]]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**Parameters**

**points1** Array of N (N >= 5) 2D 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 .

**focal** focal length of the camera. Note that this function assumes that points1 and points2 are feature points from cameras with same focal length and principal point.

**pp** principal point of the camera.

**method** Method for computing a fundamental matrix.

- **RANSAC** for the RANSAC algorithm.
- **LMEDS** for the LMedS algorithm.

**threshold** 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.

**prob** Parameter used for the RANSAC or LMedS methods only. It specifies a desirable level of confidence (probability) that the estimated matrix is correct.

**mask** 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.

This function differs from the one above that it computes camera matrix from focal length and principal point:

$$K = \begin{bmatrix} f & 0 & x_{pp} \\ 0 & f & y_{pp} \\ 0 & 0 & 1 \end{bmatrix}$$

**§ findFundamentalMat() [1/2]**

```
Mat cv::findFundamentalMat ( InputArray points1,
                            InputArray points2,
                            int method = FM_RANSAC,
                            ransacReprojThreshold =
                            double 3.,
                            double confidence = 0.99,
                            OutputArray mask = noArray()
)
```

**Python:**

```
retval, mask = cv.findFundamentalMat( points1, points2[, method[, ransacReprojThreshold[, confidence[, mask]]]] )
```

Calculates a fundamental matrix from the corresponding points in two images.

**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.  $N \geq 8$

**ransacReprojThreshold** Parameter used only 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.

**confidence** Parameter used for the RANSAC and LMedS methods only. It specifies a desirable level of confidence (probability) that the estimated matrix is correct.

**mask** 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);
```

**§ findFundamentalMat() [2/2]**

```
Mat cv::findFundamentalMat ( InputArray points1,
                            InputArray points2,
                            OutputArray mask,
                            int method = FM_RANSAC,
                            ransacReprojThreshold =
                            double 3.,
                            double confidence = 0.99
)
```

**Python:**

```
retval, mask = cv.findFundamentalMat( points1, points2[, method[, ransacReprojThreshold[, confidence[, mask]]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**§ findHomography() [1/2]**

```
Mat cv::findHomography ( InputArray srcPoints,
                        InputArray dstPoints,
                        int method = 0,
                        ransacReprojThreshold =
                        double 3,
                        OutputArray mask = noArray(),
                        const int maxIters = 2000,
                        const double confidence = 0.995
)
```

**Python:**

```
retval, mask = cv.findHomography( srcPoints, dstPoints[, method[, ransacReprojThreshold[, mask[, maxIters[, confidence]]]]])
```

Finds a perspective transformation between two planes.

**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 compute a homography matrix. The following methods are possible:

- **0** - a regular method using all the points, i.e., the least squares method
- **RANSAC** - RANSAC-based robust method
- **LMEDS** - Least-Median robust method
- **RHO** - PROSAC-based robust method

**ransacReprojThreshold** Maximum allowed reprojection error to treat a point pair as an inlier (used in the RANSAC and RHO methods only). That is, if

$$\|dstPoints_i - convertPointsHomogeneous(H * srcPoints_i)\|_2 > ransacReprojThreshold$$

then the point  $i$  is considered as 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 ( RANSAC or LMEDS ). Note that the input mask values are ignored.

**maxIters** The maximum number of RANSAC iterations.

**confidence** Confidence level, between 0 and 1.

The function finds and returns 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 ( $srcPoints_i, dstPoints_i$ ) 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 three robust methods. The methods RANSAC, LMEDS and RHO try many different random subsets of the corresponding point pairs (of four pairs each, collinear pairs are discarded), estimate the homography matrix using this subset and a simple least-squares algorithm, and then compute the quality/goodness of the computed homography (which is the number of inliers for RANSAC or the least median re-projection 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 methods RANSAC and RHO can handle practically any ratio of outliers but need 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  $h_{33} = 1$ . Note that whenever an  $H$  matrix cannot be estimated, an empty one will be returned.

**See also**

[getAffineTransform](#), [estimateAffine2D](#), [estimateAffinePartial2D](#), [getPerspectiveTransform](#), [warpPerspective](#), [perspectiveTransform](#)

**Examples:**

[samples/cpp/tutorial\\_code/features2D/Homography/decompose\\_homography.cpp](#),  
[samples/cpp/tutorial\\_code/features2D/Homography/homography\\_from\\_camera\\_displacement.cpp](#),

[samples/cpp/tutorial\\_code/features2D/Homography/pose\\_from\\_homography.cpp](#), and [samples/cpp/warpPerspective\\_demo.cpp](#).

## § [findHomography\(\)](#) [2/2]

```
Mat cv::findHomography ( InputArray srcPoints,
                        InputArray dstPoints,
                        OutputArray mask,
                        int method = 0,
                        double ransacReprojThreshold = 3
)
```

### Python:

```
retval, mask = cv.findHomography( srcPoints, dstPoints[, method[, ransacReprojThreshold[, mask[, maxIters[, confidence]]]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

## § [getOptimalNewCameraMatrix\(\)](#)

```
Mat cv::getOptimalNewCameraMatrix ( InputArray cameraMatrix,
                                    InputArray distCoeffs,
                                    Size imageSize,
                                    double alpha,
                                    Size newImgSize = Size(),
                                    Rect * validPixROI = 0,
                                    bool centerPrincipalPoint = false
)
```

### Python:

```
retval, validPixROI = cv.getOptimalNewCameraMatrix( cameraMatrix, distCoeffs, imageSize, alpha[, newImgSize[, centerPrincipalPoint]] )
```

Returns the new camera matrix based on the free scaling parameter.

### Parameters

<b>cameraMatrix</b>	Input camera matrix.
<b>distCoeffs</b>	Input vector of distortion coefficients ( $k_1, k_2, p_1, p_2, [k_3, k_4, k_5, k_6, s_1, s_2, s_3, s_4, \tau_x, \tau_y]$ ) of 4, 5, 8, 12 or 14 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
<b>imageSize</b>	Original image size.
<b>alpha</b>	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.
<b>newImgSize</b>	Image size after rectification. By default, it is set to imageSize .
<b>validPixROI</b>	Optional output rectangle that outlines all-good-pixels region in the undistorted image. See roi1, roi2 description in stereoRectify .
<b>centerPrincipalPoint</b>	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.

### Returns

`new_camera_matrix` Output new camera matrix.

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 undistorted 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 .

## § [getValidDisparityROI\(\)](#)

```
Rect cv::getValidDisparityROI ( Rect roi1,
                               Rect roi2,
                               int minDisparity,
                               int numberOfDisparities,
                               int SADWindowSize
                             )
```

**Python:**

```
retval = cv.getValidDisparityROI( roi1, roi2, minDisparity, numberOfDisparities, SADWindowSize )
```

computes valid disparity ROI from the valid ROIs of the rectified images (that are returned by `cv::stereoRectify()`)

**§ initCameraMatrix2D()**

```
Mat cv::initCameraMatrix2D ( InputArrayOfArrays objectPoints,
                           InputArrayOfArrays imagePoints,
                           Size imageSize,
                           double aspectRatio = 1.0
                         )
```

**Python:**

```
retval = cv.initCameraMatrix2D( objectPoints, imagePoints, imageSize[, aspectRatio] )
```

Finds an initial camera matrix from 3D-2D point correspondences.

**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.
- 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,  $f_x = f_y * \text{aspectRatio}$ .

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()**

```
void cv::matMulDeriv ( InputArray A,
                      InputArray B,
                      OutputArray dABdA,
                      OutputArray dABdB
                    )
```

**Python:**

```
dABdA, dABdB = cv.matMulDeriv( A, B[, dABdA[, dABdB]] )
```

Computes partial derivatives of the matrix product for each multiplied matrix.

**Parameters**

- A** First multiplied matrix.
- B** Second multiplied matrix.
- dABdA** First output derivative matrix  $d(A*B)/dA$  of size  $A.\text{rows} * B.\text{cols} \times A.\text{rows} * A.\text{cols}$ .
- dABdB** Second output derivative matrix  $d(A*B)/dB$  of size  $A.\text{rows} * B.\text{cols} \times B.\text{rows} * B.\text{cols}$ .

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()**

```
void cv::projectPoints ( InputArray objectPoints,
                        InputArray rvec,
                        InputArray tvec,
                        InputArray cameraMatrix,
                        InputArray distCoeffs,
                        OutputArray imagePoints,
                        jacobian =
                        OutputArray noArray(),
                        double aspectRatio = 0
)
```

**Python:**

```
imagePoints, jacobian = cv.projectPoints( objectPoints, rvec, tvec, cameraMatrix, distCoeffs[, imagePoints[, jacobian[, aspectRatio]]] )
```

Projects 3D points to an image plane.

**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 ( $k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6[, s_1, s_2, s_3, s_4[, \tau_x, \tau_y]]]]$ ) of 4, 5, 8, 12 or 14 elements. If the vector is 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 ( $f_x/f_y$ ) 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.

**recoverPose()** [1/3]

```
int cv::recoverPose ( InputArray E,
                      InputArray points1,
                      InputArray points2,
                      InputArray cameraMatrix,
                      OutputArray R,
                      OutputArray t,
                      InputOutputArray mask = noArray()
)
```

**Python:**

```
retval, R, t, mask = cv.recoverPose( E, points1, points2, cameraMatrix[, R[, t[, mask]]] )
retval, R, t, mask = cv.recoverPose( E, points1, points2[, R[, t[, focal[, pp[, mask]]]]]
retval, R, t, mask, triangulatedPoints = cv.recoverPose( E, points1, points2, cameraMatrix, distanceThresh[, R[, t[, mask[, triangulatedPoints]]]] )
```

Recover relative camera rotation and translation from an estimated essential matrix and the corresponding points in two images, using cheirality check.  
Returns the number of inliers which pass the check.

**Parameters**

- E** The input essential matrix.
- points1** Array of N 2D 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 .
- cameraMatrix** Camera matrix  $K = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ . Note that this function assumes that points1 and points2 are feature points from cameras with the same camera matrix.
- R** Recovered relative rotation.
- t** Recovered relative translation.
- mask** Input/output mask for inliers in points1 and points2. : If it is not empty, then it marks inliers in points1 and points2 for then given essential matrix E. Only these inliers will be used to recover pose. In the output mask only inliers which pass the cheirality check. This function decomposes an essential matrix using decomposeEssentialMat and then verifies possible pose hypotheses by doing cheirality check. The cheirality check basically means that the triangulated 3D points should have positive depth. Some details can be found in [144] .

This function can be used to process output E and mask from findEssentialMat. In this scenario, points1 and points2 are the same input for findEssentialMat. :

```
// 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] = ...;
}

// cametra_matrix with both focal lengths = 1, and principal point = (0, 0)
Mat cameraMatrix = Mat::eye(3, 3, CV_64F);

Mat E, R, t, mask;

E = findEssentialMat(points1, points2, cameraMatrix, RANSAC, 0.999, 1.0, mask);
recoverPose(E, points1, points2, cameraMatrix, R, t, mask);
```

**recoverPose()** [2/3]

```
int cv::recoverPose ( InputArray E,
                      InputArray points1,
                      InputArray points2,
                      OutputArray R,
                      OutputArray t,
                      double focal = 1.0,
                      Point2d pp =
                      Point2d(0, 0),
                      InputOutputArray mask = noArray()
)
```

**Python:**

```
retval, R, t, mask = cv.recoverPose( E, points1, points2, cameraMatrix[, R[, t[, mask]]] )
retval, R, t, mask = cv.recoverPose( E, points1, points2[, R[, t[, focal[, pp[, mask]]]]] )
retval, R, t, mask, triangulatedPoints = cv.recoverPose( E, points1, points2, cameraMatrix, distanceThresh[, R[, t[, mask[, triangulatedPoints]]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**Parameters**

**E** The input essential matrix.

**points1** Array of N 2D 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 .

**R** Recovered relative rotation.

**t** Recovered relative translation.

**focal** Focal length of the camera. Note that this function assumes that points1 and points2 are feature points from cameras with same focal length and principal point.

