

Advanced Computer Vision

Week 07

Oct. 14, 2022
Seokju Lee

Let's Learn SIFT Algorithm

Scale Invariant Feature Transform (SIFT)

and its applications for image alignment and 2D object recognition.

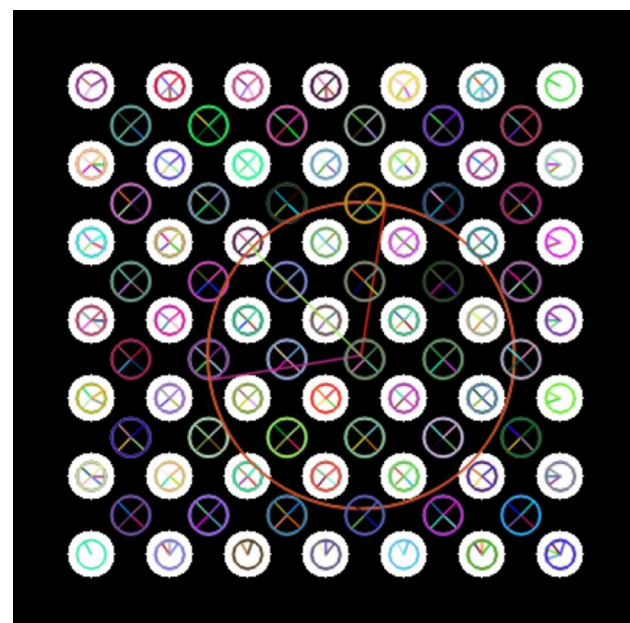
Topics:

(1) What is an Interest Point?

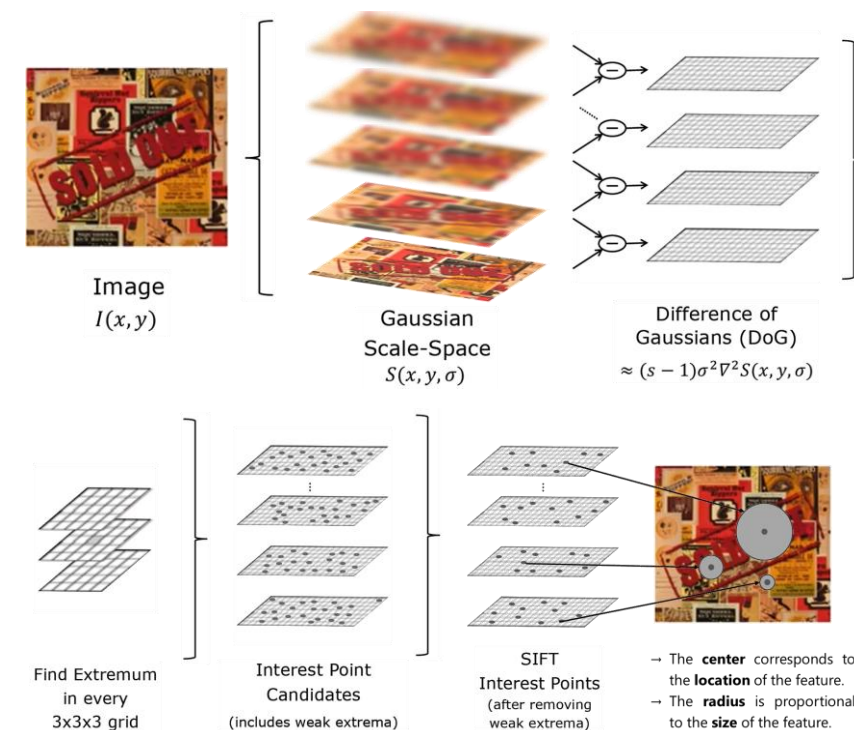
(2) Detecting Blobs

(3) SIFT Detector

(4) SIFT Descriptor



Detected SIFT features with multiscale





SIFT Descriptor



What is Descriptor?

