

UNIVERSITÀ DEGLI STUDI DELLA BASILICATA







Corso di Sistemi Informativi A.A. 2018/19

Docente

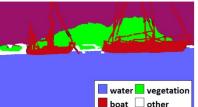
Domenico Daniele Bloisi



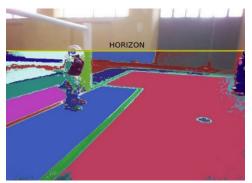














Riferimenti

Queste slide sono adattate da
 Noah Snavely - CS5670: Computer Vision
 "Lecture 5: Feature descriptors and matching"
 "Lecture 7: Transformations and warping"

 I contenuti fanno riferimento ai capitoli 3 e 4 del libro "Computer Vision: Algorithms and Applications" di Richard Szeliski, disponibile al seguente indirizzo http://szeliski.org/Book/

Problem: Feature matching



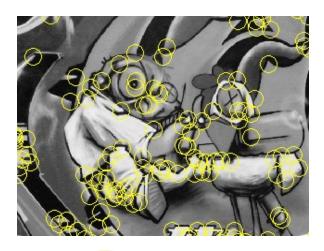
Recap

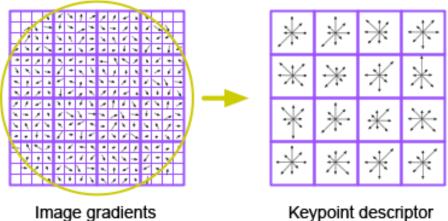
Keypoint detection: repeatable and distinctive

- Corners, blobs, stable regions
- Harris

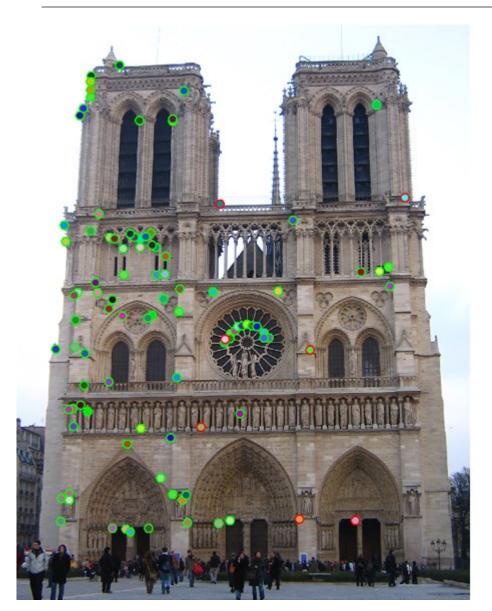
Descriptors: robust and selective

- spatial histograms of orientation
- SIFT and variants are typically good for stitching and recognition





Which features match?

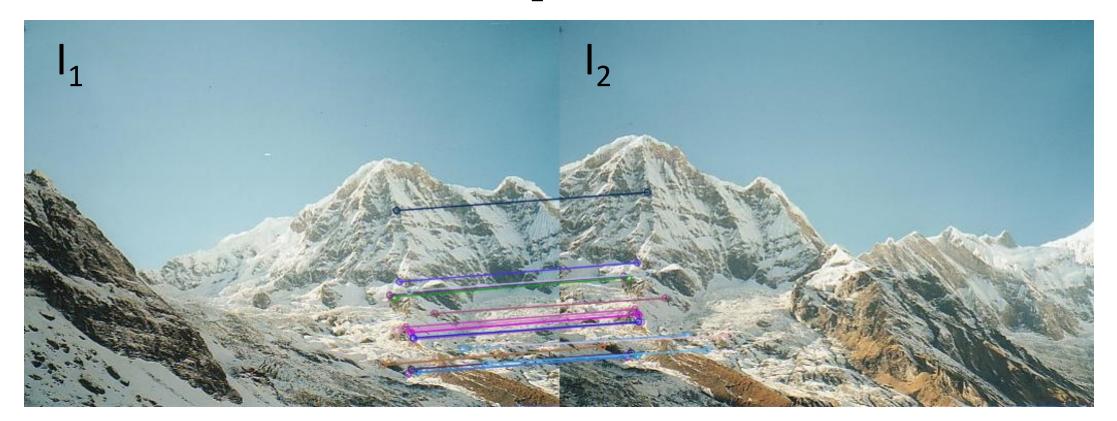




Features matching

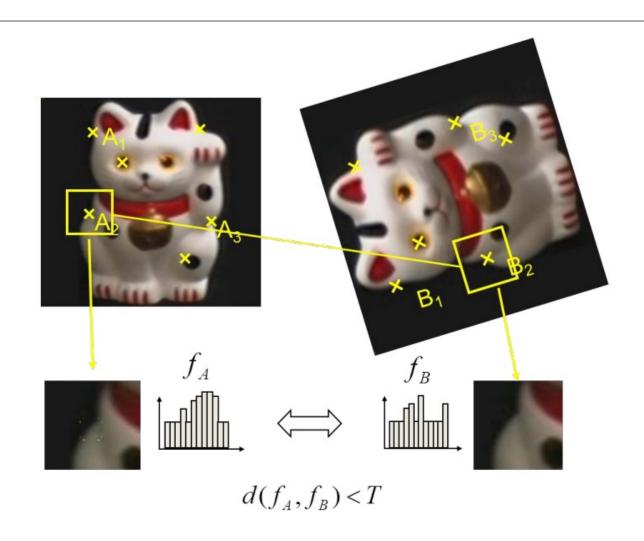
Given a feature in I_1 , how to find the best match in I_2 ?

