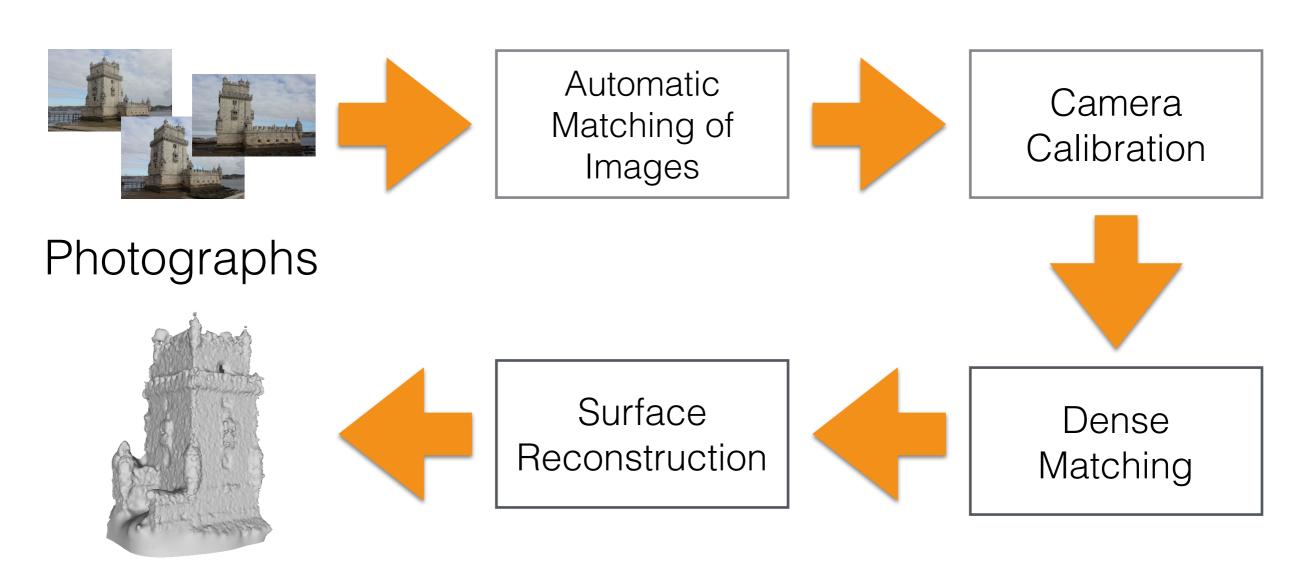
3D from Photographs: Automatic Matching of Images

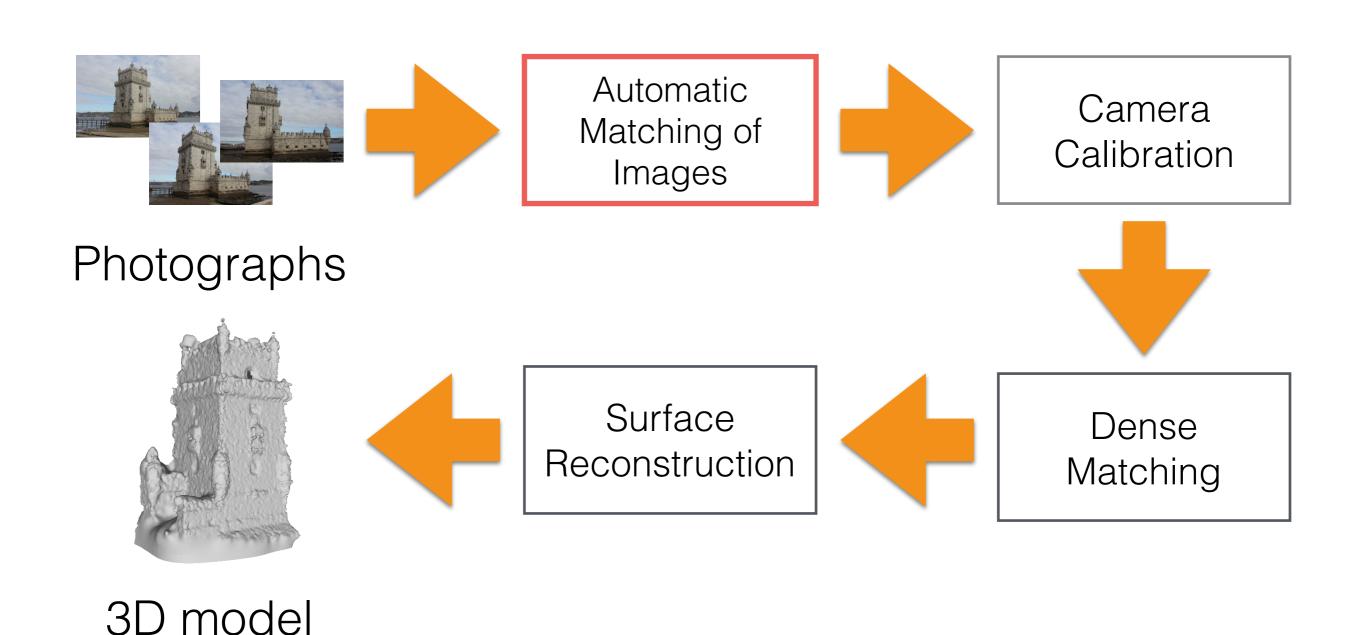
Dr Francesco Banterle francesco.banterle@isti.cnr.it

3D from Photographs



3D model

3D from Photographs



The Matching Problem

 We need to find corresponding feature across two or more views:



The Matching Problem

- Why?
 - 3D Reconstruction.
 - Image Registration.
 - Visual Tracking.
 - Object Recognition.
 - etc.

The Matching Problem: Automatic Panorama Generation



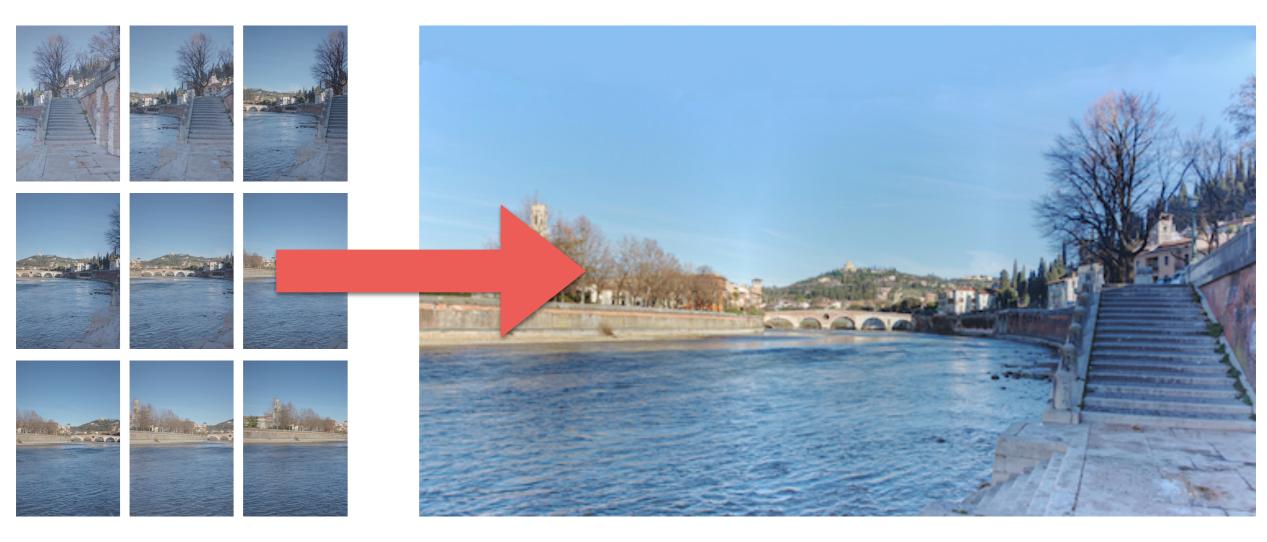
Input Photographs

The Matching Problem: Automatic Panorama Generation



Input Photographs

The Matching Problem: Automatic Panorama Generation



Input Photographs

Panorama

Extraction of Features

Features

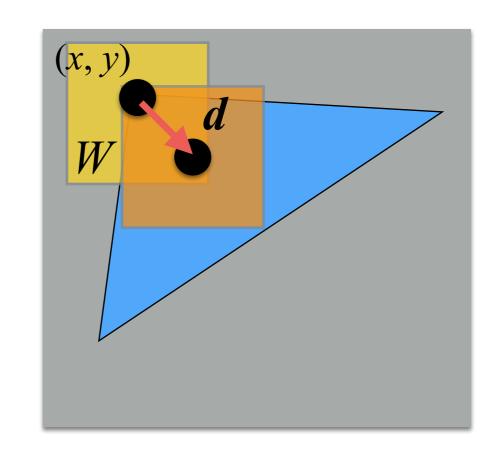
- A feature is a piece of the input image that is relevant for solving a given task.
- Features can be global or local.
- We will focus on local features that are more robust to occlusions and variations.

Extraction of Local Features

- We can extract different kind of features:
 - Flat regions or Blobs
 - Edges
 - Corners

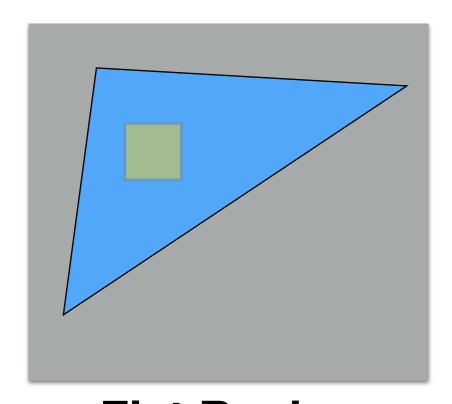
Harris Corner Detector

- Let's consider a window W centered in (x, y):
 - how do pixels change from a window in (x, y) to another one with a shift d = (u, v)?
 - Let's compare each pixel before and after moving W by d = (u, v) using the sum of squared differenced (SSD).

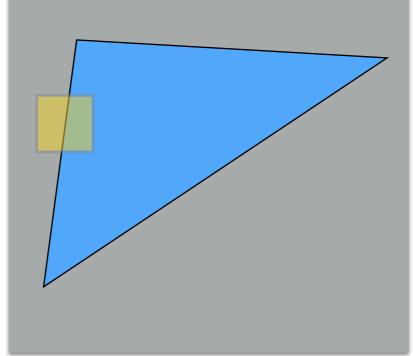


$$E(x,y) = \sum_{x_k,y_k \in W(x,y)} \left(I(x_k + u, y_k + v) - I(x_k, y_k) \right)^2$$

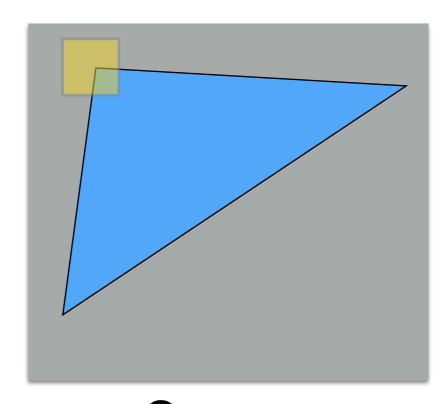
What a Corners is



Flat Region:
no change
in all directions.



Edge:
no change
along the edge.



Corner: significant change in all directions.

 Let's apply a first-order approximation, which provides good results for small motions:

$$I(x+u,y+v) \approx I(x,y) + \frac{\partial I}{\partial x}u + \frac{\partial I}{\partial y}v$$
$$\approx I(x,y) + \begin{bmatrix} I_x & I_y \end{bmatrix} \cdot \begin{bmatrix} u \\ v \end{bmatrix}$$

$$\begin{split} E(x,y) &= \sum_{x_k,y_k \in W(x,y)} \left(I(x_k + u, y_k + v) - I(x_k, y_k) \right)^2 \\ &\approx \sum_{x_k,y_k \in W(x,y)} \left(I(x_k, y_k) + I_x(x_k, y_k)u + I_y(x_k, y_k)v - I(x_k, y_k) \right)^2 \\ &= \sum_{x_k,y_k \in W(x,y)} \left(I_x(x_k, y_k)u + I_y(x_k, y_k)v \right)^2 \\ &= \sum_{x_k,y_k \in W(x,y)} \left(I_x(x_k, y_k)^2u^2 + 2I_x(x_k, y_k)I_y(x_k, y_k) + I_y(x_k, y_k)^2v^2 \right) \end{split}$$

$$E(x,y) \approx \sum_{x_k,y_k \in W(x,y)} \left(I_x(x_k, y_k)^2 u^2 + 2I_x(x_k, y_k) I_y(x_k, y_k) + I_y(x_k, y_k)^2 v^2 \right)$$

= $Au^2 + 2Buv + Cv^2$

$$A = \sum_{x_k, y_k \in W(x, y)} I_x(x_k, y_k)^2$$

$$B = \sum_{x_k, y_k \in W(x, y)} I_x(x_k, y_k) I_y(x_k, y_k)$$

$$C = \sum_{x_k, y_k \in W(x, y)} I_y(x_k, y_k)^2$$

 The surface at (x, y) can be locally approximate by a quadratic form:

$$E(x,y) \approx Au^{2} + 2Buv + Cv^{2}$$

$$\approx \begin{bmatrix} u & v \end{bmatrix} \cdot \begin{bmatrix} A & B \\ B & C \end{bmatrix} \cdot \begin{bmatrix} u \\ v \end{bmatrix}$$

$$A = \sum_{x_k, y_k \in W(x, y)} I_x(x_k, y_k)^2$$

$$B = \sum_{x_k, y_k \in W(x, y)} I_x(x_k, y_k) I_y(x_k, y_k)$$

$$C = \sum_{x_k, y_k \in W(x, y)} I_y(x_k, y_k)^2$$

• E(x,y) can be rewritten as

$$E(x,y) \approx \sum_{x_k,y_k \in W(x,y)} \begin{bmatrix} u & v \end{bmatrix} \cdot \begin{bmatrix} I_x^2(x_k,y_k) & I_x(x_k,y_k)I_y(x_k,y_k) \\ I_x(x_k,y_k)I_y(x_k,y_k) & I_y^2(x_k,y_k) \end{bmatrix} \cdot \begin{bmatrix} u \\ v \end{bmatrix}$$
$$= \begin{bmatrix} u & v \end{bmatrix} \cdot M \cdot \begin{bmatrix} u \\ v \end{bmatrix}$$

$$M = \sum_{x_k, y_k \in W(x,y)} \begin{bmatrix} I_x^2(x_k, y_k) & I_x(x_k, y_k)I_y(x_k, y_k) \\ I_x(x_k, y_k)I_y(x_k, y_k) & I_y^2(x_k, y_k) \end{bmatrix}$$