Descriptor = Set of **Distinctive** features and their **identifiers**

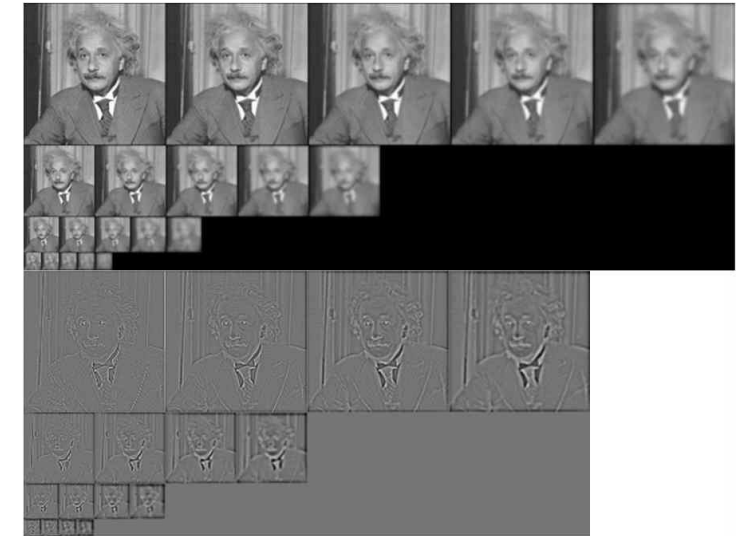
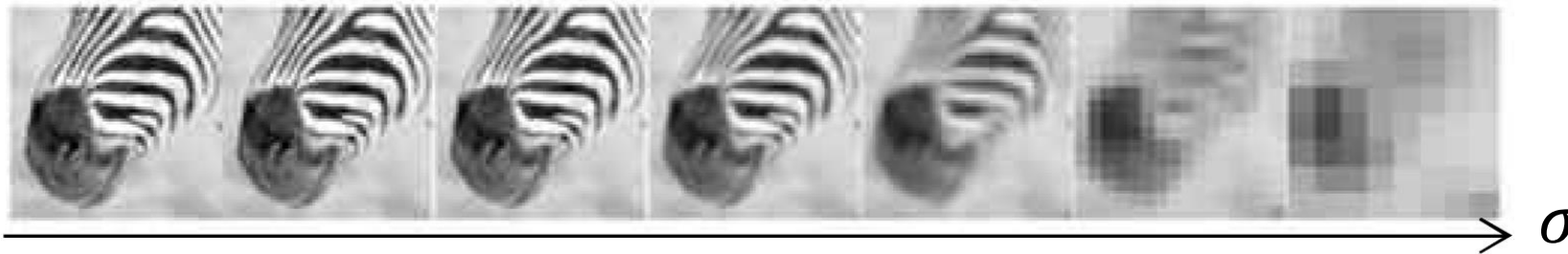


→ Must be **robust** to variances + **generalized/standardized**

SIFT: Scale Invariance

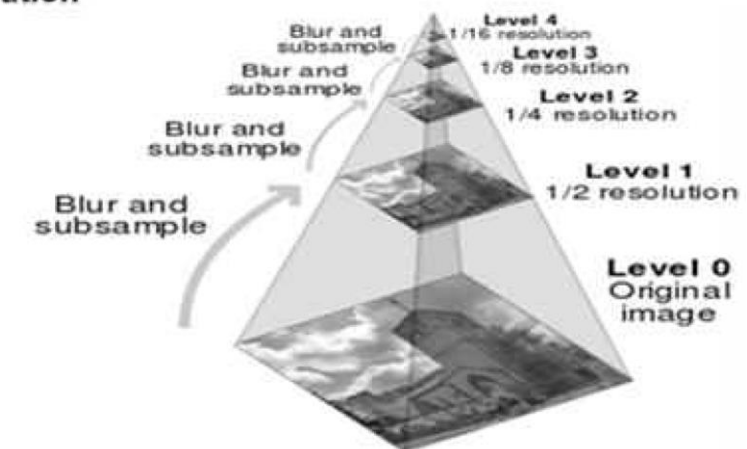
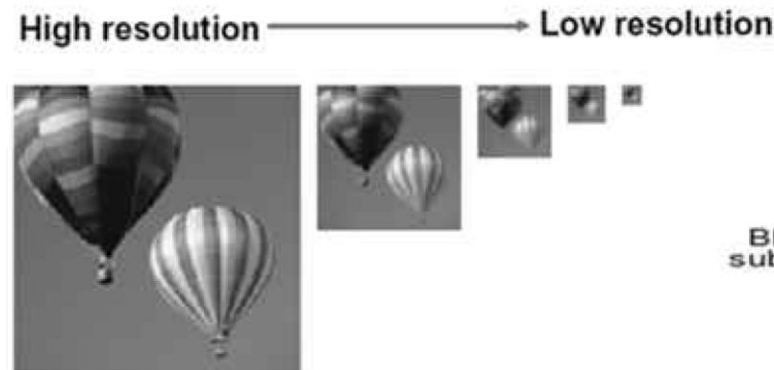
Scale space