- 1. Define distance function that compares two descriptors
- 2. Test all the features in I_2 , find the one with min distance



Overview of point feature matching

- 1. Detect a set of distinct feature points
- 2. Define a patch around each point
- 3. Extract and normalize the patch
- 4. Compute a local descriptor
- 5. Match local descriptors



Source: Trym Vegard Haavardsholm

Distance between descriptors

L₁ distance (SAD):

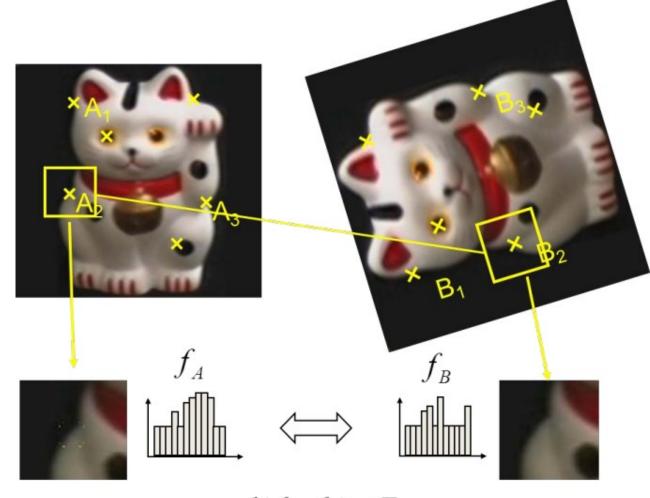
$$d(f_a, f_b) = \sum |f_a - f_b|$$

L₂ distance (SSD):

$$d(f_a, f_b) = \sum (f_a - f_b)^2$$

– Hamming distance:

$$d(f_a, f_b) = \sum XOR(f_a, f_b)$$



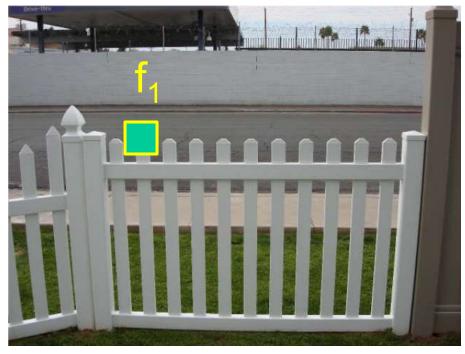
$$d(f_A, f_B) < T$$

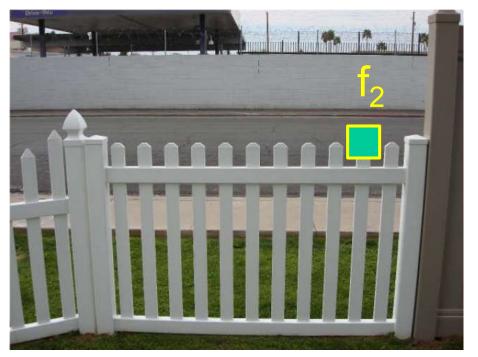
Source: Trym Vegard Haavardsholm

Features distance: SSD

How to define the difference between two features f_1 , f_2 ?

- Simple approach: L_2 distance, $||f_1 f_2||$ i.e., sum of square differences (SSD) between entries of the two descriptors
- can give small distances for ambiguous (incorrect) matches
 i.e., does not provide a way to discard ambiguous (bad) matches



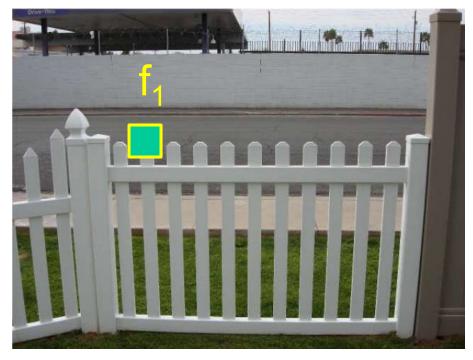


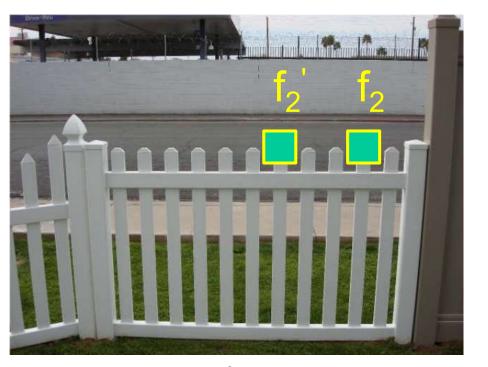
 l_1

Features distance: Ratio of SSDs

How to define the difference between two features f_1 , f_2 ?