• E(x,y) can be rewritten as

$$E(x,y) \approx \sum_{x_k,y_k \in W(x,y)} \begin{bmatrix} u & v \end{bmatrix} \cdot \begin{bmatrix} I_x^2(x_k,y_k) & I_x(x_k,y_k)I_y(x_k,y_k) \\ I_x(x_k,y_k)I_y(x_k,y_k) & I_y^2(x_k,y_k) \end{bmatrix} \cdot \begin{bmatrix} u \\ v \end{bmatrix}$$

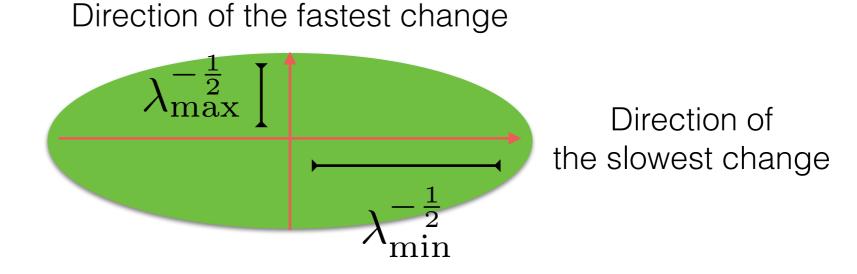
$$= \begin{bmatrix} u & v \end{bmatrix} \cdot M \cdot \begin{bmatrix} u \\ v \end{bmatrix} \qquad \text{Ellipse Equation:}$$

$$E(u,v) = k$$

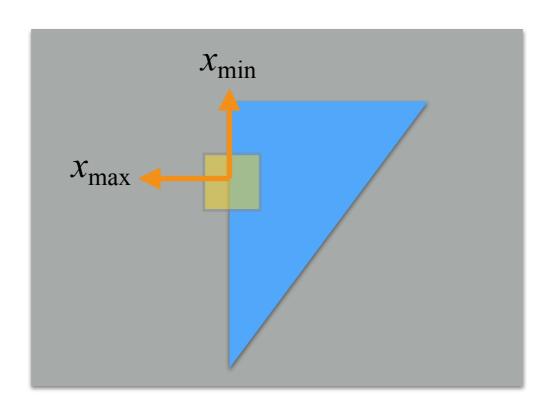
$$M = \sum_{x_k, y_k \in W(x,y)} \begin{bmatrix} I_x^2(x_k, y_k) & I_x(x_k, y_k)I_y(x_k, y_k) \\ I_x(x_k, y_k)I_y(x_k, y_k) & I_y^2(x_k, y_k) \end{bmatrix}$$

Harris Corner Detector: Second Moment Matrix

- M reveals information about the distribution of gradients around a pixel.
- The eigenvectors of *M* identify the directions of fastest and slowest change.



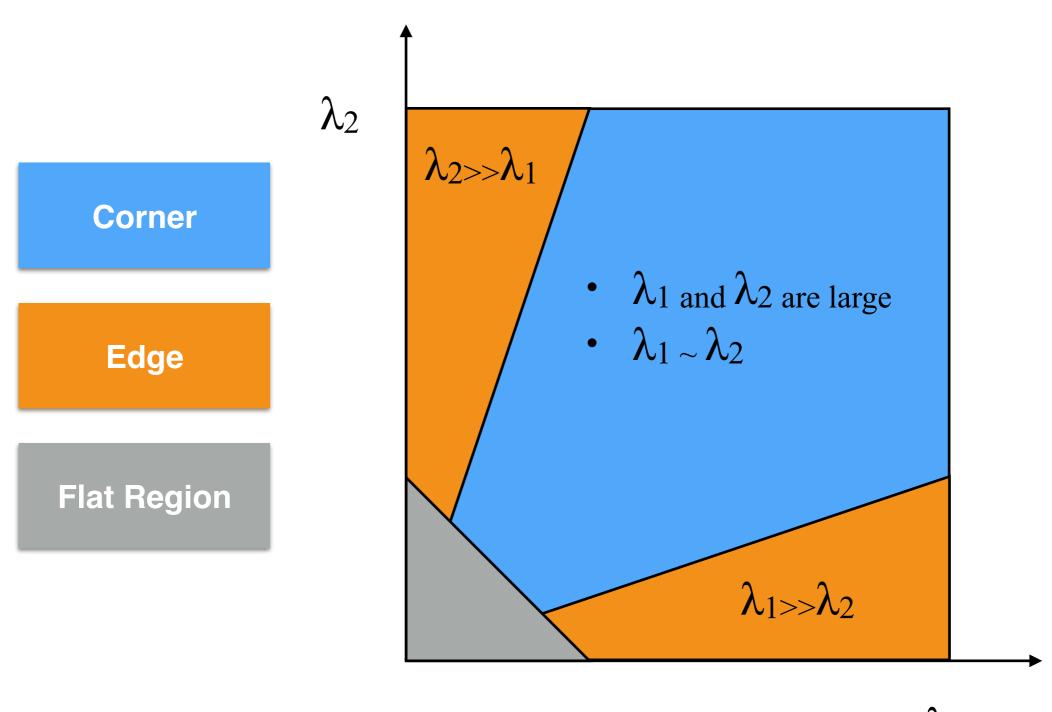
Harris Corner Detector: Second Moment Matrix



Eigenvalues and eigenvectors of M define shift directions with the smallest and largest change in E:

- x_{max} = direction of largest increase in E
- λ_{max} = amount of increase in direction x_{max}
- x_{\min} = direction of smallest increase in E
- λ_{\min} = amount of increase in direction x_{\min}

Classification



Harris Corner Detector: Corners Measure

 Instead of directly computing the eigenvalues, we use a measure that determines the "cornerness" of a pixel (i.e., how close to be a corner is):

$$R = \text{Det}(M) - k\text{Tr}(M)^{2}$$
$$\text{Det}(M) = \lambda_{1}\lambda_{2}$$
$$\text{Tr}(M) = \lambda_{1} + \lambda_{2}$$

• *k* is an empire constant with values [0.04 0.06].

Harris Corner Detector: Cornerness Measure



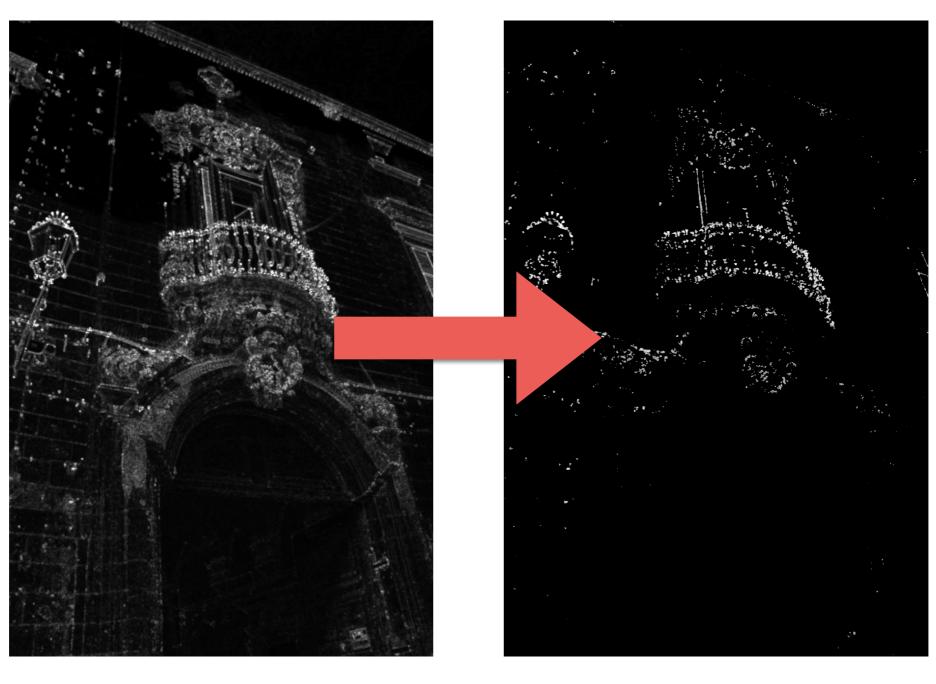
Input Image



K

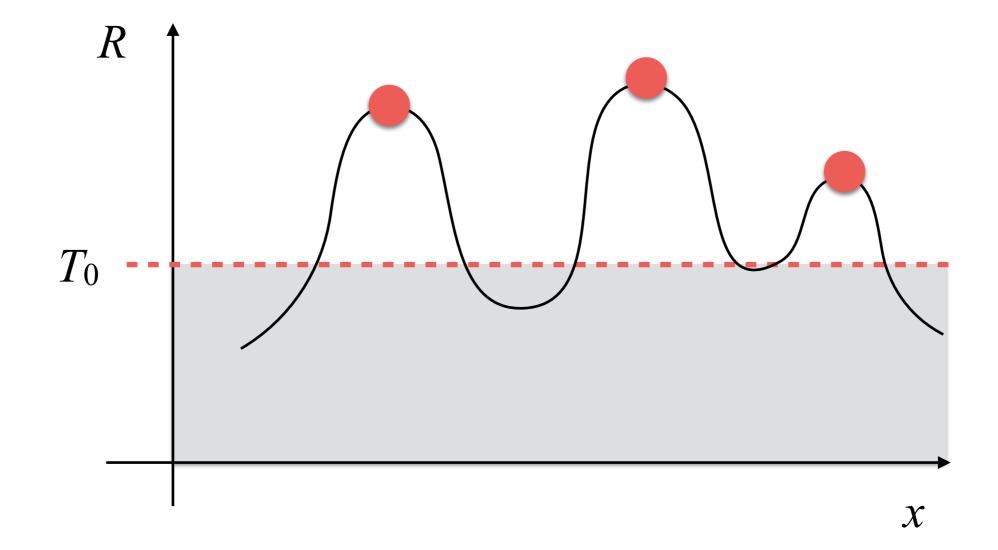
- We have to find pixels with large corner response, R, i.e., $R > T_0$.
 - Typically, T_0 in [0,1] depends on the number of points we want to extract; a default value is 0.01.

Harris Corner Detector: Thresholding

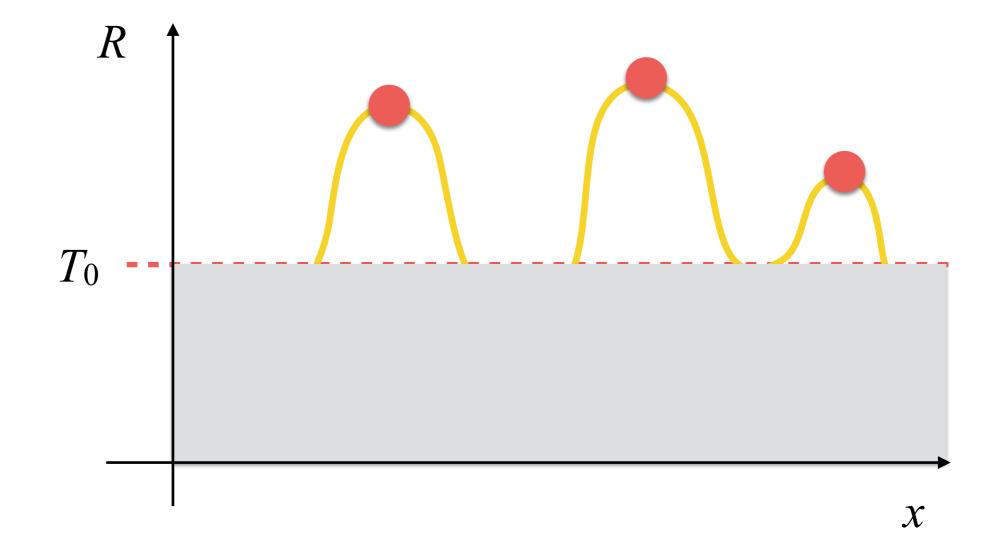


R after thresholding

 At this point, we need to suppress/remove values that are not maxima.



 At this point, we need to suppress/remove values that are not maxima, but they are over the threshold (yellow pixels).



- We set a radius (in pixel) for suppressing non-maxima;
 e.g., 3-5.
- We apply to R a maximum filter; it is similar to a median filter, but it computes the maximum instead of the median. After this we obtain a filtered image called R_{max} .
- A pixel at position (x, y) is a local maximum if and only if:

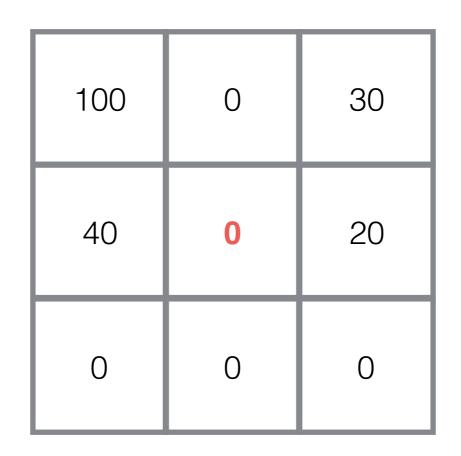
$$R_{\max}(x,y) = R(x,y) \wedge R(x,y) > T_0$$

100	0	30
40	20	20
0	0	0

The current pixel that we are evaluating is the central one! $T_0 = 5$

100	0	30
40	20	20
0	0	0

The maximum is 100!



20 < 100 so it has to be suppressed; i.e., set to 0!