Different scales are necessary for describing different objects in the image, and we may not know the correct scale/size ahead of time (generalization).



Gaussian Pyramid

Image pyramid technique is used for reducing the memory usage and computational complexity.



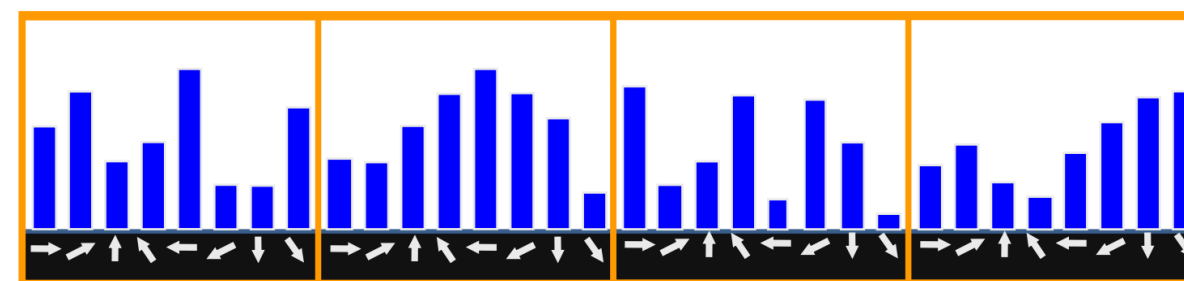
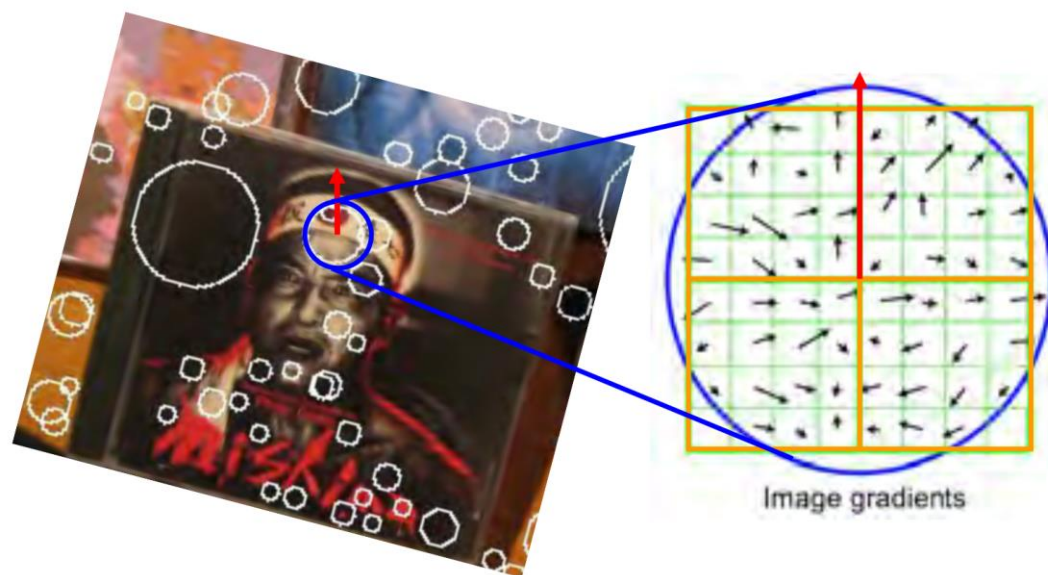
SIFT: Rotation Invariance