**pp** principal point of the camera.

**mask** Input/output mask for inliers in points1 and points2. : If it is not empty, then it marks inliers in points1 and points2 for then given essential matrix E. Only these inliers will be used to recover pose. In the output mask only inliers which pass the cheirality check.

This function differs from the one above that it computes camera matrix from focal length and principal point:

$$K = \begin{bmatrix} f & 0 & x_{pp} \\ 0 & f & y_{pp} \\ 0 & 0 & 1 \end{bmatrix}$$

**recoverPose()** [3/3]

```
int cv::recoverPose ( InputArray E,
                      InputArray points1,
                      InputArray points2,
                      InputArray cameraMatrix,
                      OutputArray R,
                      OutputArray t,
                      double distanceThresh,
                      InputOutputArray mask = noArray(),
                      OutputArray triangulatedPoints = noArray()
)
```

**Python:**

```
retval, R, t, mask = cv.recoverPose( E, points1, points2, cameraMatrix[, R[, t[, mask]]] )
retval, R, t, mask = cv.recoverPose( E, points1, points2[, R[, t[, focal[, pp[, mask]]]]] )
retval, R, t, mask, triangulatedPoints = cv.recoverPose( E, points1, points2, cameraMatrix, distanceThresh[, R[, t[, mask[, triangulatedPoints]]]] )
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**Parameters**

<b>E</b>	The input essential matrix.
<b>points1</b>	Array of N 2D points from the first image. The point coordinates should be floating-point (single or double precision).
<b>points2</b>	Array of the second image points of the same size and format as points1.
<b>cameraMatrix</b>	Camera matrix $K = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix}$ . Note that this function assumes that points1 and points2 are feature points from cameras with the same camera matrix.
<b>R</b>	Recovered relative rotation.
<b>t</b>	Recovered relative translation.
<b>distanceThresh</b>	threshold distance which is used to filter out far away points (i.e. infinite points).
<b>mask</b>	Input/output mask for inliers in points1 and points2. : If it is not empty, then it marks inliers in points1 and points2 for then given essential matrix E. Only these inliers will be used to recover pose. In the output mask only inliers which pass the cheirality check.
<b>triangulatedPoints</b>	3d points which were reconstructed by triangulation.

**S rectify3Collinear()**

```
float cv::rectify3Collinear ( InputArray cameraMatrix1,  
                             InputArray distCoeffs1,  
                             InputArray cameraMatrix2,  
                             InputArray distCoeffs2,  
                             InputArray cameraMatrix3,  
                             InputArray distCoeffs3,  
                             InputArrayOfArrays imgpt1,  
                             InputArrayOfArrays imgpt3,  
                             Size imageSize,  
                             InputArray R12,  
                             InputArray T12,  
                             InputArray R13,  
                             InputArray T13,  
                             OutputArray R1,  
                             OutputArray R2,  
                             OutputArray R3,  
                             OutputArray P1,  
                             OutputArray P2,  
                             OutputArray P3,  
                             OutputArray Q,  
                             double alpha,  
                             Size newImgSize,  
                             Rect * roi1,  
                             Rect * roi2,  
                             int flags  
)
```

**Python:**

```
retval,  
R1,  
R2,  
R3,  
P1,  
= cv.rectify3Collinear( cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, cameraMatrix3, distCoeffs3, imgpt1, imgpt3, imageSize, R12, T12, R13, T13,  
P2,  
P3,  
Q,  
roi1,  
roi2
```

computes the rectification transformations for 3-head camera, where all the heads are on the same line.

**§ reprojectImageTo3D()**

```
void cv::reprojectImageTo3D ( InputArray disparity,
                            OutputArray _3dImage,
                            InputArray Q,
                            bool handleMissingValues =
                            int ddepth = -1
                        )
```

**Python:**

```
_3dImage = cv.reprojectImageTo3D( disparity, Q[, _3dImage[, handleMissingValues[, ddepth]]) )
```

Reprojects a disparity image to 3D space.

**Parameters**

<b>disparity</b>	Input single-channel 8-bit unsigned, 16-bit signed, 32-bit signed or 32-bit floating-point disparity image. If 16-bit signed format is used, the values are assumed to have no fractional bits.
<b>_3dImage</b>	Output 3-channel floating-point image of the same size as <b>disparity</b> . Each element of <b>_3dImage</b> (x,y) contains 3D coordinates of the point (x,y) computed from the disparity map.
<b>Q</b>	$4 \times 4$ perspective transformation matrix that can be obtained with <b>stereoRectify</b> .
<b>handleMissingValues</b>	Indicates, whether the function should handle missing values (i.e. points where the disparity was not computed). If <b>handleMissingValues</b> =true, then pixels with the minimal disparity that corresponds to the outliers (see <b>StereoMatcher::compute</b> ) are transformed to 3D points with a very large Z value (currently set to 10000).
<b>ddepth</b>	The optional output array depth. If it is -1, the output image will have <b>CV_32F</b> depth. <b>ddepth</b> can also be set to <b>CV_16S</b> , <b>CV_32S</b> or <b>CV_32F</b> .

The function transforms a single-channel disparity map to a 3-channel image representing a 3D surface. That is, for each pixel (x,y) and the corresponding disparity  $d=\text{disparity}(x,y)$ , it computes:

$$\begin{aligned}[X \ Y \ Z \ W]^T &= Q * [x \ y \ \text{disparity}(x,y) \ 1]^T \\ \underline{\text{_3dImage}}(x,y) &= (X/W, \ Y/W, \ Z/W)\end{aligned}$$

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  $\{(x,y,d),\dots\}$  to 3D space, use **perspectiveTransform**.

## § Rodrigues()