Harris Corner Detector: Pruning Corners Example 2

20	0	30
40	100	20
0	0	0

The current pixel that we are evaluating is the central one! $T_0 = 5$

Harris Corner Detector: Pruning Corners Example 2

20	0	30
40	100	20
0	0	0

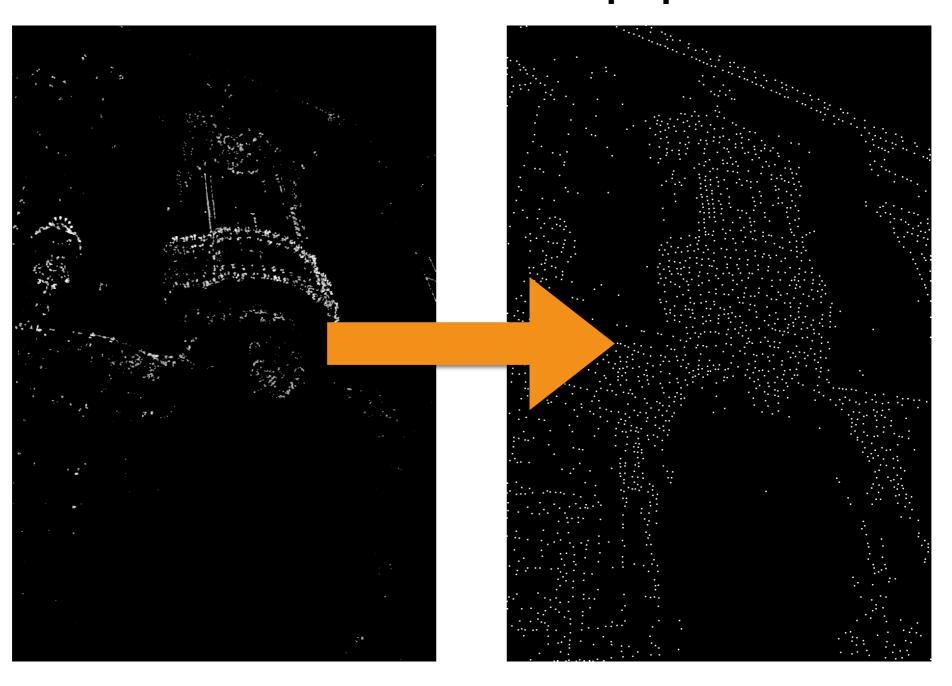
The maximum is 100!

Harris Corner Detector: Pruning Corners Example 2

20	0	30
40	100	20
0	0	0

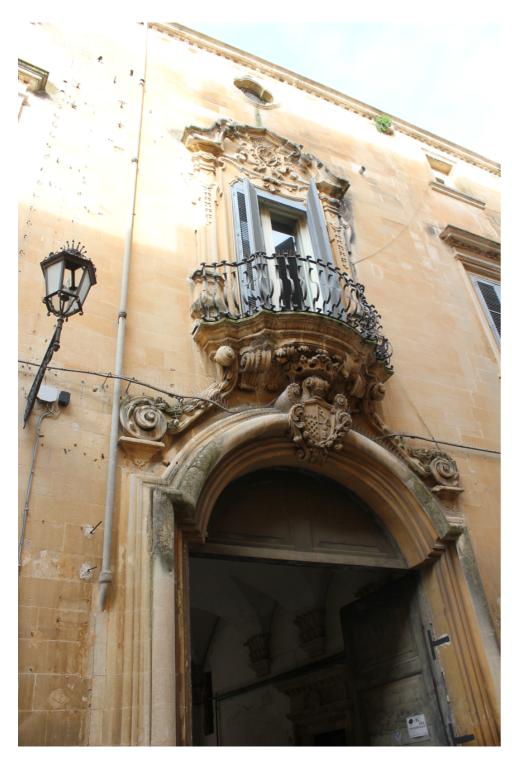
100 == 100 so it has to be kept!

Harris Corner Detector: Non-Maximal Suppression



R after thresholding Non-Maximal Suppression

Harris Corner Detector: Non-Maximal Suppression



Harris Corner Detector: Non-Maximal Suppression



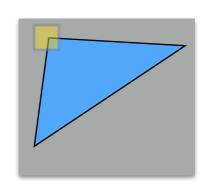
Harris Corner Detector: Non-Maximal Suppression

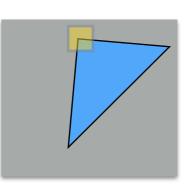




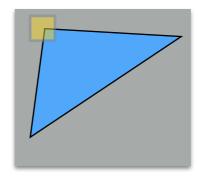
Harris Corner: Advantages

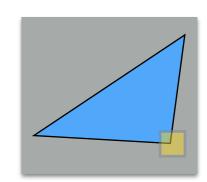
Translational invariance:





Rotation invariance:

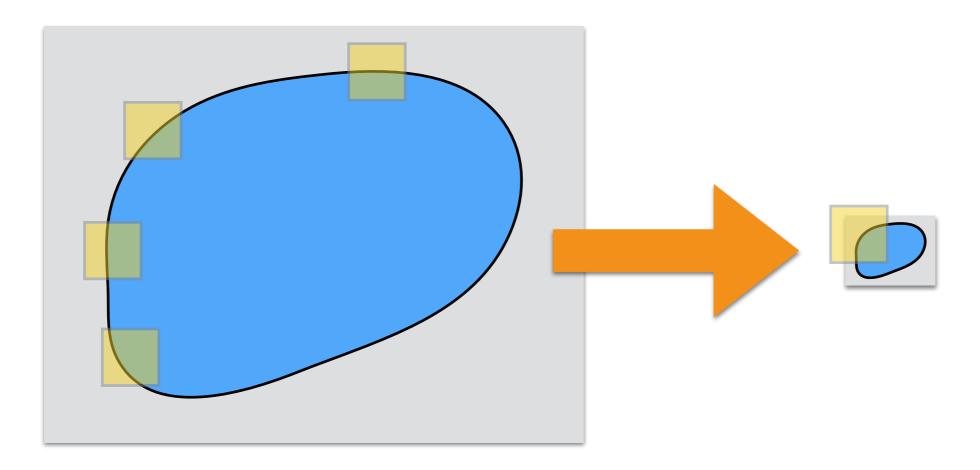




- Only derivatives are employed:
 - Intensity shift invariance: I' = I + b
 - Intensity scale invariance: I' = I a

Harris Corner: Disadvantage

Not scale invariant!



All points are classified as edges

It is now a corner!

The same feature in different images can have different size!

The Scale Problem

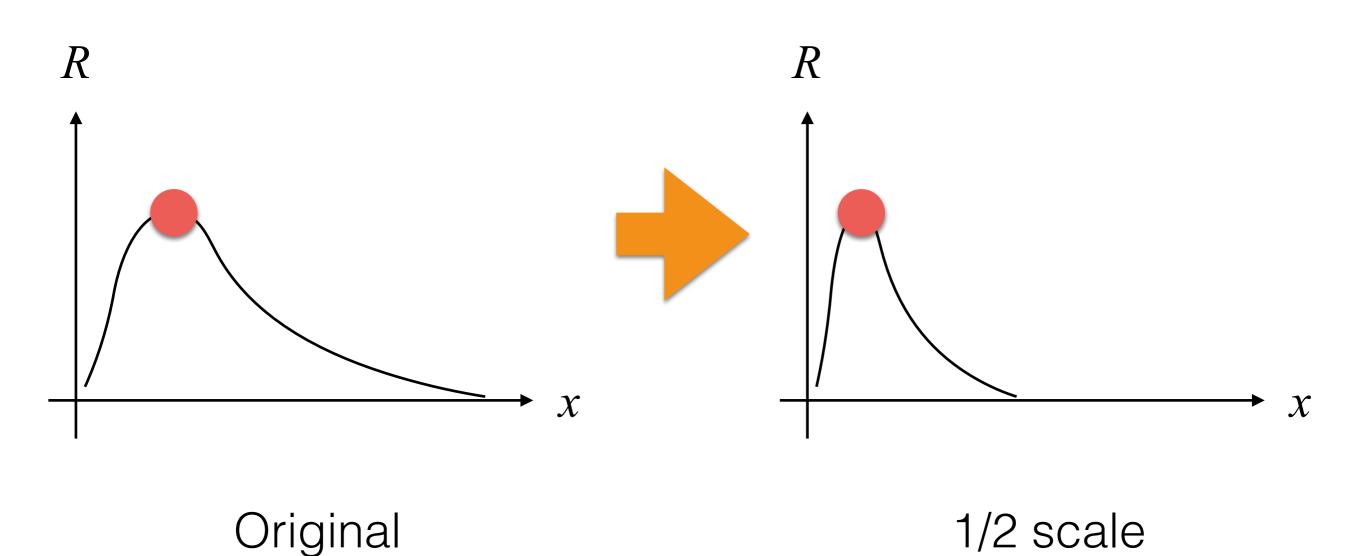




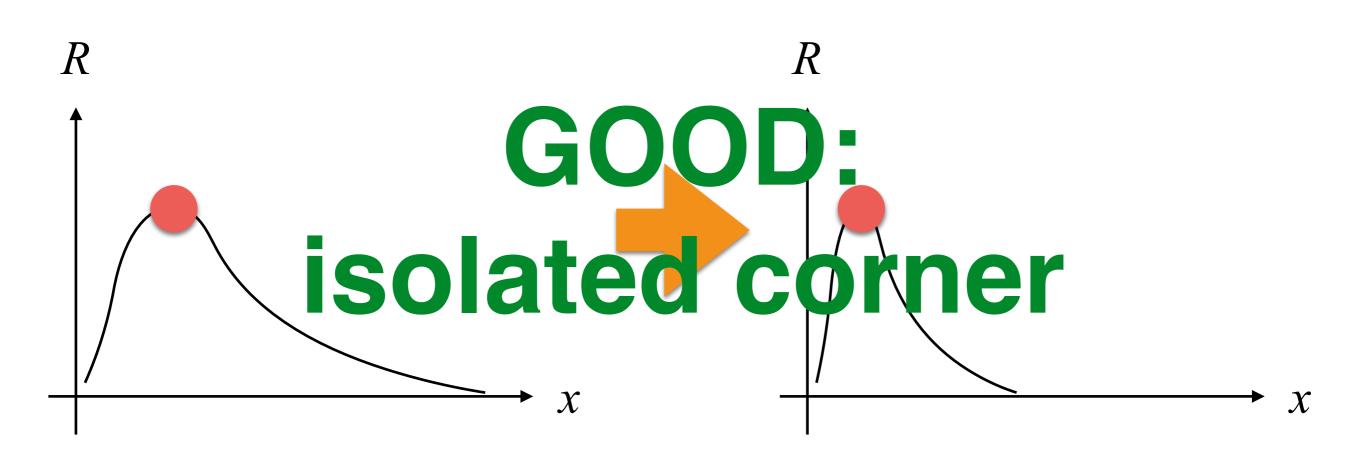
Near Object

Far Object

Scale Invariant: Stable Corners



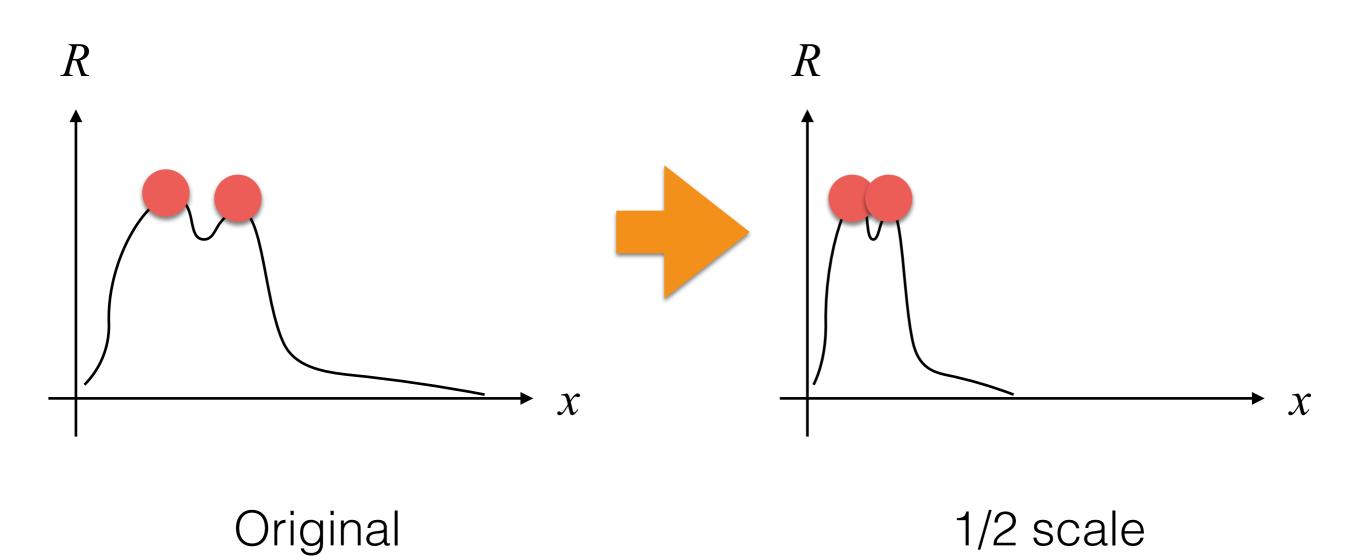
Scale Invariant: Stable Corners



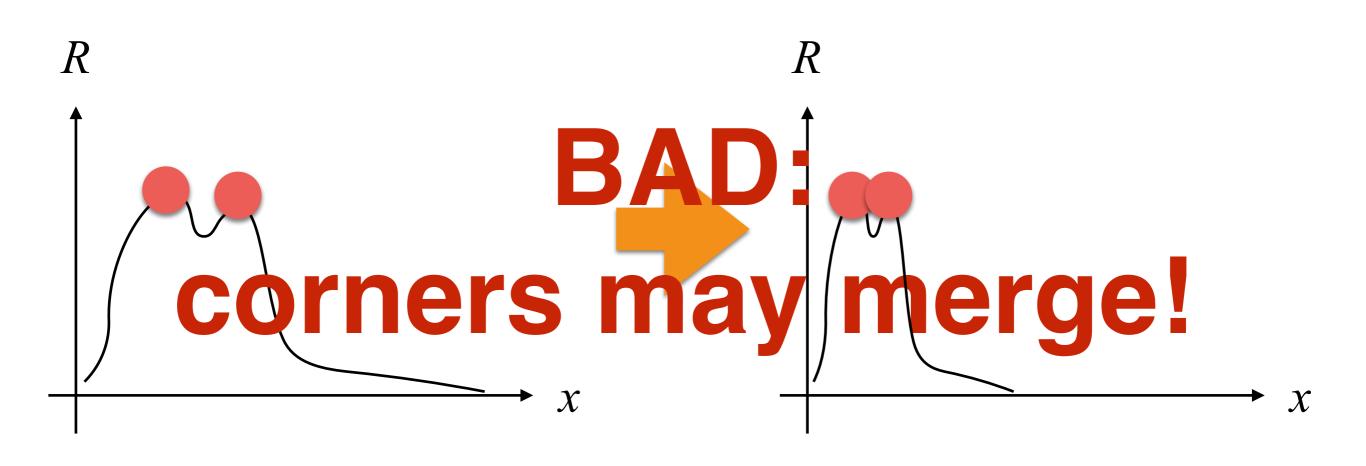
1/2 scale

Original

Scale Invariant: Unstable Corners



Scale Invariant: Unstable Corners



Original

1/2 scale

Scale Invariant: A Multi-Scale Approach

- Depending on the content of the image:
 - We need to detect the scale of corner.
 - We need to use its scale to vary the size of the window W for computing corners!