- Better approach: ratio distance = SSD(f₁, f₂) / SSD(f₁, f₂')
 - $-f_2$ is best SSD match to f_1 in I_2
 - $-f_2$ ' is 2nd best SSD match to f_1 in I_2
 - An ambiguous/bad match will have ratio close to 1
 - Look for unique matches which have low ratio





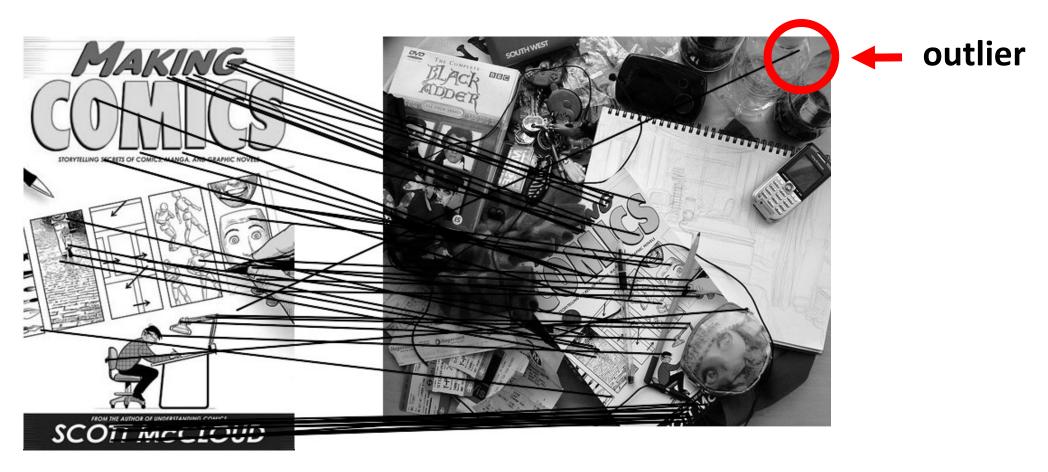
1.

Feature matching example



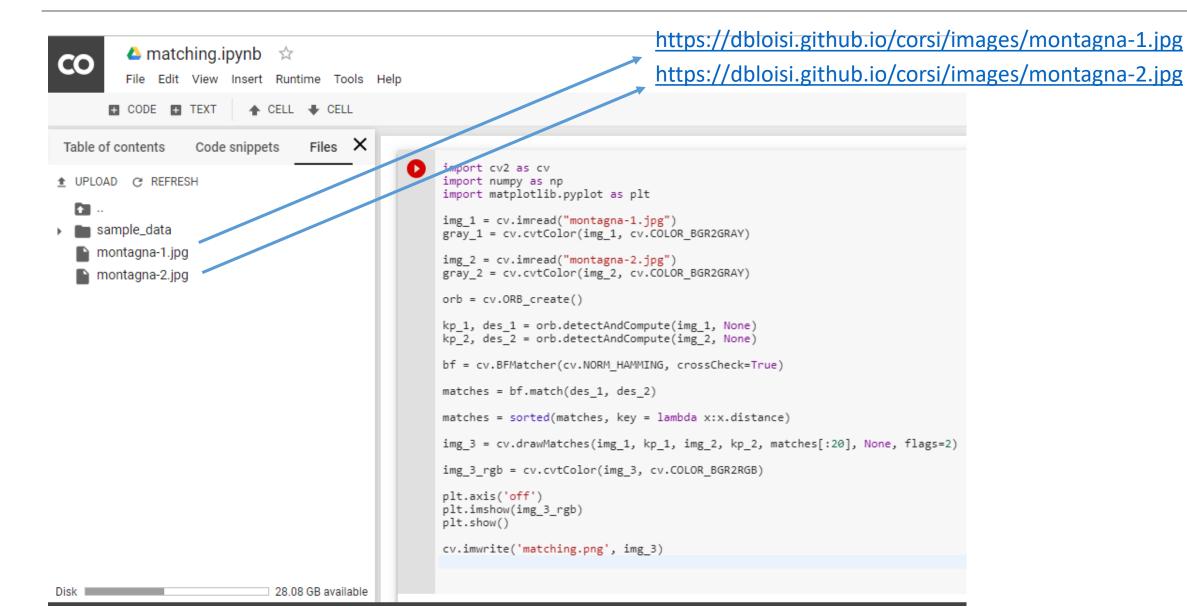
58 matches (thresholded by ratio score)

Feature matching example



51 matches (thresholded by ratio score)

Example in Colab



Example in Colab: ORB detection

```
import cv2 as cv
import numpy as np
import matplotlib.pyplot as plt
img 1 = cv.imread("montagna-1.jpg")
gray 1 = cv.cvtColor(img_1, cv.COLOR_BGR2GRAY)
img 2 = cv.imread("montagna-2.jpg")
gray_2 = cv.cvtColor(img_2, cv.COLOR_BGR2GRAY)
orb = cv.ORB_create()
kp_1, des_1 = orb.detectAndCompute(img_1, None)
kp 2, des 2 = orb.detectAndCompute(img 2, None)
```