```
void cv::Rodrigues ( InputArray src,
                     OutputArray dst,
                     OutputArray jacobian = noArray()
)
```

**Python:**

```
dst, jacobian = cv.Rodrigues( src[, dst[, jacobian]] )
```

Converts a rotation matrix to a rotation vector or vice versa.

**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.

$$\theta \leftarrow \text{norm}(r)$$

$$r \leftarrow r/\theta$$

$$R = \cos \theta I + (1 - \cos \theta) rr^T + \sin \theta \begin{bmatrix} 0 & -r_z & r_y \\ r_z & 0 & -r_x \\ -r_y & r_x & 0 \end{bmatrix}$$

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`.

**Examples:**

`samples/cpp/tutorial_code/features2D/Homography/decompose_homography.cpp`,  
`samples/cpp/tutorial_code/features2D/Homography/homography_from_camera_displacement.cpp`, and  
`samples/cpp/tutorial_code/features2D/Homography/pose_from_homography.cpp`.

**RQDecomp3x3()**

```
Vec3d cv::RQDecomp3x3 ( InputArray src,
                         OutputArray mtxR,
                         OutputArray mtxQ,
                         Qx = noArray(),
                         Qy = noArray(),
                         Qz = noArray()
)
```

**Python:**

```
retval, mtxR, mtxQ, Qx, Qy, Qz = cv.RQDecomp3x3( src[, mtxR[, mtxQ[, Qx[, Qy[, Qz]]]]] )
```

Computes an RQ decomposition of 3x3 matrices.

**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 principal axes that results in the same orientation of an object, e.g. see [173]. Returned tree rotation matrices and corresponding three Euler angles are only one of the possible solutions.

## § sampsonDistance()

```
double cv::sampsonDistance ( InputArray pt1,
                            InputArray pt2,
                            InputArray F
                           )
```

### Python:

```
retval = cv.sampsonDistance( pt1, pt2, F )
```

Calculates the Sampson Distance between two points.

The function `cv::sampsonDistance` calculates and returns the first order approximation of the geometric error as:

$$sd(pt1, pt2) = \frac{(pt2^t \cdot F \cdot pt1)^2}{((F \cdot pt1)(0))^2 + ((F \cdot pt1)(1))^2 + ((F^t \cdot pt2)(0))^2 + ((F^t \cdot pt2)(1))^2}$$

The fundamental matrix may be calculated using the `cv::findFundamentalMat` function. See [77] 11.4.3 for details.

### Parameters

- pt1** first homogeneous 2d point
- pt2** second homogeneous 2d point
- F** fundamental matrix

### Returns

The computed Sampson distance.

## § solveP3P()

```
int cv::solveP3P ( InputArray objectPoints,
                   InputArray imagePoints,
                   InputArray cameraMatrix,
                   InputArray distCoeffs,
                   OutputArrayOfArrays rvecs,
                   OutputArrayOfArrays tvecs,
                   int flags
                  )
```

### Python:

```
retval, rvecs, tvecs = cv.solveP3P( objectPoints, imagePoints, cameraMatrix, distCoeffs, flags[, rvecs[, tvecs]] )
```

Finds an object pose from 3 3D-2D point correspondences.

### Parameters

- objectPoints** Array of object points in the object coordinate space, 3x3 1-channel or 1x3/3x1 3-channel. `vector<Point3f>` can be also passed here.
- imagePoints** Array of corresponding image points, 3x2 1-channel or 1x3/3x1 2-channel. `vector<Point2f>` can be also passed here.
- cameraMatrix** Input camera matrix  $A = \begin{bmatrix} fx & 0 & cx \\ 0 & fy & cy \\ 0 & 0 & 1 \end{bmatrix}$ .
- distCoeffs** Input vector of distortion coefficients ( $k_1, k_2, p_1, p_2, [k_3, k_4, k_5, k_6, s_1, s_2, s_3, s_4, [\tau_x, \tau_y]]$ ) of 4, 5, 8, 12 or 14 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
- rvecs** Output rotation vectors (see Rodrigues ) that, together with tvecs , brings points from the model coordinate system to the camera coordinate system. A P3P problem has up to 4 solutions.
- tvecs** Output translation vectors.
- flags** Method for solving a P3P problem:
  - **SOLVEPNP\_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" ([64]).
  - **SOLVEPNP\_AP3P** Method is based on the paper of Tong Ke and Stergios I. Roumeliotis. "An Efficient Algebraic Solution to the Perspective-Three-Point Problem" ([96]).

The function estimates the object pose given 3 object points, their corresponding image projections, as well as the camera matrix and the distortion coefficients.

[§ solvePnP\(\)](#)