Compute gradient for each blurred image

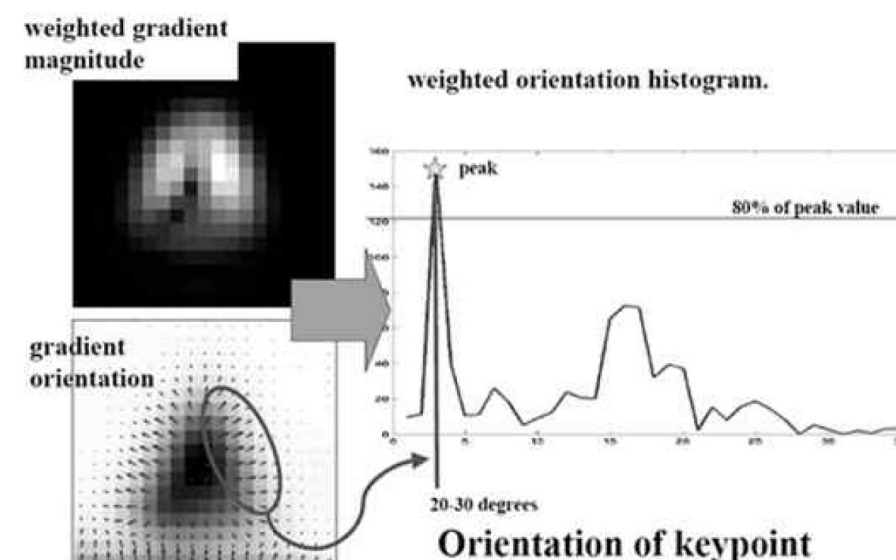
$$m(x, y) = \sqrt{(L(x+1, y) - L(x-1, y))^2 + (L(x, y+1) - L(x, y-1))^2}$$

$$\theta(x, y) = \tan^{-1}((L(x, y+1) - L(x, y-1)) / (L(x+1, y) - L(x-1, y)))$$

Histograms of gradient directions over spatial regions



- **Normalized** histogram: invariant to **rotation**, **scale**, **brightness**
- Can be directly used as a **signature** for matching SIFT Features!



Comparing SIFT Descriptors

Essentially comparing two arrays of data

Let $H_1(k)$ and $H_2(k)$ be two arrays of data of length N ,

L2 Distance :

$$d(H_1, H_2) = \sqrt{\sum_k (H_1(k) - H_2(k))^2}$$

Smaller the distance metric, **better** the match.

Perfect match when $d(H_1, H_2) = 0$

Comparing SIFT Descriptors

Essentially comparing two arrays of data

Let $H_1(k)$ and $H_2(k)$ be two arrays of data of length N ,

Normalized Correlation :

$$d(H_1, H_2) = \frac{\sum_k [(H_1(k) - \bar{H}_1)(H_2(k) - \bar{H}_2)]}{\sqrt{\sum_k (H_1(k) - \bar{H}_1)^2} \sqrt{\sum_k (H_2(k) - \bar{H}_2)^2}}$$

$$\text{where: } \bar{H}_i = \frac{1}{N} \sum_{k=1}^N H_i(k)$$

Larger the distance metric, **better** the match.

Perfect match when $d(H_1, H_2) = 1$

*Correlation?



Source image



Template

How do we **locate** the template in the image?

Minimize :

$$E[i, j] = \sum_m \sum_n (f[i + m, j + n] - t[m, n])^2$$

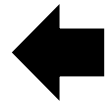
$$E[i, j] = \sum_m \sum_n (f^2[i + m, j + n] + t^2[m, n] - \underline{2f[i + m, j + n]t[m, n]})$$

→ Maximize

*Correlation?



Source image



Template

How do we **locate** the template in the image?