Scale Invariant: The Signature Function

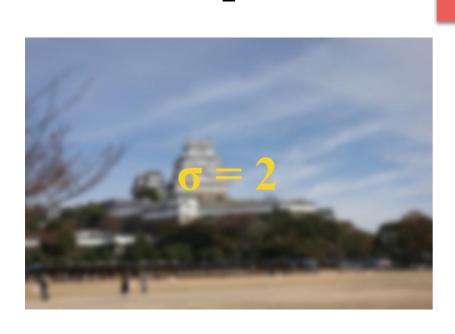
- A signature function, s, is a function giving us an idea of the local content of the image, I, around a point with coordinates (x, y) at a given scale σ .
- An example of signature function is the Difference of Gaussians (DoG):

$$s(I, x, y, \sigma) = [I \otimes G(\sigma)](x, y) - [I \otimes G(\sigma \cdot 2)](x, y)$$

where G is a Gaussian kernel.

Scale Invariant: The Signature Function







DoG





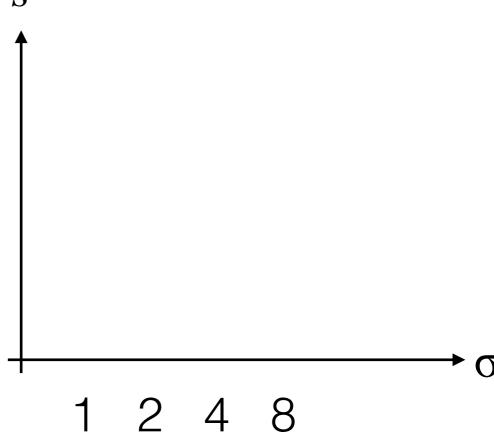
We need to find the right scale for resizing W for each image!

- The signature function, s, can give us an idea of the content of the image.
- Therefore, we need to find a maximum point of s for pixel of an input image!



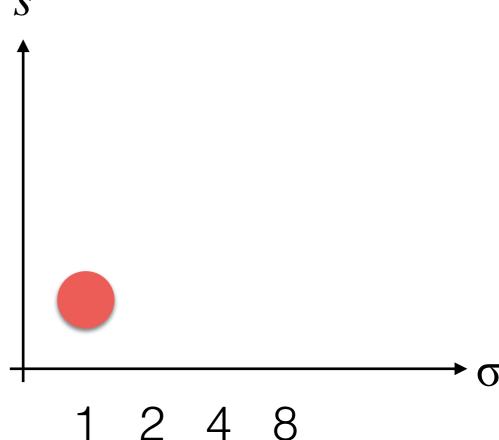
Let's build s at the red point!





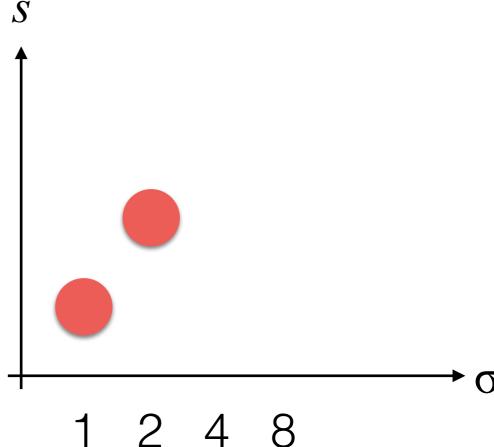
This is our start!





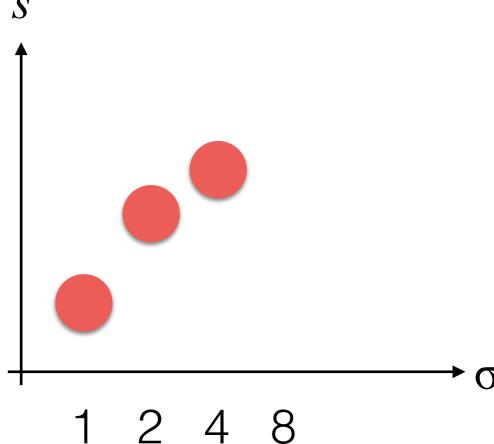
$$\sigma = 1$$





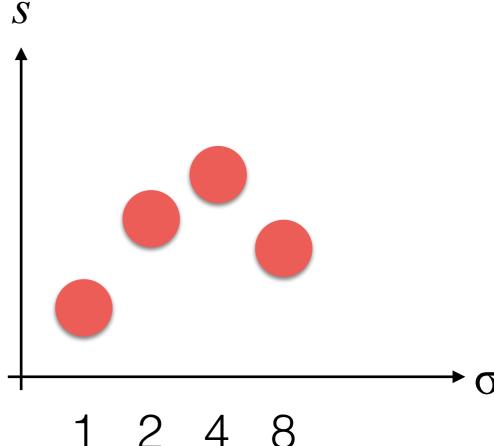
$$\sigma = 2$$





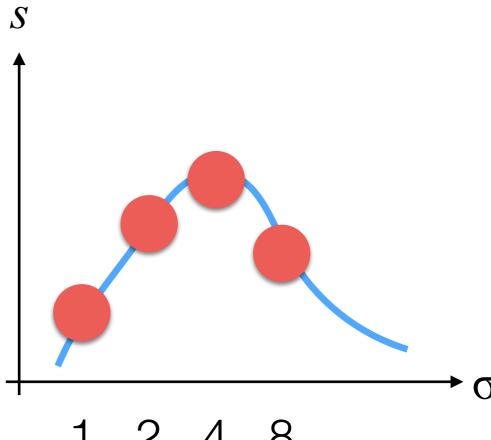
$$\sigma = 4$$



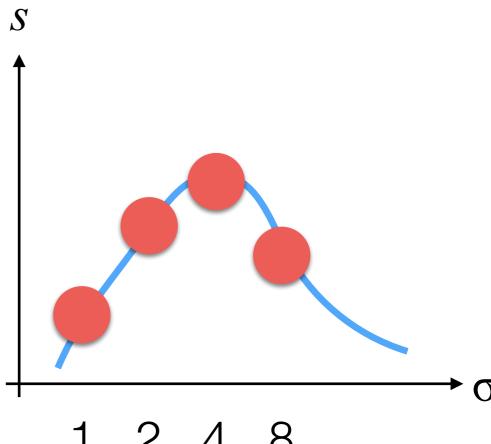


 $\sigma = 8$



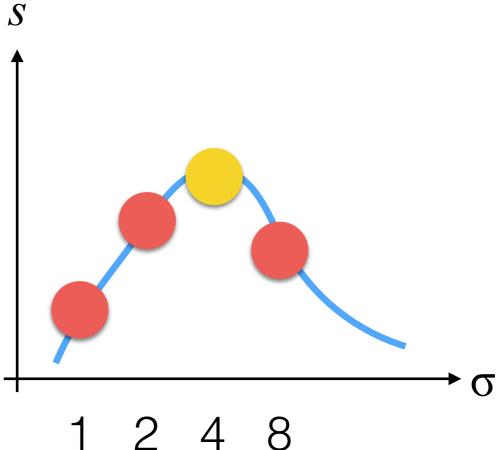






Which is σ for which s is the maximum?

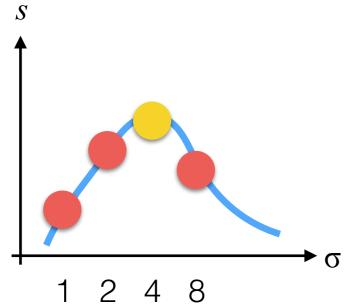


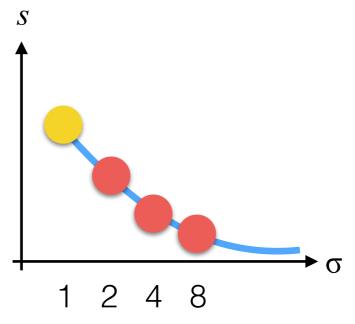


It is
$$\sigma = 4$$









Extraction of Features

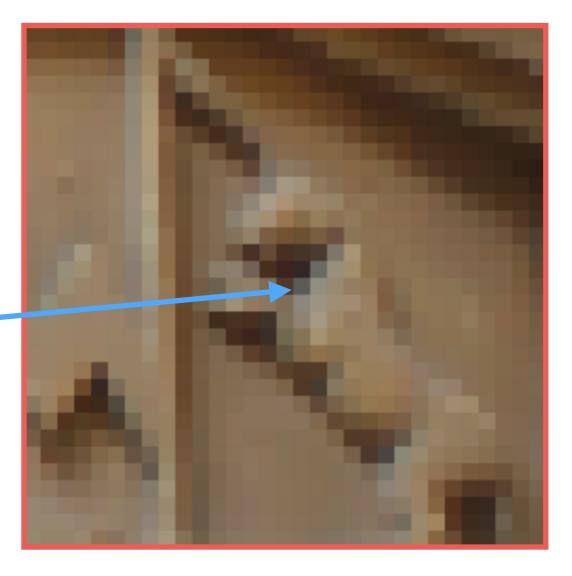
- General overview:
 - We compute the scale for each pixel using the sigma value at which we have the maximum value of the signature function.
 - We compute the Harris Corner using the scale to increase the size of the local window; i.e., the scale of the window will be multiplied by the sigma value.

 Once we found our features (i.e., corners), we need to describe in a meaningful and possibly unique way.

Why?

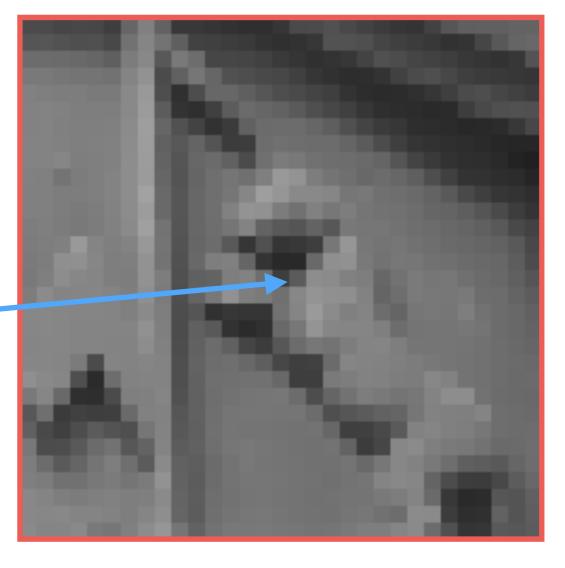
 We want compare corners between images in order to find correspondences between images.





A patch, P, is a sub-image centered in a given point (u, v).





A patch, P, is a sub-image centered in a given point (u, v).

- There are many local features descriptors in literature:
 - BRIEF/ORB descriptor.
 - SIFT descriptor.
 - SURF descriptor.
 - etc.

- Good properties that we want are invariance to:
 - Illumination changes.
 - Rotation.

BRIEF Descriptor

• The descriptor creates a vector of *n* binary values:

BRIEF
$$(P) = \mathbf{b} = [0, 1, 0, 0, \dots, 1]^{\top}$$

• For efficiency, it is encoded as a number:

$$n_{\mathbf{b}} = \sum_{I=1}^{n} 2^{i-1} b_i$$

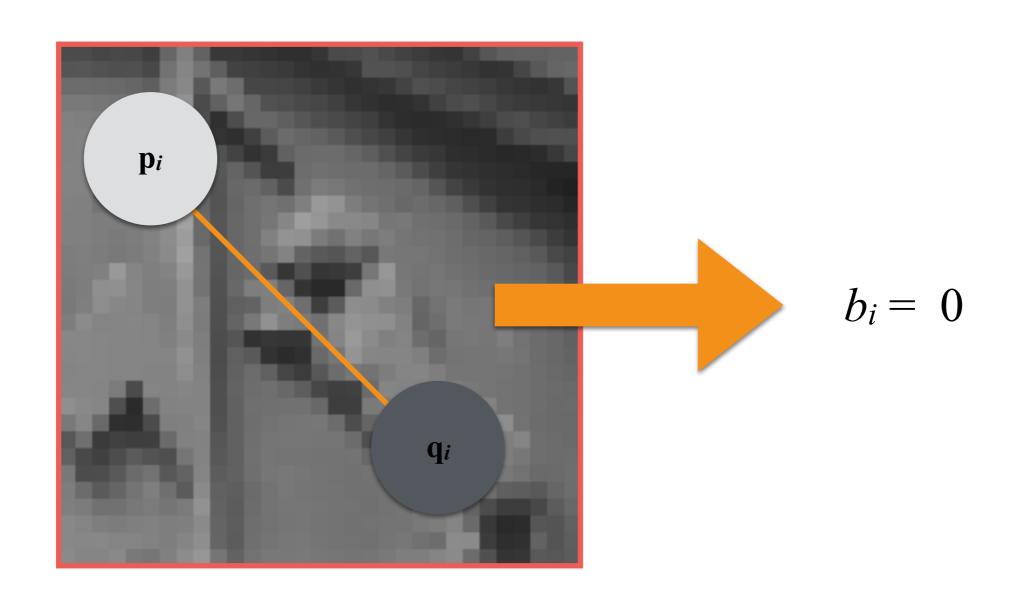
BRIEF Descriptor

 Given a patch, P, of size SxS an element of b is defined as

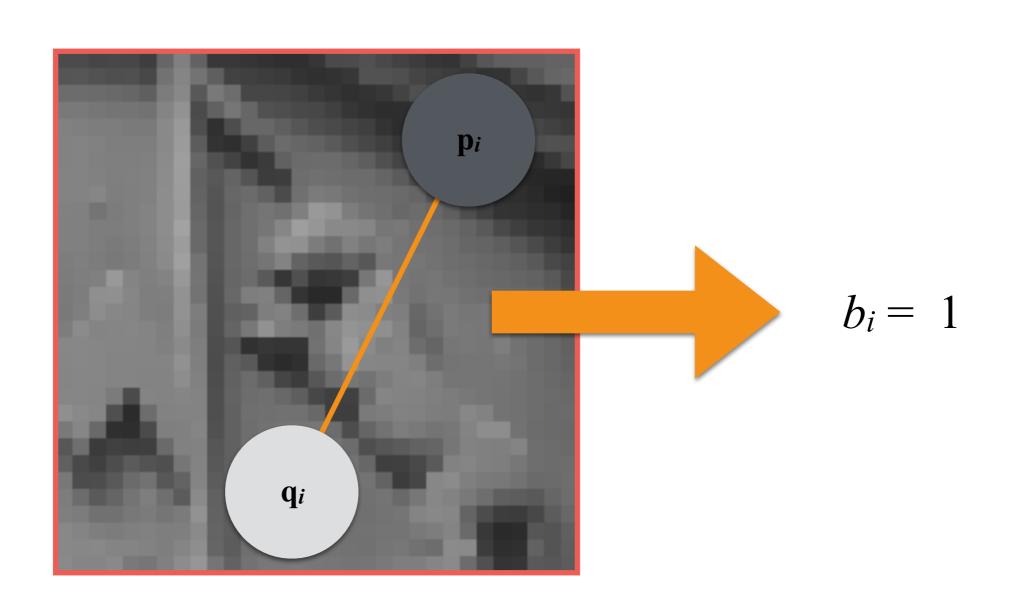
$$b_i(\mathbf{q}_i, \mathbf{p}_i) = \begin{cases} 1 & \text{if } P(\mathbf{p}_i) < P(\mathbf{q}_i), \\ 0 & \text{otherwise} \end{cases}$$

• where \mathbf{p}_i and \mathbf{q}_i are the coordinates (x, y) of two random points in P.