```
bool cv::solvePnP ( InputArray objectPoints,
                    InputArray imagePoints,
                    InputArray cameraMatrix,
                    InputArray distCoeffs,
                    OutputArray rvec,
                    OutputArray tvec,
                    bool useExtrinsicGuess = false,
                    int flags = SOLVEPNP_ITERATIVE
)
```

**Python:**

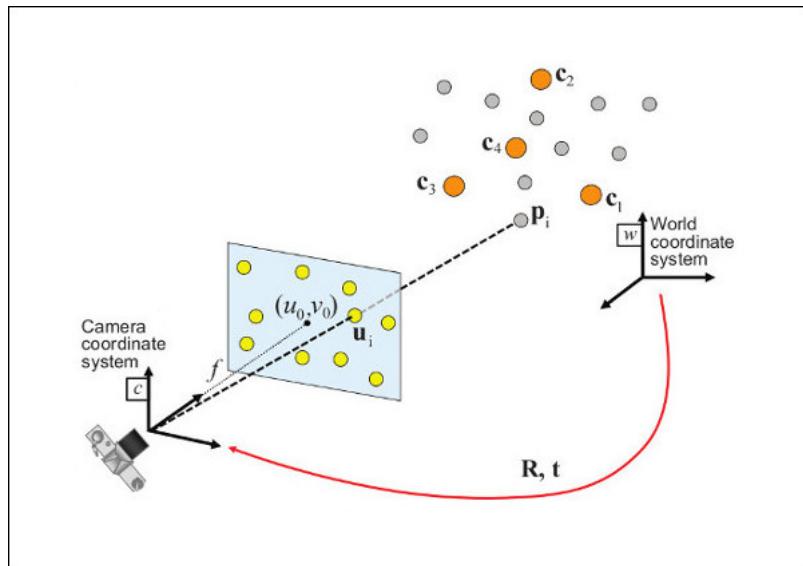
```
retval, rvec, tvec = cv.solvePnP( objectPoints, imagePoints, cameraMatrix, distCoeffs[, rvec[, tvec[, useExtrinsicGuess[, flags]]]] )
```

Finds an object pose from 3D-2D point correspondences.

**Parameters**

<b>objectPoints</b>	Array of object points in the object coordinate space, Nx3 1-channel or 1xN/Nx1 3-channel, where N is the number of points. vector<Point3f> can be also passed here.
<b>imagePoints</b>	Array of corresponding image points, Nx2 1-channel or 1xN/Nx1 2-channel, where N is the number of points. vector<Point2f> can be also passed here.
<b>cameraMatrix</b>	Input camera matrix $A = \begin{bmatrix} fx & 0 & cx \\ 0 & fy & cy \\ 0 & 0 & 1 \end{bmatrix}$ .
<b>distCoeffs</b>	Input vector of distortion coefficients ( $k_1, k_2, p_1, p_2, [k_3, k_4, k_5, k_6, s_1, s_2, s_3, s_4, [\tau_x, \tau_y]]$ ) of 4, 5, 8, 12 or 14 elements. If the vector is NULL/empty, the zero distortion coefficients are assumed.
<b>rvec</b>	Output rotation vector (see <a href="#">Rodrigues</a> ) that, together with tvec , brings points from the model coordinate system to the camera coordinate system.
<b>tvec</b>	Output translation vector.
<b>useExtrinsicGuess</b>	Parameter used for <a href="#">SOLVEPNP_ITERATIVE</a> . 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.
<b>flags</b>	Method for solving a PnP problem: <ul style="list-style-type: none"> <li>• <b>SOLVEPNP_ITERATIVE</b> 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 .</li> <li>• <b>SOLVEPNP_P3P</b> 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" (<a href="#">[64]</a>). In this case the function requires exactly four object and image points.</li> <li>• <b>SOLVEPNP_AP3P</b> Method is based on the paper of T. Ke, S. Roumeliotis "An Efficient Algebraic Solution to the Perspective-Three-Point Problem" (<a href="#">[96]</a>). In this case the function requires exactly four object and image points.</li> <li>• <b>SOLVEPNP_EPNP</b> Method has been introduced by F.Moreno-Noguer, V.Lepetit and P.Fua in the paper "EPnP: Efficient Perspective-n-Point Camera Pose Estimation" (<a href="#">[107]</a>).</li> <li>• <b>SOLVEPNP_DLS</b> Method is based on the paper of Joel A. Hesch and Stergios I. Roumeliotis. "A Direct Least-Squares (DLS) Method for PnP" (<a href="#">[84]</a>).</li> <li>• <b>SOLVEPNP_UPNP</b> Method is based on the paper of A.Penate-Sanchez, J.Andrade-Cetto, F.Moreno-Noguer. "Exhaustive Linearization for Robust Camera Pose and Focal Length Estimation" (<a href="#">[148]</a>). In this case the function also estimates the parameters <math>f_x</math> and <math>f_y</math> assuming that both have the same value. Then the cameraMatrix is updated with the estimated focal length.</li> <li>• <b>SOLVEPNP_AP3P</b> Method is based on the paper of Tong Ke and Stergios I. Roumeliotis. "An Efficient Algebraic Solution to the Perspective-Three-Point Problem" (<a href="#">[96]</a>). In this case the function requires exactly four object and image points.</li> </ul>

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, see the figure below (more precisely, the X-axis of the camera frame is pointing to the right, the Y-axis downward and the Z-axis forward).



Points expressed in the world frame  $\mathbf{X}_w$  are projected into the image plane  $[u, v]$  using the perspective projection model  $\Pi$  and the camera intrinsic parameters matrix  $\mathbf{A}$ :

$$\begin{bmatrix} u \\ v \\ 1 \end{bmatrix} = \mathbf{A} \Pi^c \mathbf{M}_w \begin{bmatrix} X_w \\ Y_w \\ Z_w \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} u \\ v \\ 1 \end{bmatrix} = \begin{bmatrix} f_x & 0 & c_x \\ 0 & f_y & c_y \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix} \begin{bmatrix} r_{11} & r_{12} & r_{13} & t_x \\ r_{21} & r_{22} & r_{23} & t_y \\ r_{31} & r_{32} & r_{33} & t_z \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} X_w \\ Y_w \\ Z_w \\ 1 \end{bmatrix}$$

The estimated pose is thus the rotation (`rvec`) and the translation (`tvec`) vectors that allow to transform a 3D point expressed in the world frame into the camera frame:

$$\begin{bmatrix} X_c \\ Y_c \\ Z_c \\ 1 \end{bmatrix} = {}^c \mathbf{M}_w \begin{bmatrix} X_w \\ Y_w \\ Z_w \\ 1 \end{bmatrix}$$

$$\begin{bmatrix} X_c \\ Y_c \\ Z_c \\ 1 \end{bmatrix} = \begin{bmatrix} r_{11} & r_{12} & r_{13} & t_x \\ r_{21} & r_{22} & r_{23} & t_y \\ r_{31} & r_{32} & r_{33} & t_z \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} X_w \\ Y_w \\ Z_w \\ 1 \end{bmatrix}$$

#### Note

- An example of how to use `solvePnP` for planar augmented reality can be found at [opencv\\_source\\_code/samples/python/plane\\_ar.py](#)
- If you are using Python:
  - Numpy array slices won't work as input because `solvePnP` requires contiguous arrays (enforced by the assertion using `cv::Mat::checkVector()` around line 55 of modules/calib3d/src/solvepnp.cpp version 2.4.9)
  - The P3P algorithm requires image points to be in an array of shape (N,1,2) due to its calling of `cv::undistortPoints` (around line 75 of modules/calib3d/src/solvepnp.cpp version 2.4.9) which requires 2-channel information.
  - Thus, given some data  $D = np.array(...)$  where  $D.shape = (N,M)$ , in order to use a subset of it as, e.g., `imagePoints`, one must effectively copy it into a new array: `imagePoints = np.ascontiguousarray(D[:,2]).reshape((N,1,2))`
- The methods `SOLVEPNP_DLS` and `SOLVEPNP_UPNP` cannot be used as the current implementations are unstable and sometimes give completely wrong results. If you pass one of these two flags, `SOLVEPNP_EPNP` method will be used instead.
- The minimum number of points is 4 in the general case. In the case of `SOLVEPNP_P3P` and `SOLVEPNP_AP3P` methods, it is required to use exactly 4 points (the first 3 points are used to estimate all the solutions of the P3P problem, the last one is used to retain the best solution that minimizes the reprojection error).
- With `SOLVEPNP_ITERATIVE` method and `useExtrinsicGuess=true`, the minimum number of points is 3 (3 points are sufficient to compute a pose but there are up to 4 solutions). The initial solution should be close to the global solution to converge.

#### Examples:

[samples/cpp/tutorial\\_code/features2D/Homography/decompose\\_homography.cpp](#), and  
[samples/cpp/tutorial\\_code/features2D/Homography/homography\\_from\\_camera\\_displacement.cpp](#).

#### § solvePnPRansac()