Example in Colab: brute force matching

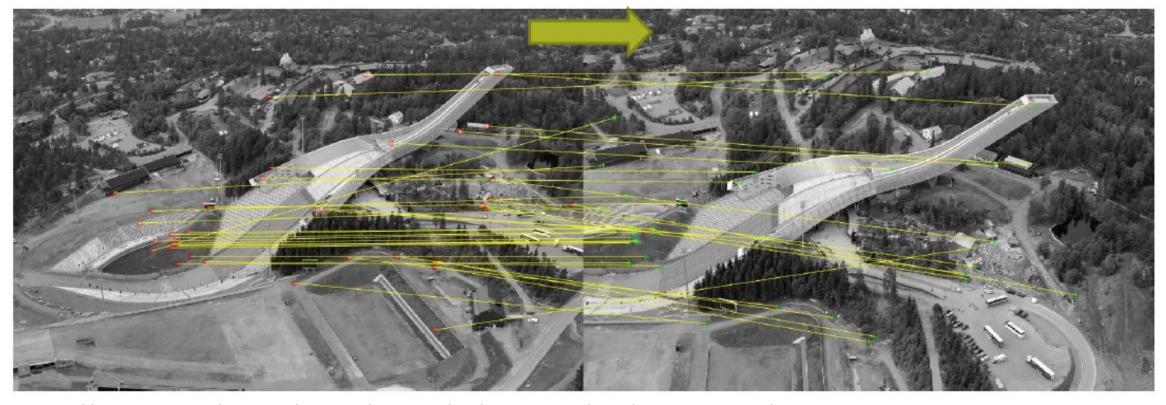
Brute-Force matcher is simple. It takes the descriptor of one feature in first set and is matched with all other features in second set using some distance calculation. And the closest one is returned.

```
bf = cv.BFMatcher(cv.NORM_HAMMING, crossCheck=True)
matches = bf.match(des_1, des_2)
matches = sorted(matches, key = lambda x:x.distance)
```

Cross check test

- Choose matches (f_a, f_b) so that
 - f_b is the best match for f_a in I_b
 - And f_a is the best match for f_b in I_a

Alternative to ratio test

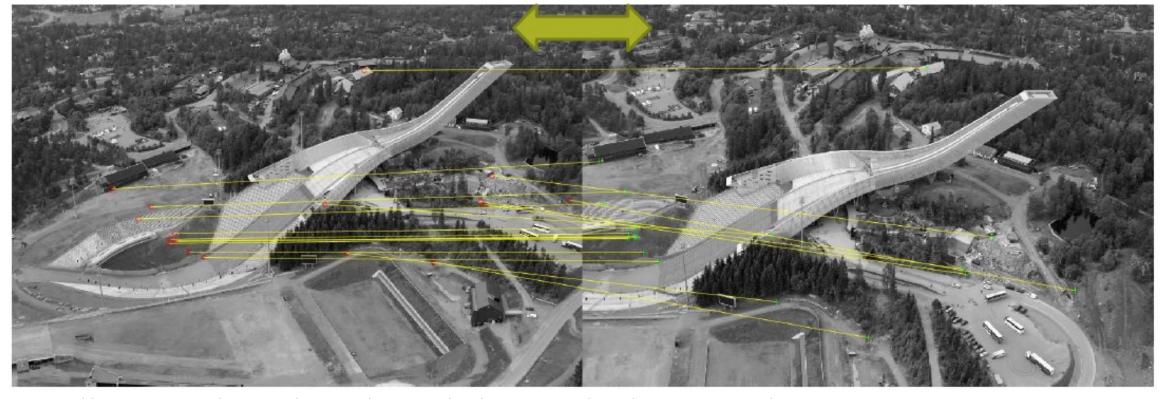


https://www.uio.no/studier/emner/matnat/its/UNIK4690/v17/forelesninger/lecture 4 2 feature matching.pdf

Cross check test

- Choose matches (f_a, f_b) so that
 - f_b is the best match for f_a in I_b
 - And f_a is the best match for f_b in I_a

Alternative to ratio test



https://www.uio.no/studier/emner/matnat/its/UNIK4690/v17/forelesninger/lecture 4 2 feature matching.pdf

Example in Colab: sorting

```
bf = cv.BFMatcher(cv.NORM_HAMMING, crossCheck=True)
matches = bf.match(des_1, des_2)
matches = sorted(matches, key = lambda x:x.distance)
```

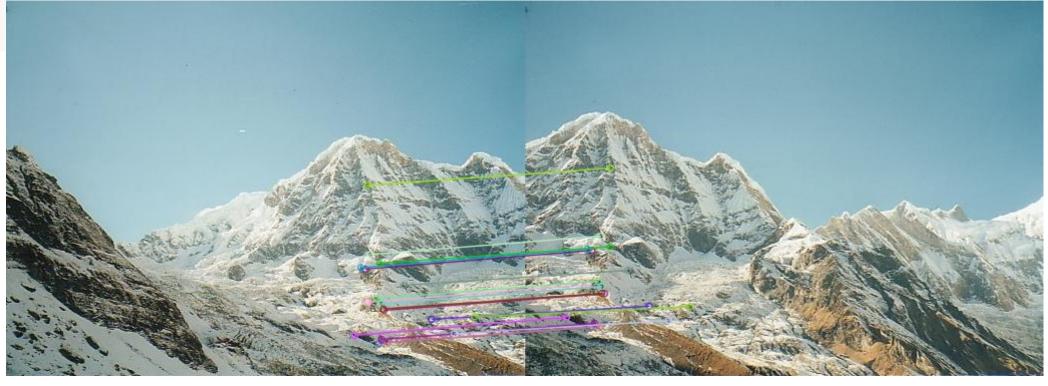
Matches are sorted in ascending order of their distances so that best matches (with low distance) come to front.