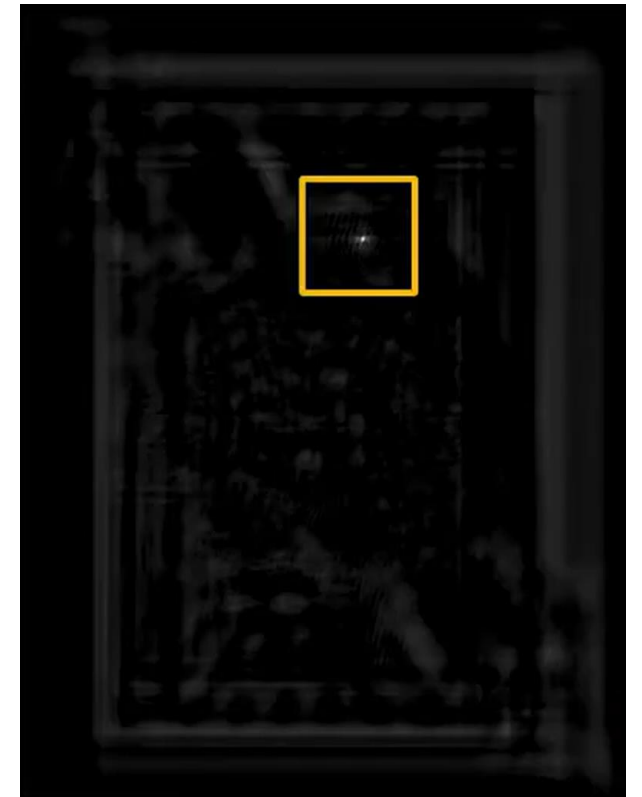
Maximize :