```

bool cv::solvePnP( InputArray objectPoints,
                  InputArray imagePoints,
                  InputArray cameraMatrix,
                  InputArray distCoeffs,
                  OutputArray rvec,
                  OutputArray tvec,
                  bool useExtrinsicGuess = false,
                  int iterationsCount = 100,
                  float reprojectionError = 8.0,
                  double confidence = 0.99,
                  OutputArray inliers = noArray(),
                  int flags = SOLVEPNP_ITERATIVE
)

```

**Python:**

```

retval,
rvec,
tvec, = cv.solvePnP( objectPoints, imagePoints, cameraMatrix, distCoeffs[, rvec[, tvec[, useExtrinsicGuess[, iterationsCount[, reprojectionError[, confidence[ inliers

```

Finds an object pose from 3D-2D point correspondences using the RANSAC scheme.

**Parameters**

<b>objectPoints</b>	Array of object points in the object coordinate space, Nx3 1-channel or 1xN/Nx1 3-channel, where N is the number of points. vector<Point3f> passed here.
<b>imagePoints</b>	Array of corresponding image points, Nx2 1-channel or 1xN/Nx1 2-channel, where N is the number of points. vector<Point2f> can be also
<b>cameraMatrix</b>	Input camera matrix $A = \begin{bmatrix} fx & 0 & cx \\ 0 & fy & cy \\ 0 & 0 & 1 \end{bmatrix}$ .
<b>distCoeffs</b>	Input vector of distortion coefficients ( $k_1, k_2, p_1, p_2[, k_3[, k_4, k_5, k_6[, s_1, s_2, s_3, s_4[, \tau_x, \tau_y]]]]$ ) of 4, 5, 8, 12 or 14 elements. If the vector contains less than 4 elements, the zero distortion coefficients are assumed.
<b>rvec</b>	Output rotation vector (see Rodrigues ) that, together with tvec , brings points from the model coordinate system to the camera coordinate system.
<b>tvec</b>	Output translation vector.
<b>useExtrinsicGuess</b>	Parameter used for SOLVEPNP_ITERATIVE. 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.
<b>iterationsCount</b>	Number of iterations.
<b>reprojectionError</b>	Inlier threshold value used by the RANSAC procedure. The parameter value is the maximum allowed distance between the observed and the projected 2D points to consider it an inlier.
<b>confidence</b>	The probability that the algorithm produces a useful result.
<b>inliers</b>	Output vector that contains indices of inliers in objectPoints and imagePoints .
<b>flags</b>	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 such a pose that minimizes reprojection error, that is, the sum of squared distances between the observed projections imagePoints and the projected (using projectP objectPoints). The use of RANSAC makes the function resistant to outliers.

**Note**

- An example of how to use solvePNPRansac for object detection can be found at opencv\_source\_code/samples/cpp/tutorial\_code/calib3d/real\_time\_pose\_estimation.cpp
- The default method used to estimate the camera pose for the Minimal Sample Sets step is **SOLVEPNP\_EPNP**. Exceptions are:
  - if you choose **SOLVEPNP\_P3P** or **SOLVEPNP\_AP3P**, these methods will be used.
  - if the number of input points is equal to 4, **SOLVEPNP\_P3P** is used.
- The method used to estimate the camera pose using all the inliers is defined by the flags parameters unless it is equal to **SOLVEPNP\_P3P** or **SOLVEPNP\_AP3P** case, the method **SOLVEPNP\_EPNP** will be used instead.

**§ stereoCalibrate()** [1/2]