BRIEF Descriptor: Example



BRIEF Descriptor: Example



BRIEF Descriptor: Test

- Let's say we have two descriptor **b**¹ and **b**². How do we check if they are describing the same corner?
- We count the number of different bits in the two vectors (Hamming distance):

$$D_H(\mathbf{b}^1, \mathbf{b}^2) = \sum_{i=1}^n \neg xor(b_i^1, b_i^2)$$

- Higher the closer:
 - This is a very computationally efficient distance function.

BRIEF Descriptor: Test

A	В	A XOR B = [(NOT A) AND B] OR [(NOT B) AND A]	NOT (A XOR B)
0	0	0	1
0	1	1	0
1	0	1	0
1	1	0	1

BRIEF Descriptor: Point-Set

- The optimal number of points' couple (i.e., the size of the descriptor; *n*) is **256**.
- This value was computed from experiments testing different lengths: 16, 32, 64, 128, 256, and 512.
- Points can be generated in different ways:
 - Uniform distribution in the patch
 - $(\mathbf{p}_i, \mathbf{q}_i) \sim \text{ i.i.d. Gaussian} \left(0, \frac{S^2}{25}\right)$

BRIEF Descriptor: Point-Set

Points are pre-computed, only once, generating a set:

$$A = \begin{bmatrix} \mathbf{p}_0, & \mathbf{p}_1, & \dots & \mathbf{p}_n \\ \mathbf{q}_0, & \mathbf{q}_1, & \dots & \mathbf{q}_n \end{bmatrix}$$

- This set is always used for the extraction of all descriptors in all photos!
 - If this is not done, we cannot do comparisons because we are comparing different tests (e.g., comparing apples and organges):
 - We need to keep consistency

- Advantages:
 - Computationally fast.
 - Invariant to illumination changes.
 - Compact!
 - Patent free.
- Disadvantage:
 - Rotation is an issue:
 - The method can handle rotations up to 10-15 degrees only.

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 - Patent free.
- Disadvantage:
 - Rotation is an issue:



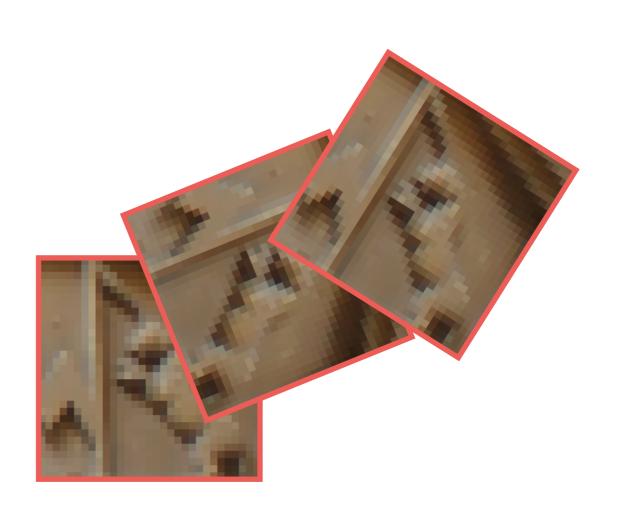
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- Disadvantage:
 - Rotation is an issue:
 - The method can handle rotations up to 10-15 degrees only.



ORB Descriptor

- The descriptor is a modified version of BRIEF and it can handle rotations!
- The first step of the algorithm is to compute the orientation of the current patch *P*.
- Idea: we determine the "center of mass" of the image, and we compute the angle between this "center of mass" and the center of the patch. This is a hint for the orientation of the patch.

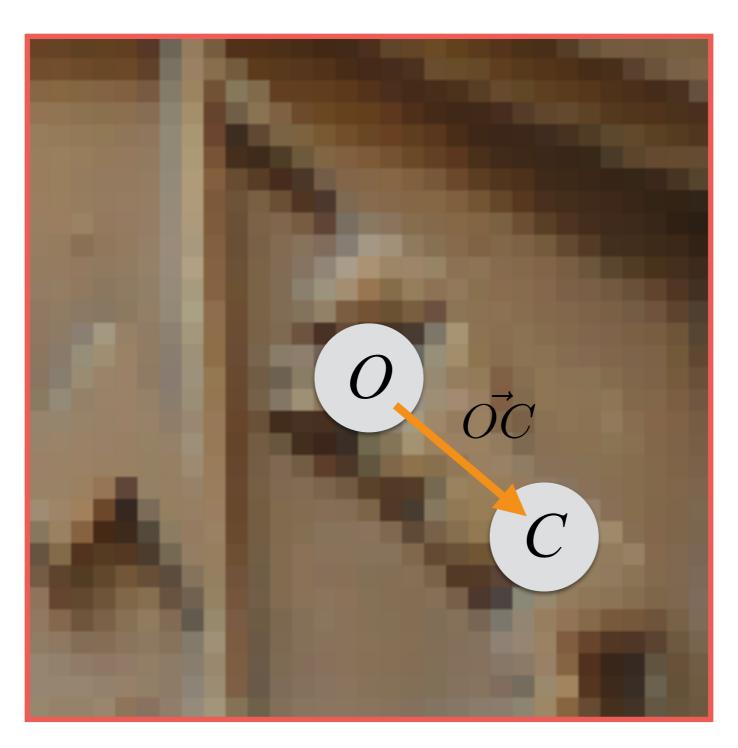
 We compute the patch orientation using Rosin moments of a patch:

$$m_{a,b} = \sum_{x,y \in P} x^a y^b P(x,y)$$

• From this, we define the centroid, *C*, as

$$C = \left(\frac{m_{1,0}}{m_{0,0}}, \frac{m_{0,1}}{m_{0,0}}\right)$$

 Now, we can create a vector from corner's center, O, to the centroid, C. This allows us to calculate the angle of rotation.

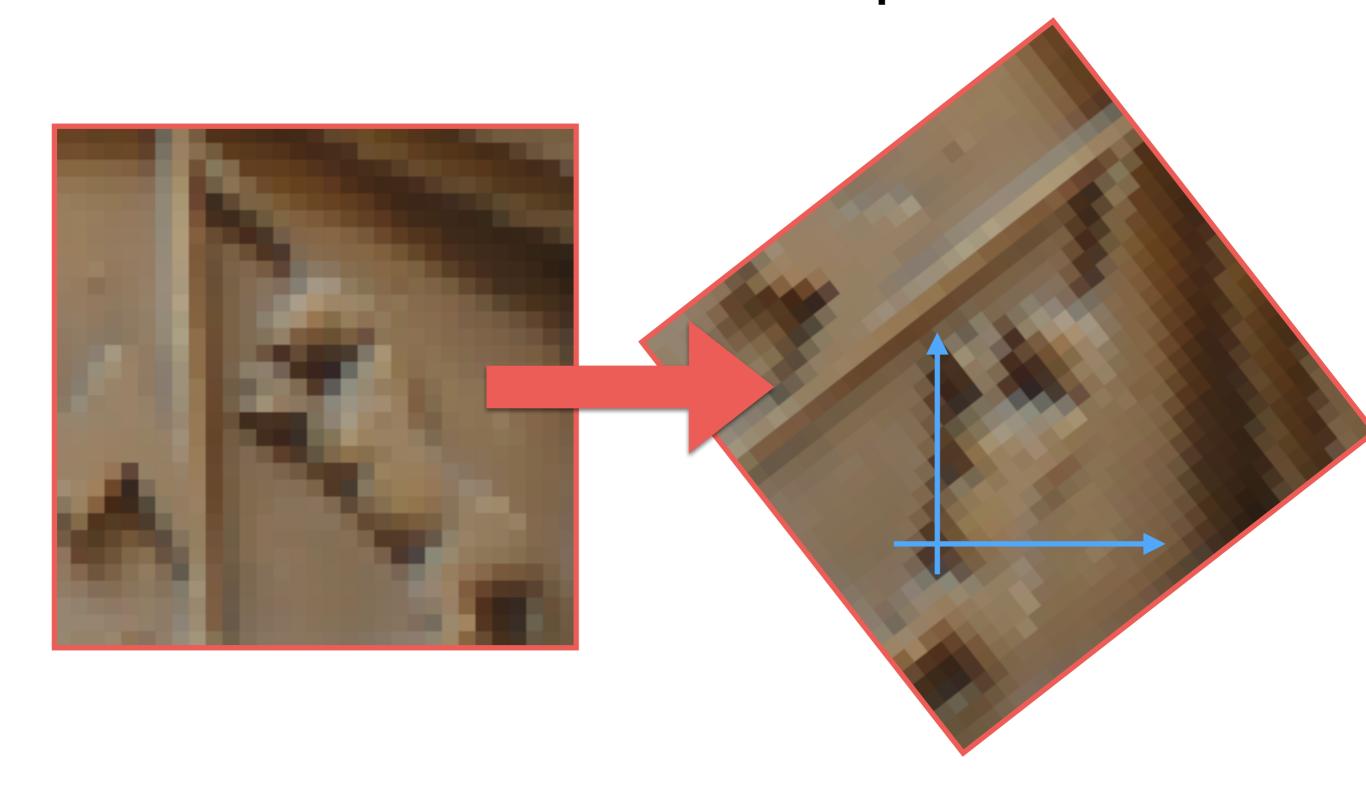


 From this vector, the orientation of the patch can be computed simply as

$$\theta = \text{atan2}(m_{0,1}, m_{1,0})$$

- From this, we can rotate the patch *P*, but this operation is very computationally expensive:
 - We need to rotate each single point in the patch!

ORB Descriptor



 Instead of rotating the whole patch, we can rotate only the points stored in A as

$$A_{\theta} = R_{\theta} \cdot A$$

- where R_{θ} is a 2D rotation matrix.
- NOTE: we need to rotate less points!

ORB Descriptor

- Advantages:
 - Computationally fast.
 - Invariant to illumination changes.
 - Compact!
 - Invariant to rotation.
 - Patent free.
- Disadvantage:
 - Not robust as SIFT.

- It is the state-of-the-art descriptor.
- It was introduced in 1999, but it is still the king.

- The first step is to compute the orientation of P.
- We compute the horizontal (P_x) and vertical (P_y) gradients of the P.
- For each pixel at coordinates (i, j) in the patch we compute its orientation and magnitude:

$$m(i,j) = \sqrt{P_x(i,j)^2 + P_y(i,j)^2}$$

$$\theta(i,j) = \operatorname{atan2}\left(P_y(i,j), P_x(i,j)\right)$$

- A histogram, H, of directions is created for each orientation taking into account its magnitude.
- We repeat this process for all gradients in the patch!
 - Note that H is initialized as a vector of zeros.

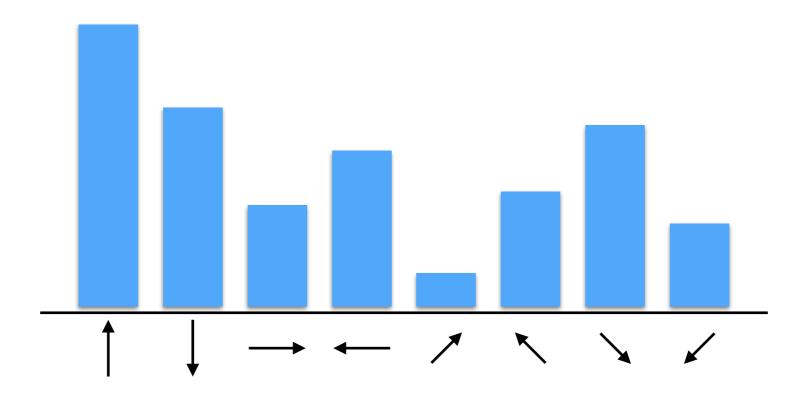
- Let's say, we have a histogram H with 18 bins (b = 18).
 - This means each bin has a size (k) in degree of 20° (k = 360 / b = 360 / 18).
- Now, we have to insert a gradient (m = 10 and $\theta = 45^{\circ}$) from our patch in H we need to process a gradient in the patch.
 - First, we compute the index of the bin to update:

$$i = \left\lfloor \frac{\theta}{k} \right\rfloor = \left\lfloor \frac{45}{20} \right\rfloor$$

Then, we update H as

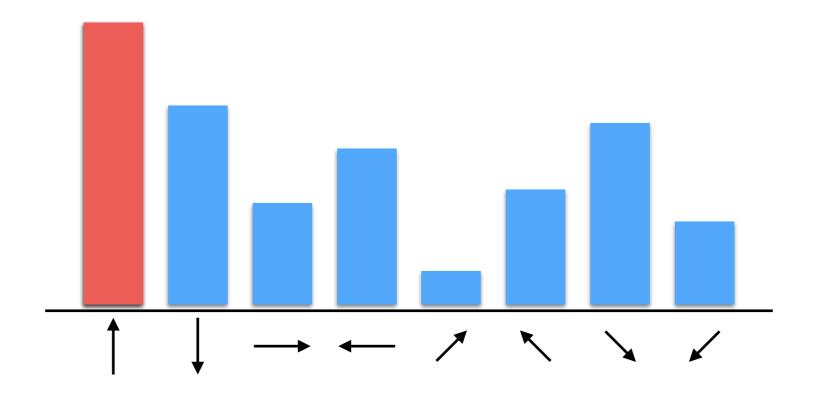
$$H(i) = H(i) + m = H(i) + 10$$

• Finally, we get this (an example with 8 bins; i.e., 8 directions):



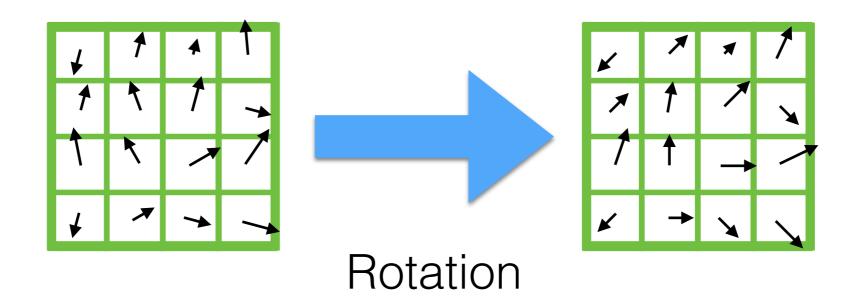
- The patch orientation, α , is given by the highest peak:
 - If we have two equal peaks, we take the as winner the first one in histogram.