$$R[i, j] = \sum_m \sum_n t[m, n] f[i + m, j + n] = t \otimes f$$

→ Cross-correlation

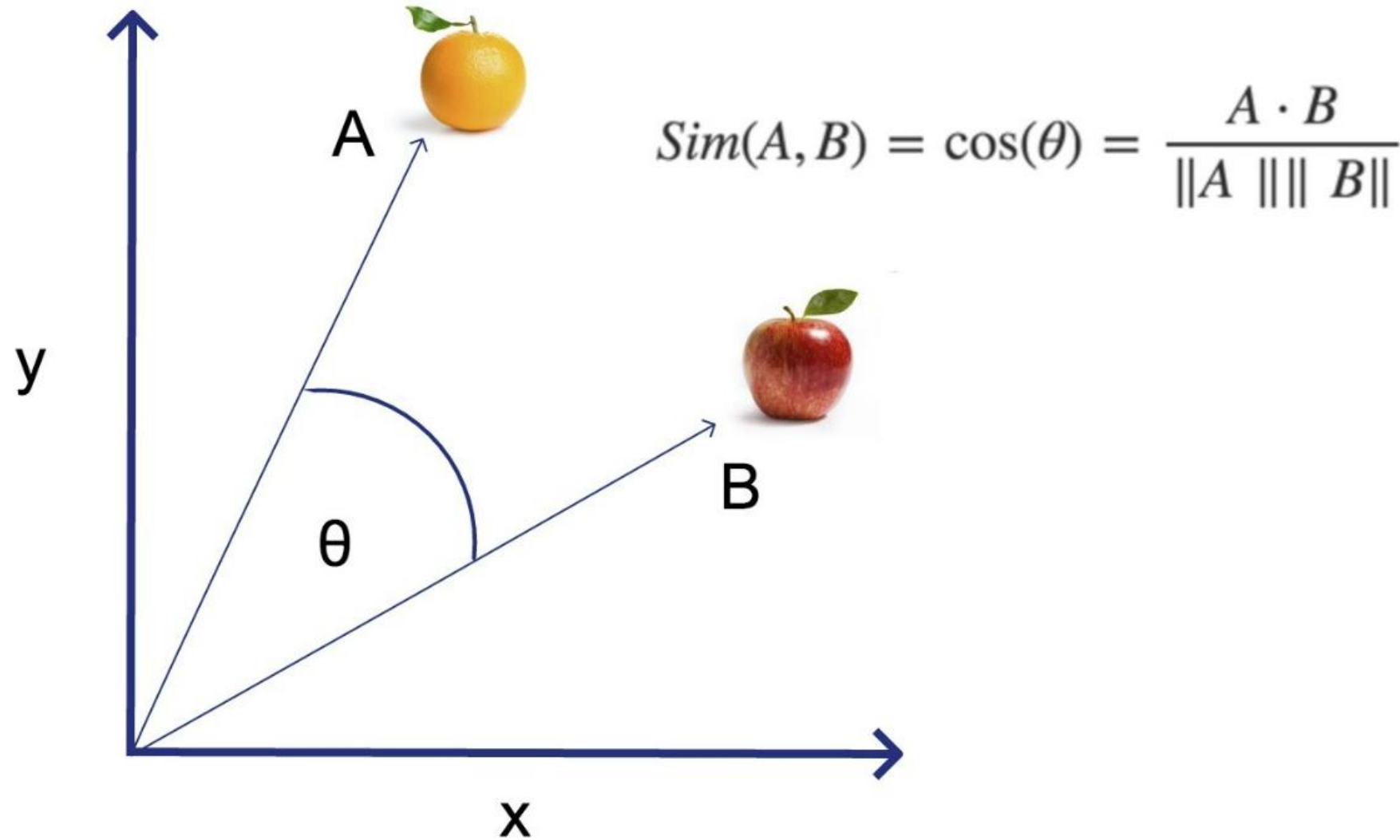
c.f.) convolution

$$\text{cosine similarity} = S_C(A, B) := \cos(\theta) = \frac{\mathbf{A} \cdot \mathbf{B}}{\|\mathbf{A}\| \|\mathbf{B}\|} = \frac{\sum_{i=1}^n A_i B_i}{\sqrt{\sum_{i=1}^n A_i^2} \sqrt{\sum_{i=1}^n B_i^2}},$$



→ Result of cross-correlation

*Cosine Similarity



Comparing SIFT Descriptors

Essentially comparing two arrays of data

Let $H_1(k)$ and $H_2(k)$ be two arrays of data of length N ,

Normalized Correlation :

$$d(H_1, H_2) = \frac{\sum_k [(H_1(k) - \bar{H}_1)(H_2(k) - \bar{H}_2)]}{\sqrt{\sum_k (H_1(k) - \bar{H}_1)^2} \sqrt{\sum_k (H_2(k) - \bar{H}_2)^2}}$$

$$\text{where: } \bar{H}_i = \frac{1}{N} \sum_{k=1}^N H_i(k)$$

Larger the distance metric, **better** the match.