```

double cv::stereoCalibrate ( InputArrayOfArrays objectPoints,
                            InputArrayOfArrays imagePoints1,
                            InputArrayOfArrays imagePoints2,
                            InputOutputArray cameraMatrix1,
                            InputOutputArray distCoeffs1,
                            InputOutputArray cameraMatrix2,
                            InputOutputArray distCoeffs2,
                            Size imageSize,
                            InputOutputArray R,
                            InputOutputArray T,
                            OutputArray E,
                            OutputArray F,
                            OutputArray perViewErrors,
                            int flags = CALIB_FIX_INTRINSIC,
                            TermCriteria criteria = TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 1e-6)
)

```

**Python:**

```

retval,
cameraMatrix1,
distCoeffs1,
= cv.stereoCalibrate(          objectPoints, imagePoints1, imagePoints2, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize[, R[, T[, distCoeffs2, R, T, E, F]]]
cameraMatrix2,
distCoeffs2, R,
T, E, F
retval,
cameraMatrix1,
distCoeffs1,
cameraMatrix2, = cv.stereoCalibrateExtended( objectPoints, imagePoints1, imagePoints2, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize, R, T, distCoeffs2, R,
T, E, F,
perViewErrors

```

Calibrates the stereo camera.

**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 CALIB\_USE\_INTRINSIC\_GUESS , CALIB\_FIX\_ASPECT\_RATIO , 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 ( $k_1, k_2, p_1, p_2, [k_3, k_4, k_5, k_6, s_1, s_2, s_3, s_4, \tau_x, \tau_y]$ ) of 4, 5, 8, 12 or 14 elements. The ou

**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.

**perViewErrors** Output vector of the RMS re-projection error estimated for each pattern view.

**flags** Different flags that may be zero or a combination of the following values:

- **CALIB\_FIX\_INTRINSIC** Fix cameraMatrix? and distCoeffs? so that only R, T, E , and F matrices are estimated.
- **CALIB\_USE\_INTRINSIC\_GUESS** Optimize some or all of the intrinsic parameters according to the specified flags. Initial values are pr
- **CALIB\_USE\_EXTRINSIC\_GUESS** R, T contain valid initial values that are optimized further. Otherwise R, T are initialized to the medi
- dimension separately).
- **CALIB\_FIX\_PRINCIPAL\_POINT** Fix the principal points during the optimization.
- **CALIB\_FIX\_FOCAL\_LENGTH** Fix  $f_x^{(j)}$  and  $f_y^{(j)}$  .
- **CALIB\_FIX\_ASPECT\_RATIO** Optimize  $f_y^{(j)}$  . Fix the ratio  $f_x^{(j)}/f_y^{(j)}$
- **CALIB\_SAME\_FOCAL\_LENGTH** Enforce  $f_x^{(0)} = f_x^{(1)}$  and  $f_y^{(0)} = f_y^{(1)}$  .
- **CALIB\_ZERO\_TANGENT\_DIST** Set tangential distortion coefficients for each camera to zeros and fix there.
- **CALIB\_FIX\_K1,...,CALIB\_FIX\_K6** Do not change the corresponding radial distortion coefficient during the optimization. If CALIB\_USE coefficient from the supplied distCoeffs matrix is used. Otherwise, it is set to 0.

- **CALIB\_RATIONAL\_MODEL** Enable coefficients k4, k5, and k6. To provide the backward compatibility, this extra flag should be explicitly used. If the flag is not set, the function computes and returns only 5 distortion coefficients.
- **CALIB\_THIN\_PRISM\_MODEL** Coefficients s1, s2, s3 and s4 are enabled. To provide the backward compatibility, this extra flag should be explicitly used. If the flag is not set, the function computes and returns only 5 distortion coefficients.
- **CALIB\_FIX\_S1\_S2\_S3\_S4** The thin prism distortion coefficients are not changed during the optimization. If **CALIB\_USE\_INTRINSIC\_MODEL** is supplied, distCoeffs matrix is used. Otherwise, it is set to 0.
- **CALIB\_TILTED\_MODEL** Coefficients tauX and tauY are enabled. To provide the backward compatibility, this extra flag should be explicitly used. If the flag is not set, the function computes and returns only 5 distortion coefficients.
- **CALIB\_FIX\_TAUX\_TAUY** The coefficients of the tilted sensor model are not changed during the optimization. If **CALIB\_USE\_INTRINSIC\_MODEL** is supplied, distCoeffs matrix is used. Otherwise, it is set to 0.

**criteria** Termination criteria for the iterative optimization algorithm.

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 are known, then the function can estimate the relative position and orientation of an object relative to the first camera and to the second camera, ( $R_1, T_1$ ) and ( $R_2, T_2$ ), respectively (this can be done with `solvePnP`), then those poses definitely intersect. Given ( $R_1, T_1$ ), it should be possible to compute ( $R_2, T_2$ ). You only need to know the position and orientation of the second camera relative to the first camera. The function `stereoCalibrate` computes ( $R, T$ ) so that:

$$R_2 = R * R_1$$

$$T_2 = R * T_1 + T,$$

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, 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`), it is recommended to do so and then pass **CALIB\_FIX\_INTRINSIC** flag to the function along with the computed intrinsic parameters. Otherwise, if all the parameters are not known, it is recommended to restrict some parameters, for example, pass **CALIB\_SAME\_FOCAL\_LENGTH** and **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 intrinsic parameters for both cameras, the extrinsic parameters for the second camera, and the fundamental matrix.

[§ `stereoCalibrate\(\)`](#) [2/2]

```
double cv::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,
                            int flags = CALIB_FIX_INTRINSIC,
                            TermCriteria criteria = TermCriteria(TermCriteria::COUNT+TermCriteria::EPS, 30, 1e-6)
                        )
```

**Python:**