Example in Colab: result

```
img_3 = cv.drawMatches(img_1, kp_1, img_2, kp_2, matches[:20], None, flags=2)
img_3_rgb = cv.cvtColor(img_3, cv.COLOR_BGR2RGB)

plt.axis('off')
plt.imshow(img_3_rgb)
plt.show()

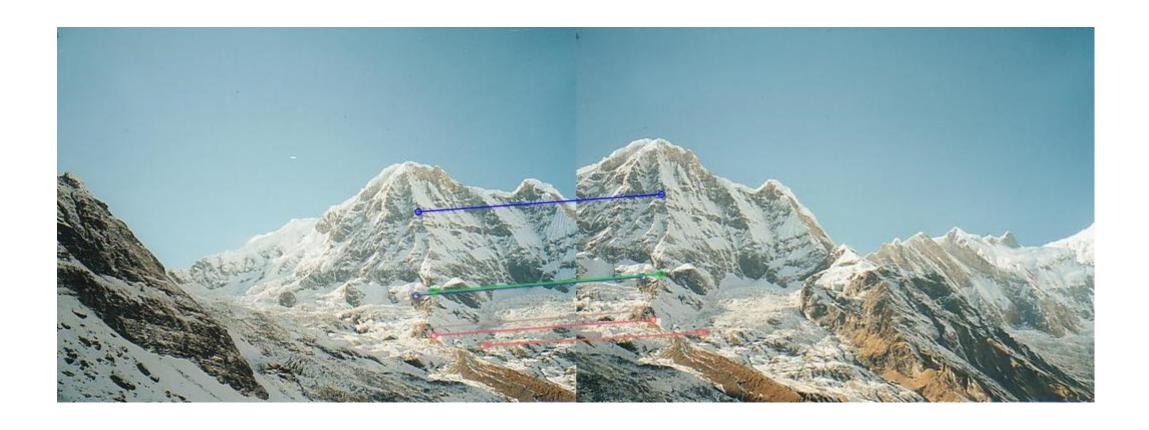
cv.imwrite('matching.png', img_3)
```



Example in Colab: ratio test

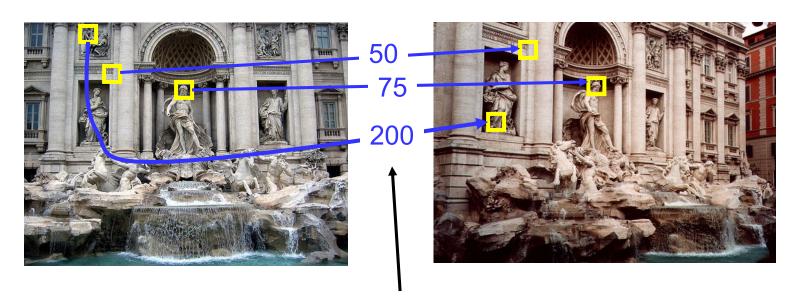
```
orb = cv.ORB_create()
kp 1, des 1 = orb.detectAndCompute(img 1, None)
kp 2, des 2 = orb.detectAndCompute(img 2, None)
bf = cv.BFMatcher()
                                           BFMatcher.knnMatch() to get k best matches.
matches = bf.knnMatch(des_1, des_2, k=2)
                                           In this example, we will take k=2 so that we
# Apply ratio test
                                           can apply ratio test
good = []
for m,n in matches:
   if m.distance < 0.7*n.distance:
        good.append([m])
img 3 = cv.drawMatchesKnn(img 1, kp 1, img 2, kp 2, good[:20], None, flags=2)
img 3 rgb = cv.cvtColor(img 3, cv.COLOR BGR2RGB)
plt.axis('off')
plt.imshow(img 3 rgb)
plt.show()
cv.imwrite('matching.png', img 3)
```

Example in Colab: ratio test results



Evaluating the results

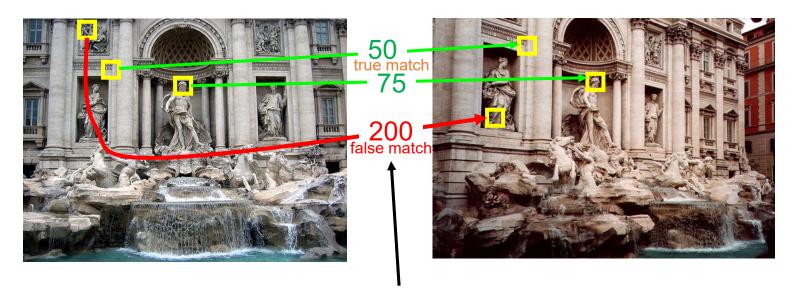
How can we measure the performance of a feature matcher?



feature distance

True/false positives

How can we measure the performance of a feature matcher?