Perfect match when $d(H_1, H_2) = 1$

Comparing SIFT Descriptors

Essentially comparing two arrays of data

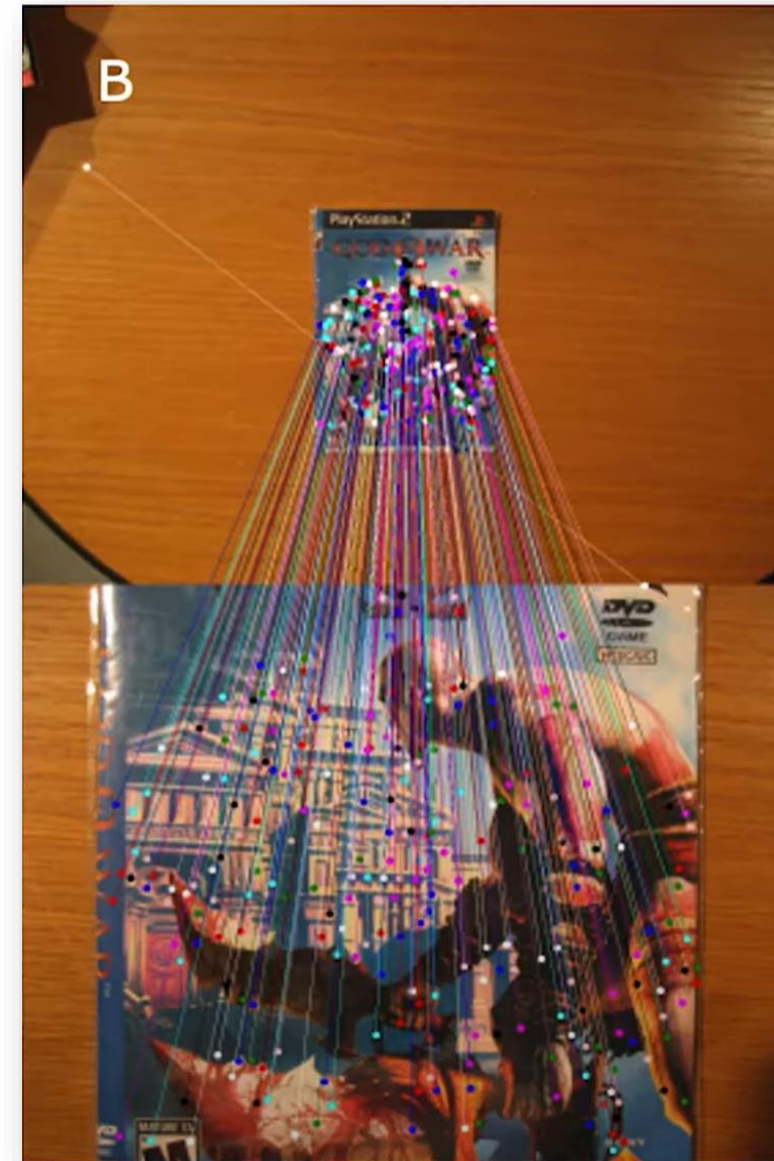
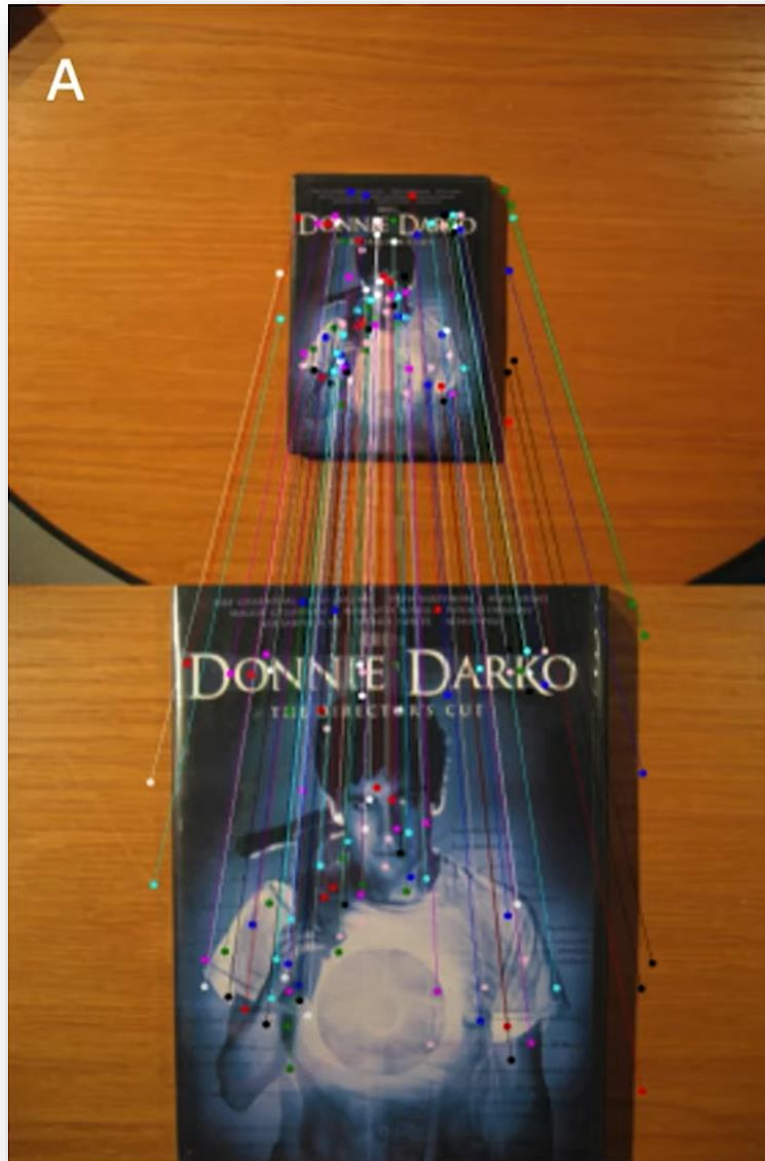
Let $H_1(k)$ and $H_2(k)$ be two arrays of data of length N ,