```
retval,  
cameraMatrix1,  
distCoeffs1,  
= cv.stereoCalibrate(          objectPoints, imagePoints1, imagePoints2, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize[, R[,  
cameraMatrix2,  
distCoeffs2, R,  
T, E, F  
retval,  
cameraMatrix1,  
distCoeffs1,  
cameraMatrix2, = cv.stereoCalibrateExtended( objectPoints, imagePoints1, imagePoints2, cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize, R, T  
distCoeffs2, R,  
T, E, F,  
perViewErrors
```

This is an overloaded member function, provided for convenience. It differs from the above function only in what argument(s) it accepts.

**s stereoRectify()**

```
void cv::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,
                        flags =
int          CALIB_ZERO_DISPARITY,
double        alpha = -1,
Size           newSize = Size(),
Rect *         validPixROI1 = 0,
Rect *         validPixROI2 = 0
)
```

**Python:**

```
R1, R2, P1,
P2, Q,
= cv.stereoRectify( cameraMatrix1, distCoeffs1, cameraMatrix2, distCoeffs2, imageSize, R, T, R1, R2, P1, P2, Q, flags, alpha, newSize)
validPixROI1,
validPixROI2
```

Computes rectification transforms for each head of a calibrated stereo camera.

**Parameters**

- cameraMatrix1** First camera matrix.
- distCoeffs1** First camera distortion parameters.
- cameraMatrix2** Second camera matrix.
- 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 `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:

- **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:

$$P1 = \begin{bmatrix} f & 0 & cx_1 & 0 \\ 0 & f & cy & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

$$P2 = \begin{bmatrix} f & 0 & cx_2 & T_x * f \\ 0 & f & cy & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix},$$

where  $T_x$  is a horizontal shift between the cameras and  $cx_1 = cx_2$  if `CALIB_ZERO_DISPARITY` is set.

- **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:

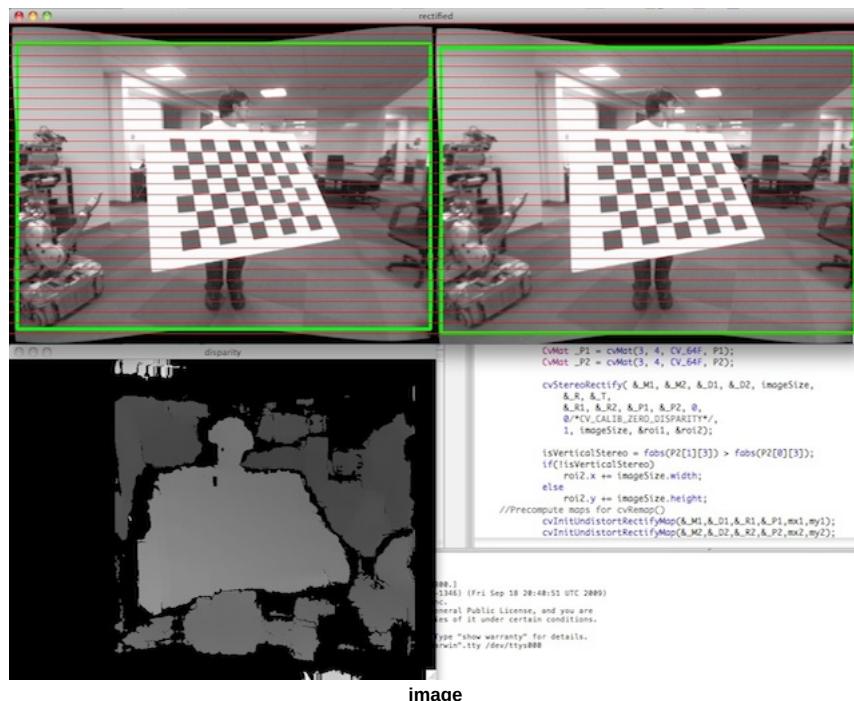
$$P1 = \begin{bmatrix} f & 0 & cx & 0 \\ 0 & f & cy_1 & 0 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

$$P2 = \begin{bmatrix} f & 0 & cx & 0 \\ 0 & f & cy_2 & T_y * f \\ 0 & 0 & 1 & 0 \end{bmatrix},$$

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 pixel:



`s stereoRectifyUncalibrated()`

```
bool cv::stereoRectifyUncalibrated ( InputArray points1,
                                     InputArray points2,
                                     InputArray F,
                                     Size imgSize,
                                     OutputArray H1,
                                     OutputArray H2,
                                     double threshold = 5
)
```

**Python:**

```
retval, H1, H2 = cv.stereoRectifyUncalibrated( points1, points2, F, imgSize[, H1[, H2[, threshold]]] )
```

Computes a rectification transform for an uncalibrated stereo camera.

**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  $|points2[i]^T * F * points1[i]| > threshold$ ) 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 [78].

**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()**

```
void cv::triangulatePoints ( InputArray projMatr1,
                            InputArray projMatr2,
                            InputArray projPoints1,
                            InputArray projPoints2,
                            OutputArray points4D
)
```

**Python:**

```
points4D = cv.triangulatePoints( projMatr1, projMatr2, projPoints1, projPoints2[, points4D] )
```

Reconstructs points by triangulation.

**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`.

**Note**

Keep in mind that all input data should be of float type in order for this function to work.

**See also**

[reprojectImageTo3D](#)

## § validateDisparity()

```
void cv::validateDisparity ( InputOutputArray disparity,
                            InputArray cost,
                            int minDisparity,
                            int numberOfDisparities,
                            int disp12MaxDisp = 1
                        )
```

### Python:

```
disparity = cv.validateDisparity( disparity, cost, minDisparity, numberOfDisparities[, disp12MaxDisp] )
```

validates disparity using the left-right check. The matrix "cost" should be computed by the stereo correspondence algorithm