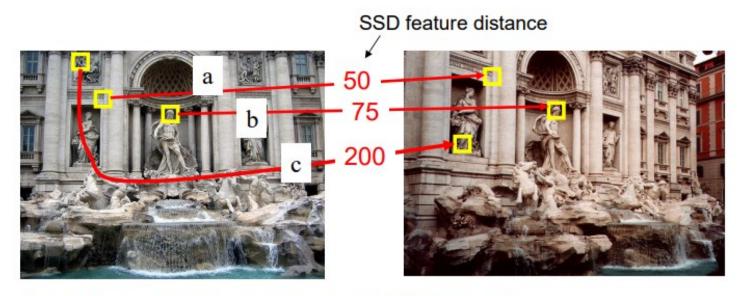


feature distance

The distance threshold affects performance

- True positives = # of detected matches that are correct
 - Suppose we want to maximize these—how to choose threshold?
- False positives = # of detected matches that are incorrect
 - Suppose we want to minimize these—how to choose threshold?

Effect of threshold T



Decision rule: Accept match if SSD < T

Example: Large T

 $T = 250 \Rightarrow a, b, c$ are all accepted as matches

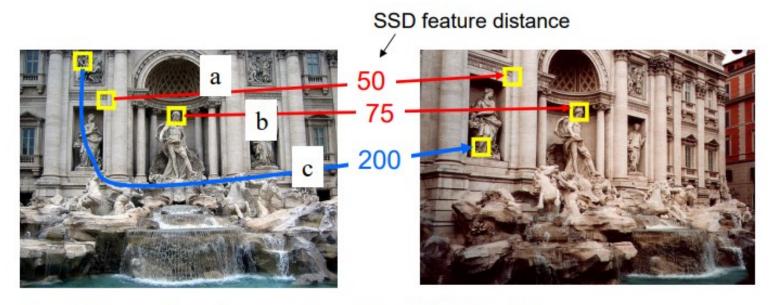
a and b are true matches ("true positives")

they are actually matches

c is a false match ("false positive")

- actually not a match

Effect of threshold T



Decision rule: Accept match if SSD < T

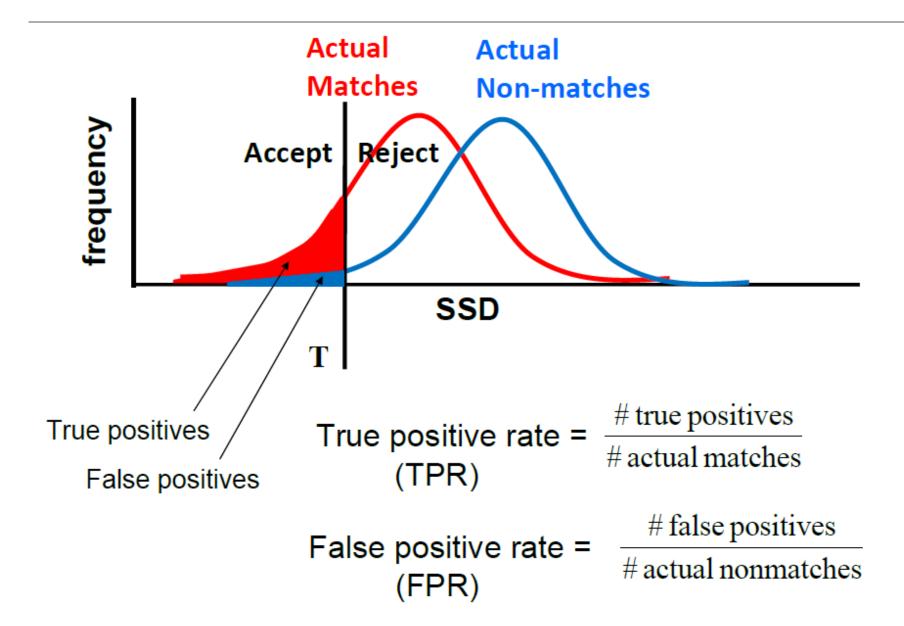
Example: Smaller T

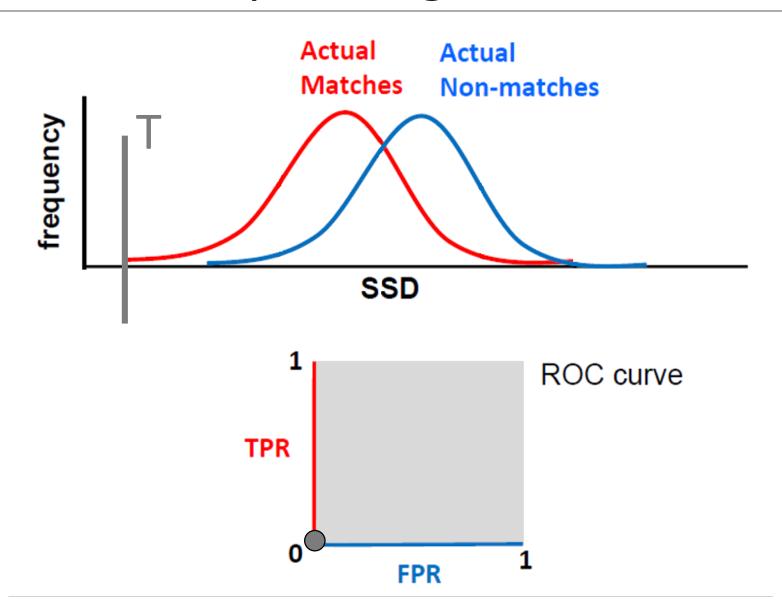
 $T = 100 \Rightarrow$ only a and b are accepted as matches