• Finally, we get this (an example with 8 bins; i.e., 8 directions):



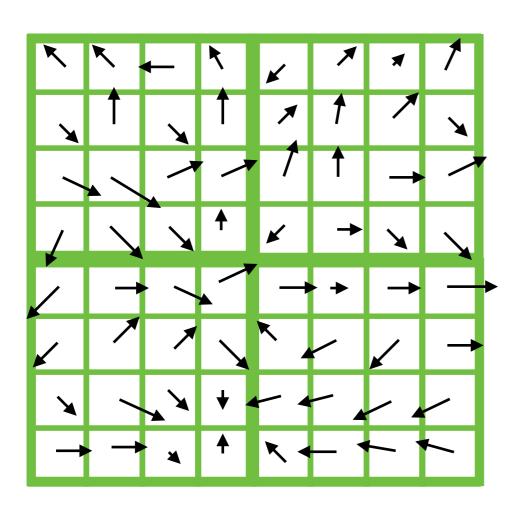
- The patch orientation, α , is given by the highest peak:
 - If we have two equal peaks, we take the as winner the first one in histogram.

- Once we have α , we can rotate all gradients in the patch using it.
 - This ensures to be invariant to rotations!

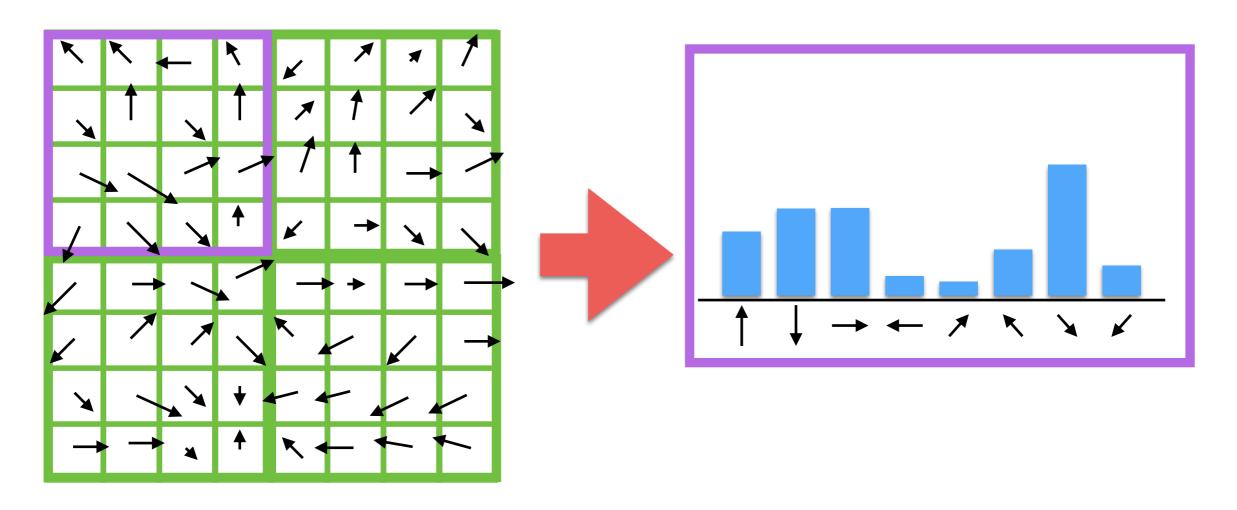


- Why do we rotate the gradients? It is computationally faster:
 - In theory, we should rotate the patch and then recompute the gradients.
 - This is computationally expensive!

- At this point, we divide the patch into 4x4 blocks.
 For each block, we compute a new histogram of directions.
- The final SIFT descriptor is the concatenation (flattening) of all these histograms.

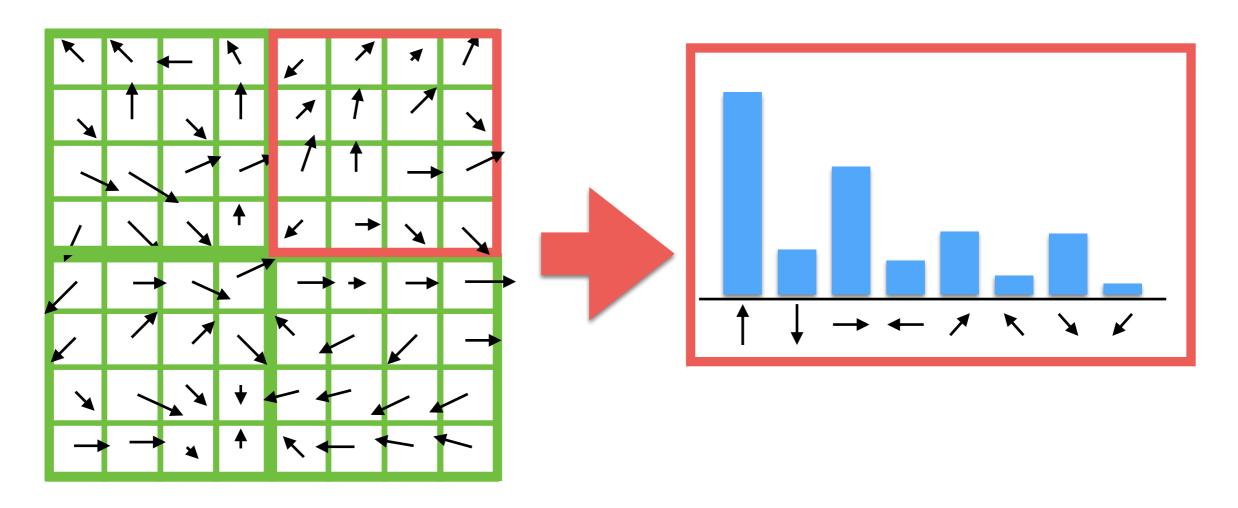


Patch and its gradients



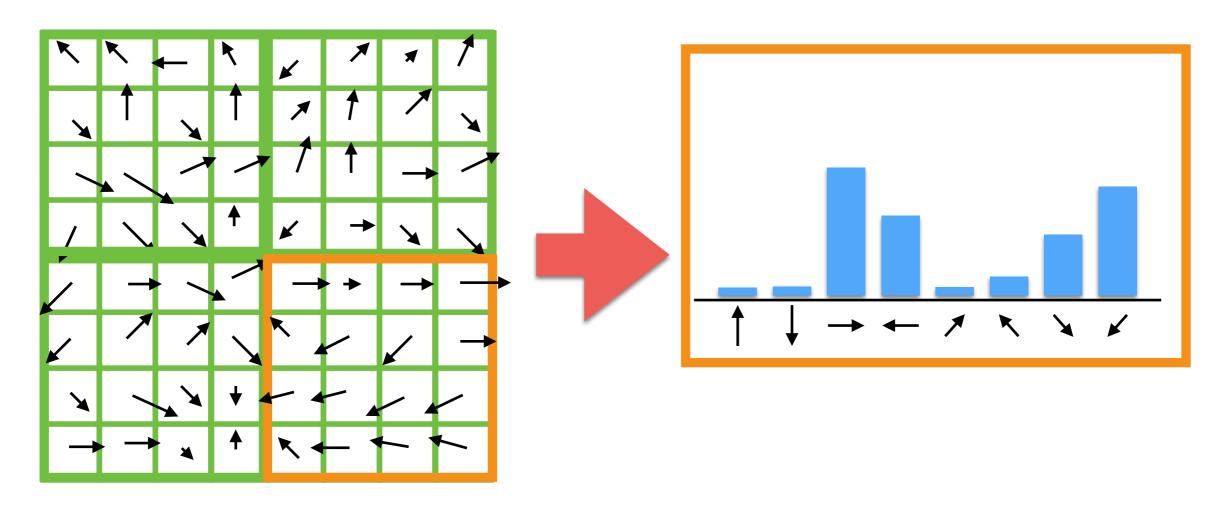
Patch and its gradients

We compute the histogram for the first block in violet



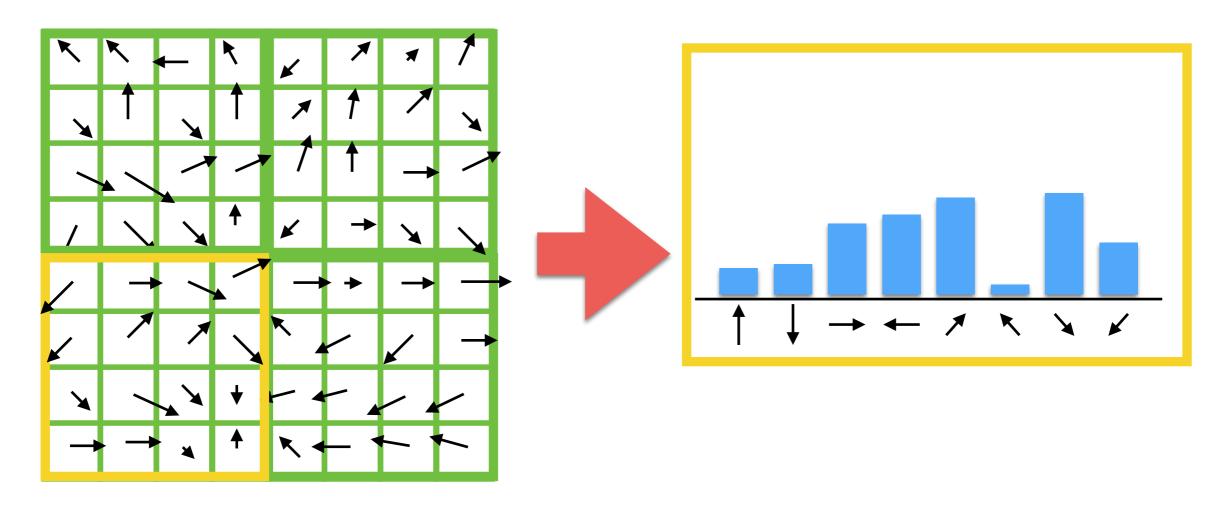
Patch and its gradients

We compute the histogram for the second block in red



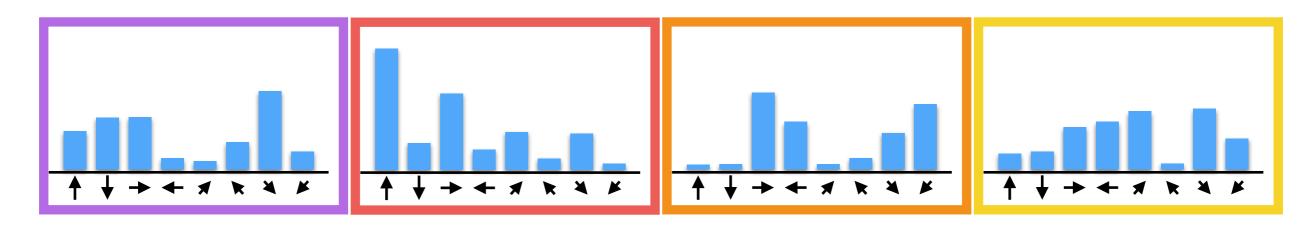
Patch and its gradients

We compute the histogram for the third block in orange



Patch and its gradients

We compute the histogram for the fourth block in yellow



The *final descriptor* is the concatenation of the histogram of all blocks. Note that this can be encoded as a vector; in this example the vector has size equal to:

$$4x8 = 32$$

4 because we have 2x2 Blocks

8 because we have 8 direction for each histogram.

SIFT Descriptor: Test

 We test the differences as distance between histograms:

$$D_2(\mathbf{h}^1, \mathbf{h}^2) = \sqrt{\sum_{i=1}^n (h_i^1 - h_i^2)^2}$$

- The lower the closer:
 - This is the opposite compared to BRIEF/ORB.

- Advantages:
 - Invariant to illumination changes.
 - Invariant to rotation.
- Disadvantages:
 - Slower than BRIEF/ORB.
 - More memory than binary methods.
 - Patented! It is patent-free from 12th of Aprile 2020!

Matching Images

Matching: An Image Against Another One

- **Input**: two descriptor lists (*they can be of equal or different size*), **desc**₁ and **desc**₂, respectively of image I_1 and I_2 .
- Output: a vector with indices of matches for each list:
 - The output is called M_{12} if we match I_1 against I_2
 - The output is called M_{21} if we match I_2 against I_1

Matching: How the Output is Encoded Example 1

- Let's say we have 4 descriptors in **desc**₁
- Let's say we have 3 descriptors in desc₂
- Let's say that we want to match I_1 against I_2 , this means that we want to compute \mathbf{M}_{12} .

