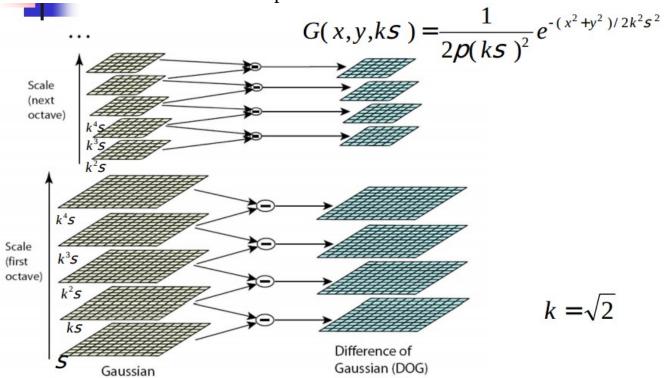
#### Scale Invariant Feature Transform

- 1. Scale space peak selection
- this step is used for finding potential location of features
  - use different gaussians for capturing different scales (different scale space)
  - subtract gaussians to get blob detector (laplacian of gaussian) because log can be approximated by dog and dog is computationally faster.
  - find if pixel is maximum/minimum in 3x3x3 (8 pixels in same level, 9 above, 9 below) neighbourhood to select it as extrema point. (peak selection)
  - use different octaves to reduce computation. For this, downscale the image, find extrema and upscale the location. But use the downscale location in the next 3/4 steps. ?



#### 2. Key point localization

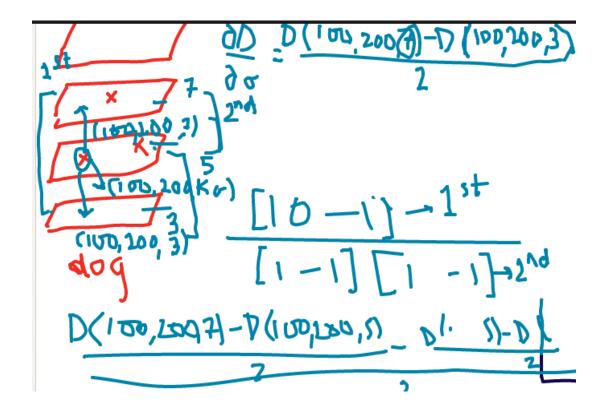
-previous step found potential points i.e. approximate location and could be features.

#### (a) Initial outlier rejection

- get proper location
- use taylor series expansion to find maximum of surface
- earlier step found approximate point in the neighbourhood (because of smoothing). This step finds the exact point in that neighbourhood.
  - for all the potential points (x,y,scale,octave), perform the following steps
  - take the dog values
  - calculate first derivative wrt x,y,sigma

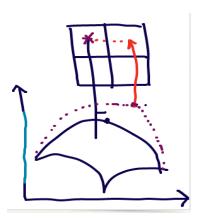
$$\begin{bmatrix}
9a \\
9D \\
9D
\\
9D
\\
5
\\
D(x'A+1'a)-D(x'A-1'a)
\\
5
\\
D(x+1'A'a)-D(x-1'a)
\\
X = (x'A'a) \longrightarrow Dad$$

calculate second derivative



• find the proper offset in (x,y, sigma)(shift in location) of extreme. Add the offset if bigger than 0.5. This is the new location of that point.

\*round instead of ceil



substitute the offset in taylor series and threshold the value. Discard if

#### less than 0.03

-this step rejects the points with less contrast.

## (b) Further outlier rejection

- -This step is rejecting edges.(in xy plane of dog plane forget sigma)
- -So use harris corner value.
  - Form the hessian matrix for the dog surface at that point

H = [Dxx Dxy]

Dxy Dyy]

- find eigenvalues lambda1, lambda2
- calculate corner value as

r = lambda1/lambda2

• reject the points if r > 10

## 3. Orientation assignment

- -find dominant orientations of the keypoint. This is for converting the feature into rotation invariant.
  - find the magnitude and orientation for every point in 4x4 neighbourhood around that point for that scale.

L = smoothed image at scale = gaussian (not dog)

$$dx = L(x+1,y) - L(x-1,y)$$
 [-1 0 1]

$$dy = L(x,y+1) - L(x,y-1)$$
 [-1 0 1]

magnitude(x,y) = sqrt(dx\*dx + dy\*dy)

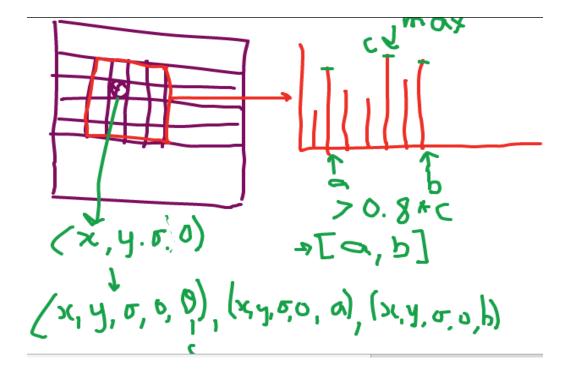
orientation(x,y) = atan(dy/dx)

• create a weighted direction histogram.

36 bins (meaning each bin covers 10 degrees i.e. 0-10, 10-20, 20-30, etc.) orientation' = orientation/10 -> assign to the bin

weight = magnitude and spatial gaussian filter with sigma = 1.5 \* scale of point

• find the dominant directions in the histogram. Maximum = dominant direction. Any direction having > 0.8\*domiant direction, consider them as potential points also.



#### 4. Keypoint descriptor

-describing the key point as a high dimensional vector

- take 16x16 pixel window around the point
- compute magnitude and relative orientation (=orientation -dominant direction) for every pixel
- 16x16 divide into 4x4 regions. So, every block contains 4x4 pixels

4x4 pixels	4x4	4x4	4x4

- for every block, construct weighted histogram (8 bins, so 0-45,45-90, etc., weight=magnitude and spatial gaussian)
- concatenate all the histograms. 8Bins x 16 blocks=> 128 vector

# postprocessing

• normalize 128D vector to unit vector

• for non-linear intensity transform, remove any value greater than 0.2 in the normalized vector and again renormalize?

$$[0.2,0,0,0.1] \rightarrow [0.66,0,0,0.33]$$

## 5. keypoints matching

-match the keypoints against database or another image

feature1 -> 128D vector in image 1 feature2 -> 128D vector in image2

euclidean distance = sqrt(  $(f1[1]-f2[1])^2 + (f1[2]-f2[2])^2 + ...$ )

```
image1 – featuresi1, featuresi2, features i3,... image2 – featuresj1, featuresj2, features j3,...
```

- take featuresi1
- compute euclidean distance wrt featuresj1, featuresj2, featuresj3,...
- find feature with minimum euclidean distance --> matching feature1
- ratio = distance best match / distance 2<sup>nd</sup> best match
- consider the second best match (2<sup>nd</sup> minimum distance) if ratio <=0.8
- threshold distance, so both have to be less than threshold distance