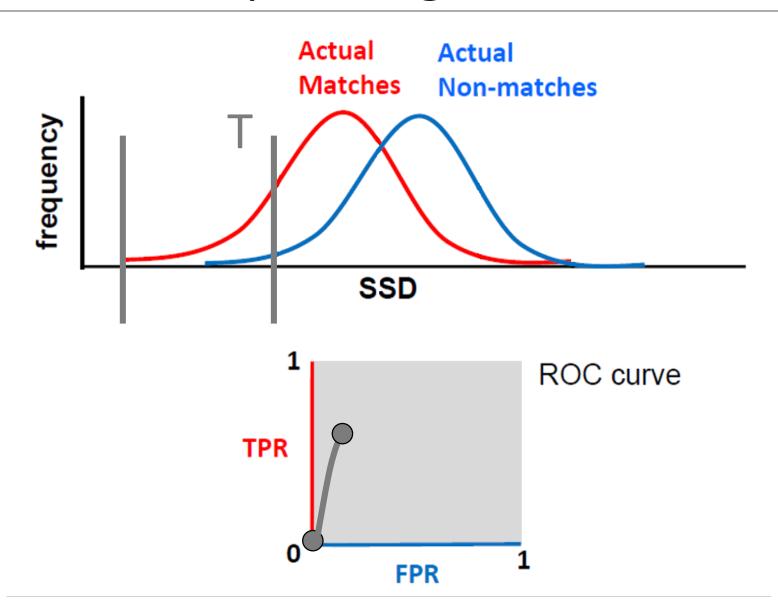
a and b are true matches ("true positives")

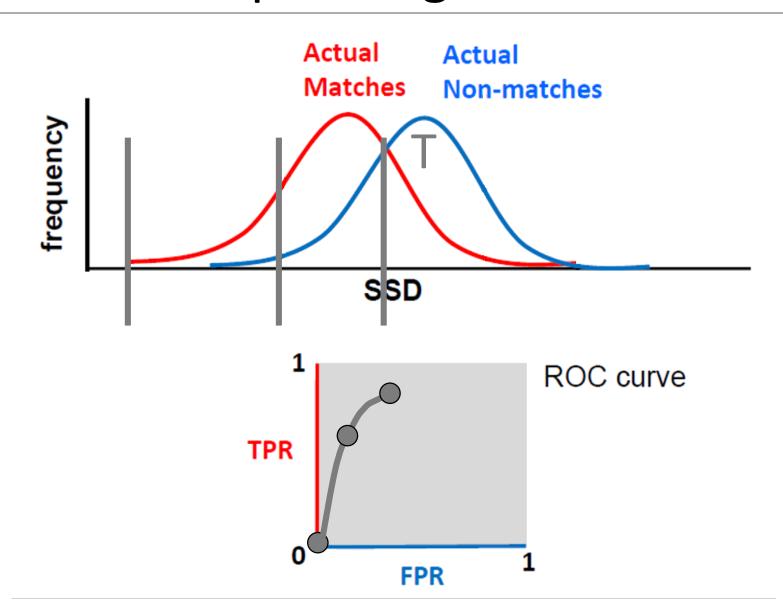
c is no longer a "false positive" (it is a "true negative")