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_3^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

$$M_{12} = []$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the first descriptor of **desc**₁ matches with the second descriptor of **desc**₂.

$$M_{12} = []$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the first descriptor of **desc**₁ matches with the second descriptor of **desc**₂.

$$\mathbf{M}_{12} = [2,]$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the second descriptor of **desc**₁ matches with the third descriptor of **desc**₂.

$$M_{12} = [2,]$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the second descriptor of **desc**₁ matches with the third descriptor of **desc**₂.

$$\mathbf{M}_{12} = [2, 3,]$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the third descriptor of **desc**₁ matches with the first descriptor of **desc**₂.

$$\mathbf{M}_{12} = [2, 3,]$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix}$$

We find out that the third descriptor of **desc**₁ matches with the first descriptor of **desc**₂.

$$\mathbf{M}_{12} = [2, 3, 1,]$$

$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2^1 \ d_3^1 \ d_4^1 \end{bmatrix} \qquad \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \ d_3^2 \end{bmatrix}$$

We find out that the fourth descriptor of **desc**₁ matches with the second descriptor of **desc**₂.

$$\mathbf{M}_{12} = [2, 3, 1, 2]$$

Matching: How the Output is Encoded Example 2

- Let's say we have 3 descriptors in **desc**₁
- Let's say we have 2 descriptors in desc₂
- Let's say that we match I_1 against I_2 , obtaining \mathbf{M}_{12} . Then, we match I_2 against I_1 obtaining \mathbf{M}_{21} .

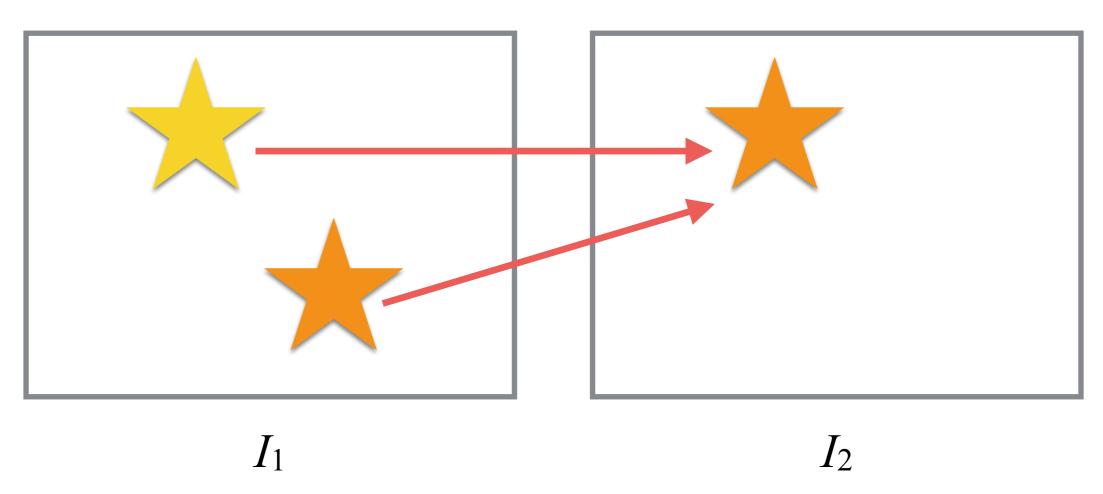
$$\mathbf{desc}_1 = egin{bmatrix} d_1^1 \ d_2 \end{bmatrix} & \mathbf{desc}_2 = egin{bmatrix} d_1^2 \ d_2^2 \end{bmatrix}$$

$$\mathbf{M}_{12} = [1,1,2] \qquad \mathbf{M}_{21} = [3,2]$$

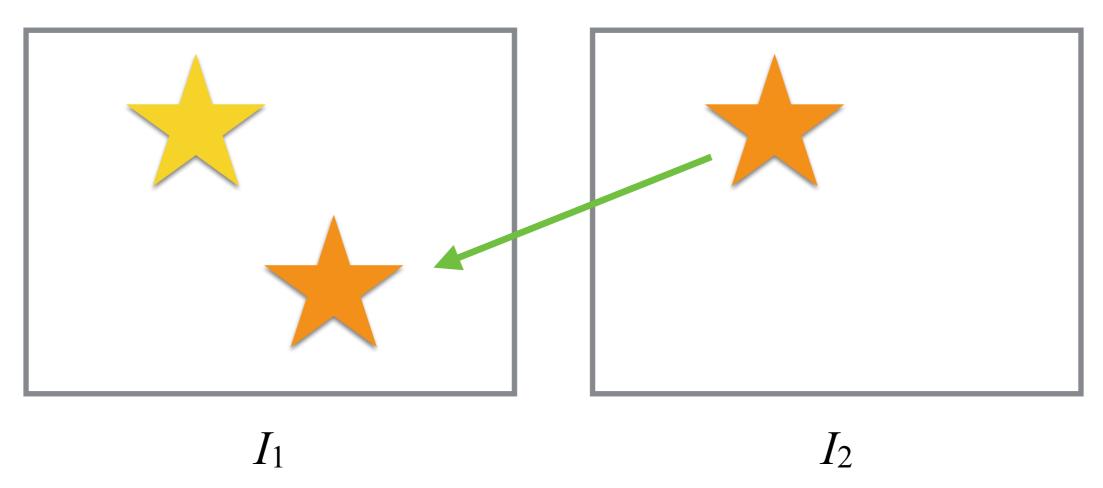
- From this example, we can notice that:
 - The matching operator is **NOT** an invertible function:
 - Therefore, M_{12} and M_{21} can be very different!
 - Why? Let's see it!



Let's say that stars in this example are feature-point.



When we match I_1 against I_2 , we have two matches; between the two stars in I_1 and the one in I_2 . This is because we need to match the yellow star with something in the other image no matter what.



When we match I_2 against I_1 , we do not have a match between the **yellow** star in I_1 and the **orange** one in I_2 because the other **orange** star in I_1 is closer than the **yellow** star!

- A simple method to find a matched descriptor in desc₂ for each descriptor in desc₁:
 - For each descriptor in desc₁, we test it against all descriptors in desc₂, and we keep as matched one the *closest* (in terms of distance; either Hamming or Euclidean).

For each descriptor \mathbf{d}^{1}_{i} in \mathbf{desc}_{1} :

```
matched(i) = -1;
matched_dist = BOTTOM;
For each descriptor \mathbf{d}^{2}_{i} in \mathbf{desc}_{2}:
  if Closer( D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{i}), matched_dist)
    matched(i) = j;
    matched_dist = D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{i});
```

D(,) is a distance function; it can be Hamming, Euclidean, etc.

endif

For each descriptor \mathbf{d}^{1}_{i} in \mathbf{desc}_{1} :

```
matched(i) = -1;
matched_dist = BOTTOM;
```

For each descriptor \mathbf{d}^{2}_{j} in \mathbf{desc}_{2} :

```
if Closer( D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{j}), matched_dist)
```

```
matched(i) = j;

matched_dist = D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{j});
```

D(,) is a distance function; it can be Hamming, Euclidean, etc.

endif

For each descriptor \mathbf{d}^{1}_{i} in \mathbf{desc}_{1} :

```
matched(i) = -1;
matched_dist = BOTTOM;
```

For each descriptor \mathbf{d}^{2}_{j} in \mathbf{desc}_{2} :

```
if Closer( D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{j}), matched_dist)
```

```
matched(i) = j;
```

```
matched_dist = D(\mathbf{d}^{1}_{i}, \mathbf{d}^{2}_{j});
```

```
endif
```

BOTTOM = +Inf for SIFT **BOTTOM** = 0 for BRIEF/ORB

D(,) is a distance function; it can be Hamming, Euclidean, etc.

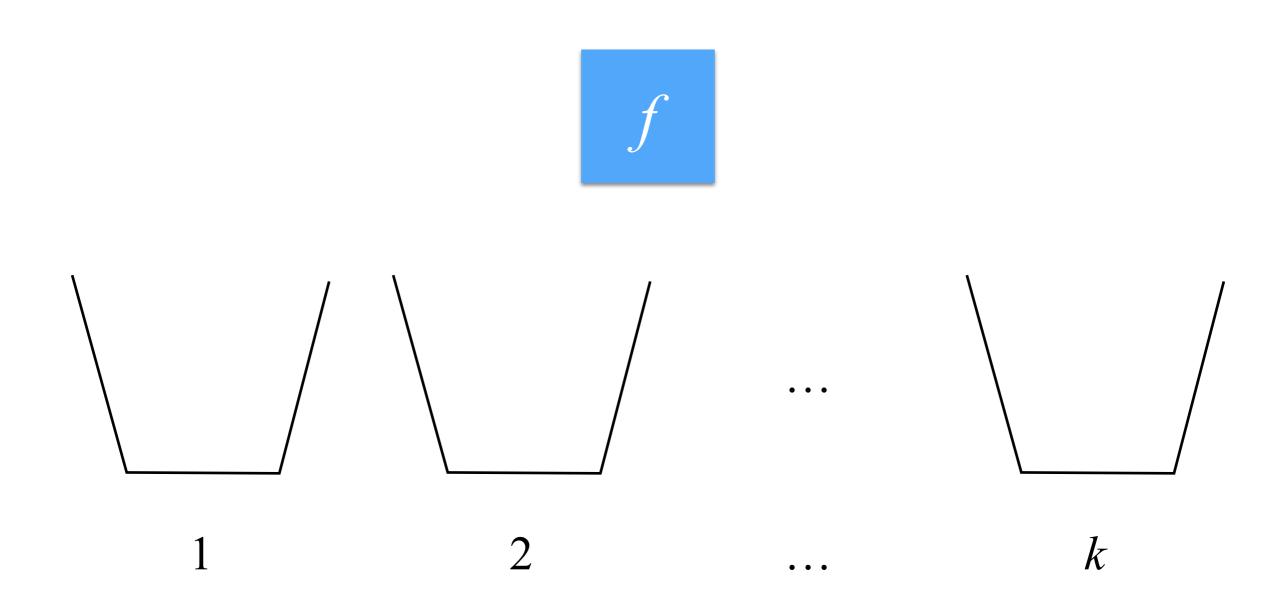
- Advantage:
 - It is exhaustive and finds the best solution!
- Disadvantage:
 - This method is very slow:
 - Let's say we have n descriptors in \mathbf{desc}_2 and n in \mathbf{desc}_2 . In the worst case, we need to compare roughly n^2 descriptors. This becomes an issue when we have more than 100 descriptors per image!

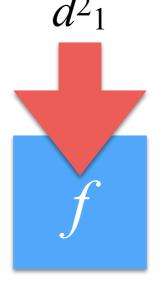
- How can we improve (approximating results)?
- Hashing: the idea is to group similar descriptors in k groups or buckets that have a constant size.

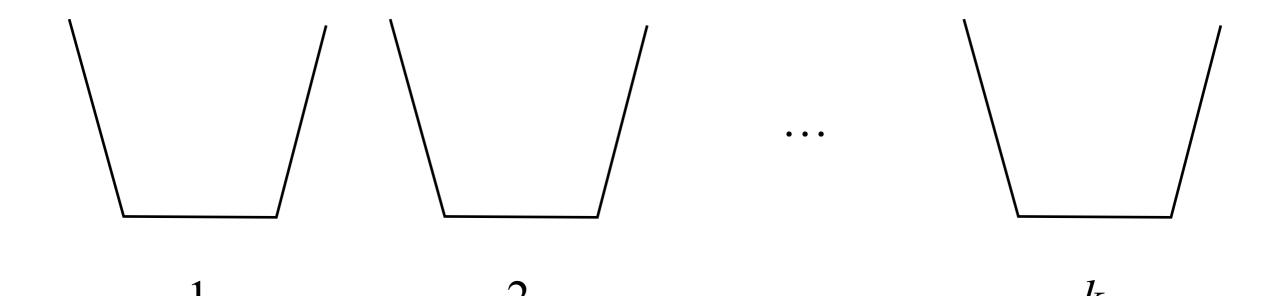
- We create k bucket.
 - Each descriptor in $desc_2$ of I_2 is assigned to a bucket using a function f, called hash function. This is defined as:

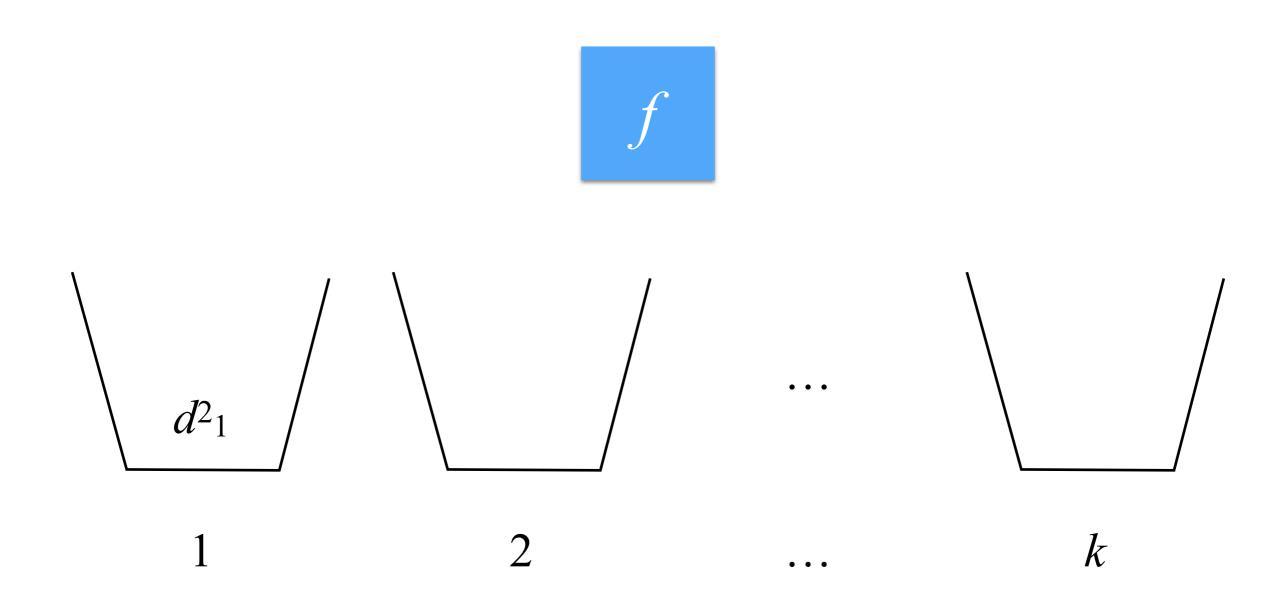
```
f: descriptor \longrightarrow [1, k] (positive integer numbers!)
```

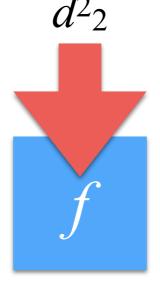
- This means that f cover generates a number in [1, k] given a descriptor.
 - For example, an *f* for BRIEF/ORB, where the descriptor is a 256-bit number, is the modulo operation.