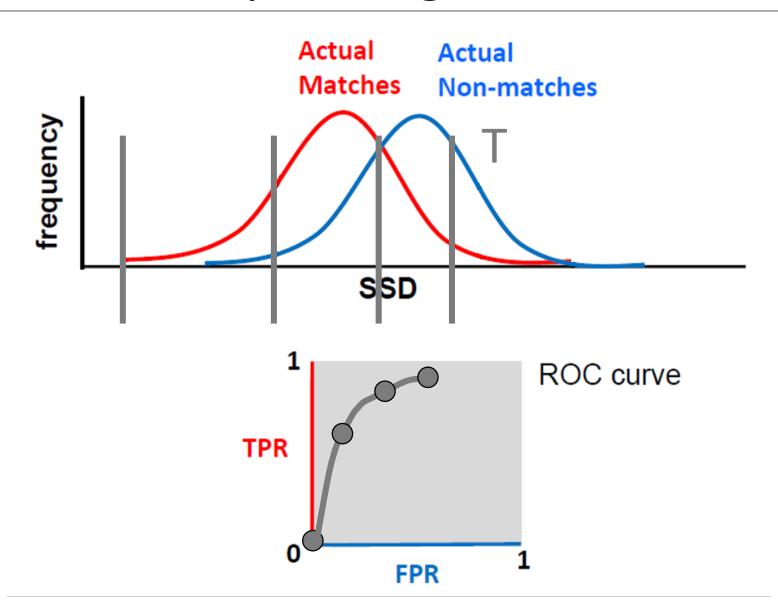
True positives and false positives

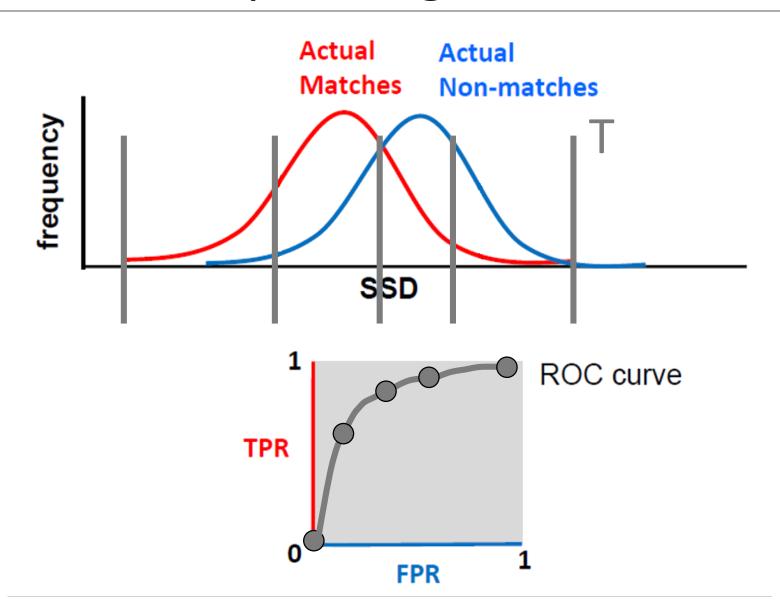




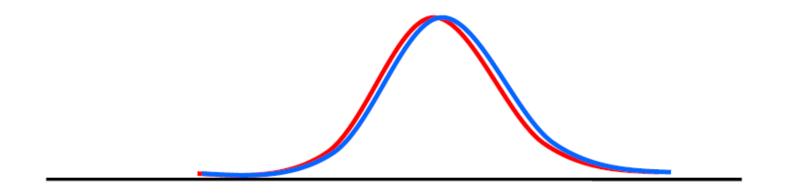


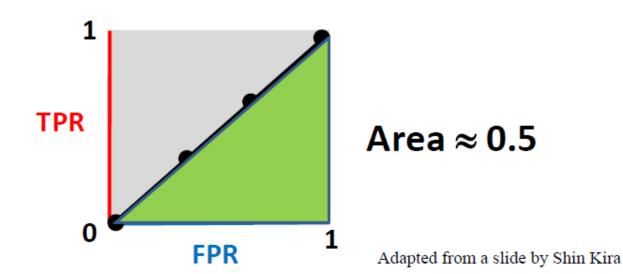




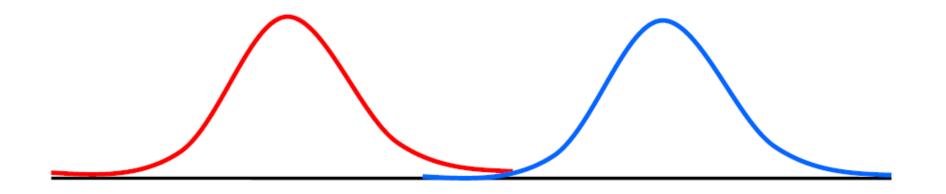


If the features selected were bad...





If the features selected were good...

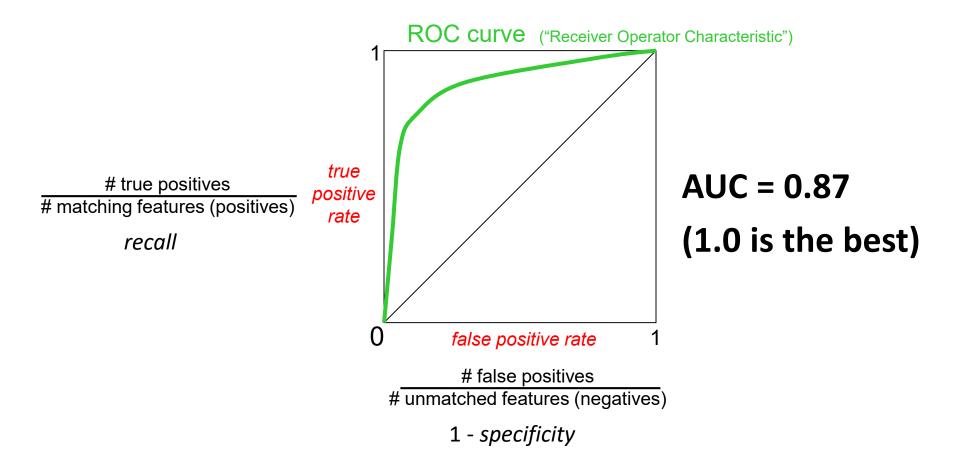




Adapted from a slide by Shin Kira

Area under the curve

Single number: Area Under the Curve (AUC)





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Feature Matching