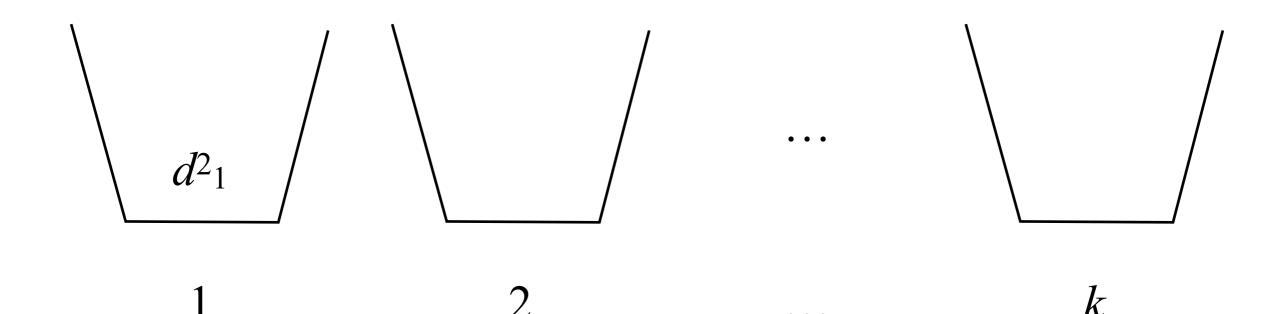


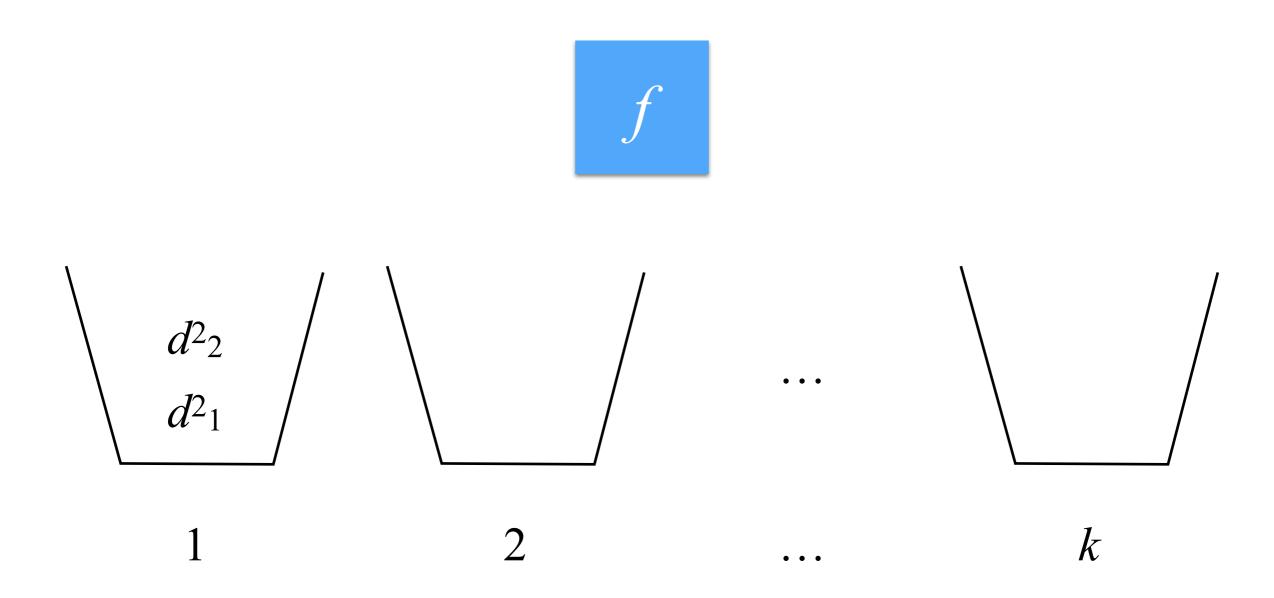




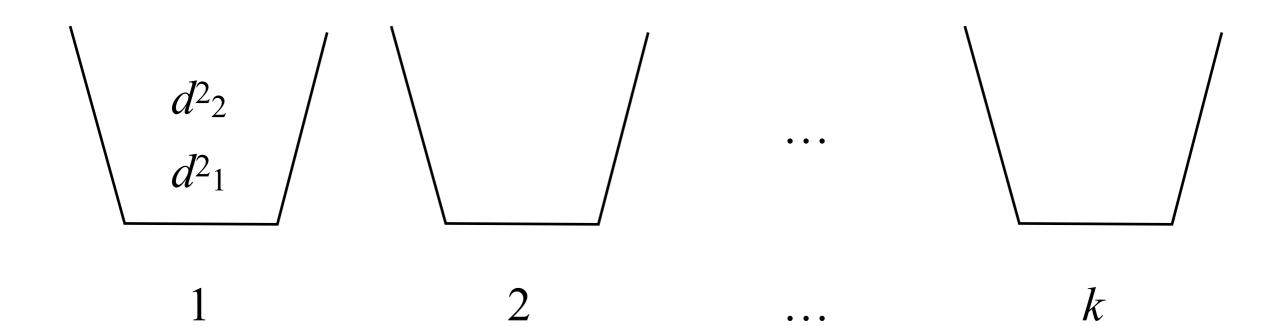


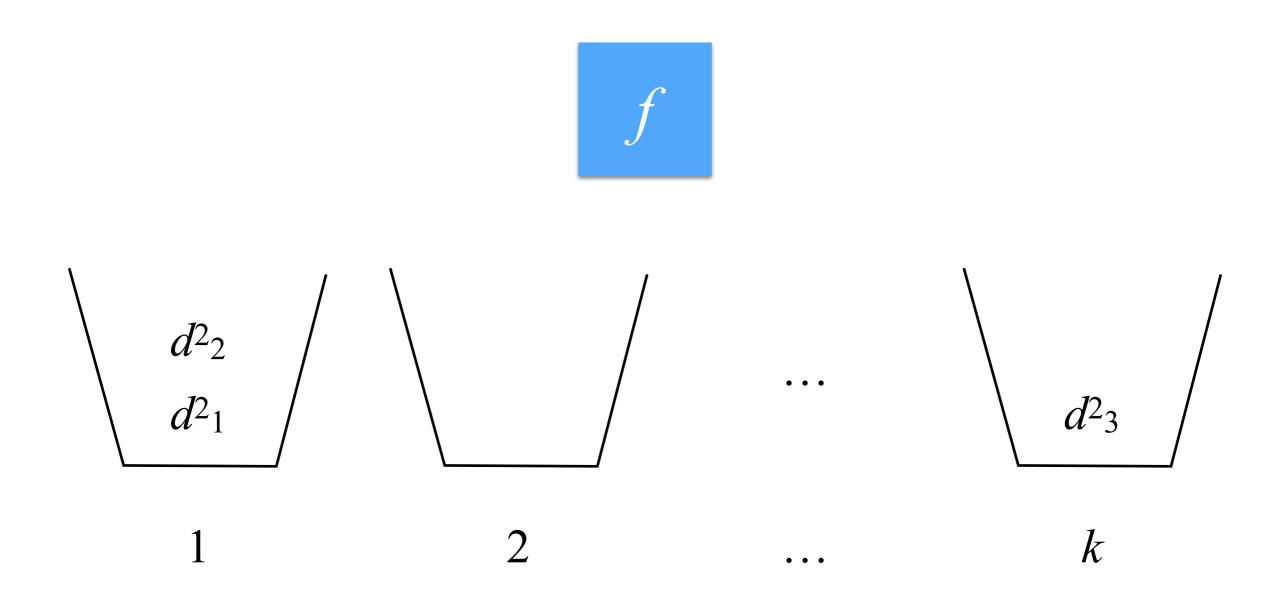






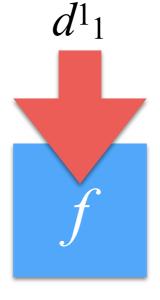


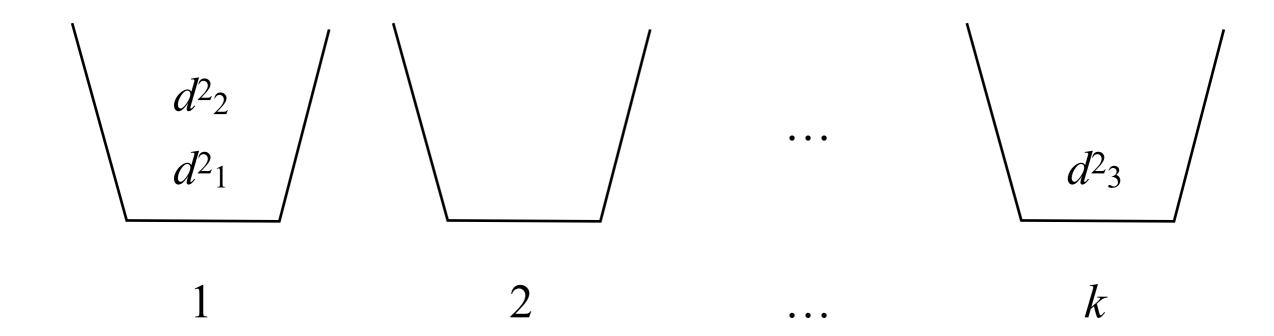


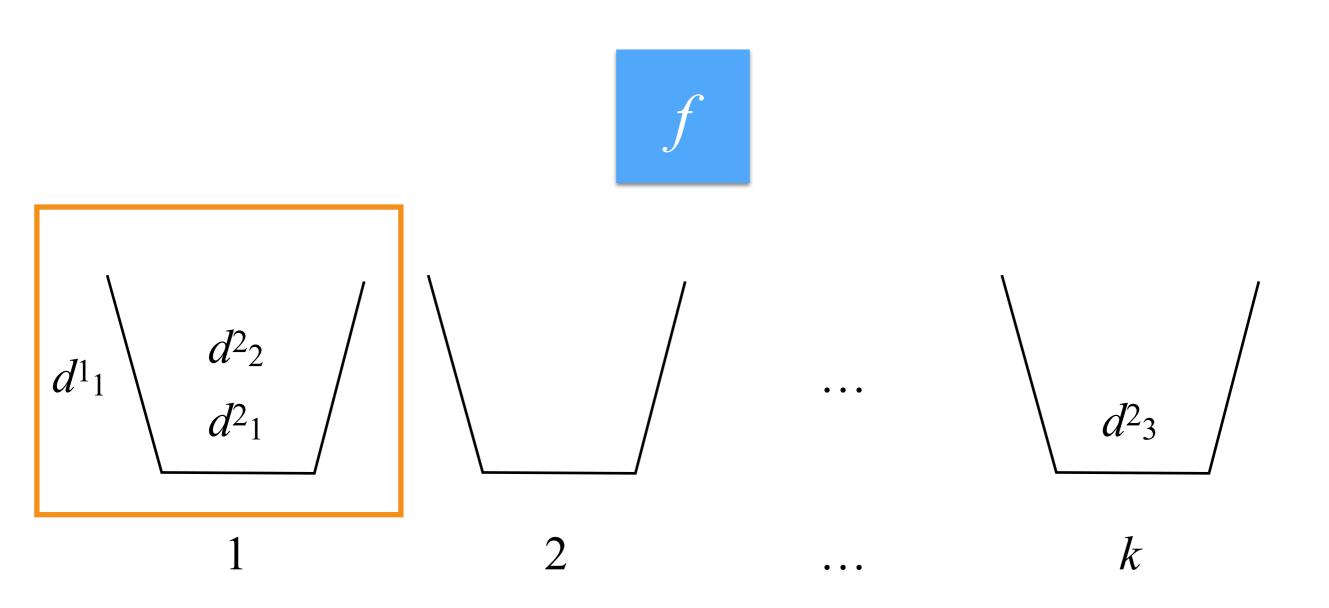


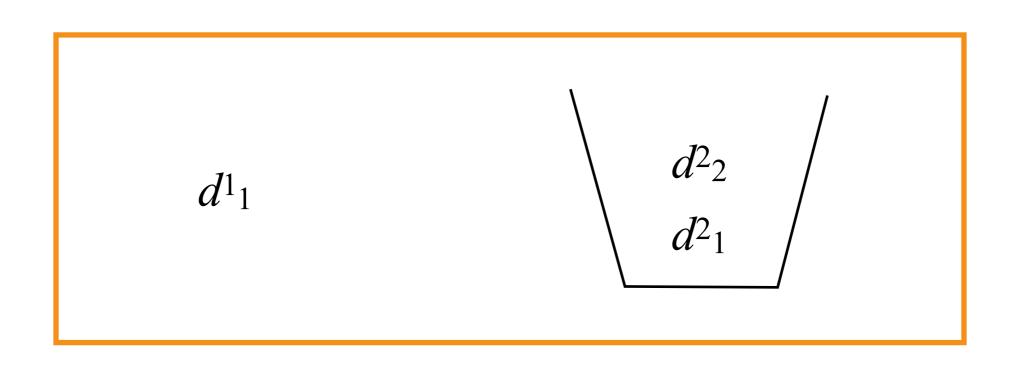
etc.

- Now, we have all descriptors of I_2 into buckets.
- To find a match for a descriptor d_i of I_1 , we apply f to d_i . In this way, we obtain a bucket number, let's call it r.
- We run the brute force method between d_i and all the descriptors that are in r.









We run brute-force: we compare d_1 with descriptor in the bucket.

- Advantages:
 - It is faster, we run the brute force method for a subset of descriptors.
- Disadvantages:
 - It is not exact, it is *approximate*; i.e., we test only a sub-set of descriptors.
 - If f is not well crafted, we may have distant descriptors in the same bucket.



Matching

- Once we have know matches between images, we can understand which images are near each others!
 - This is important for stable algorithms for triangulation of points and determining cameras' poses!

that's all folks!