Intersection :

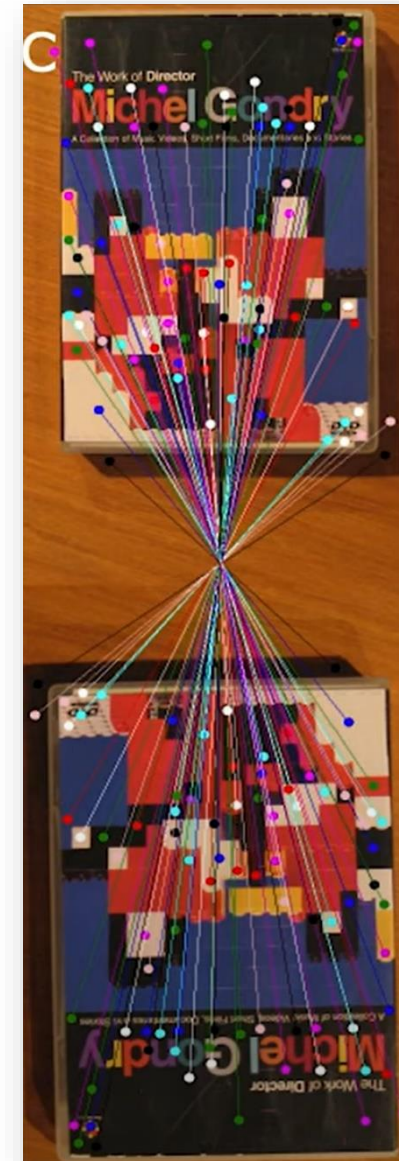
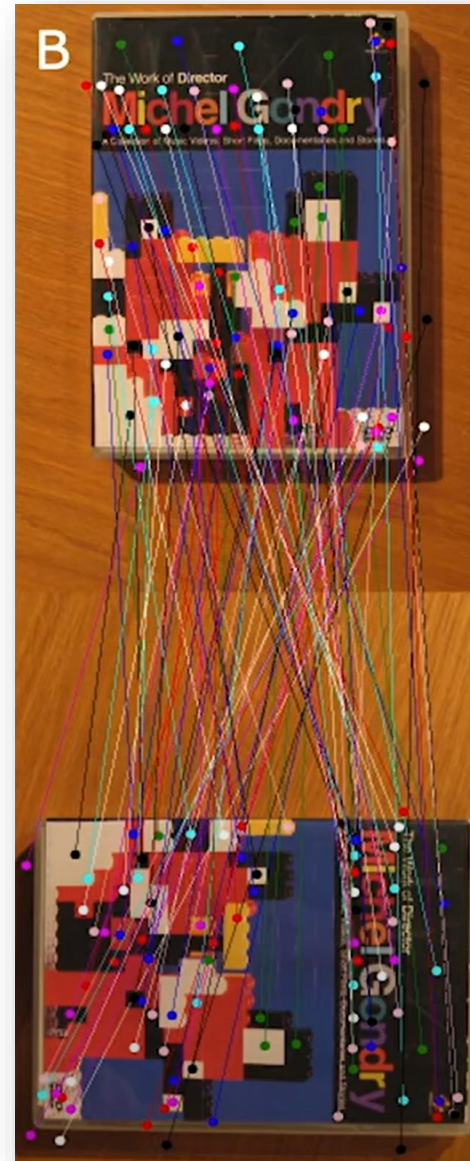
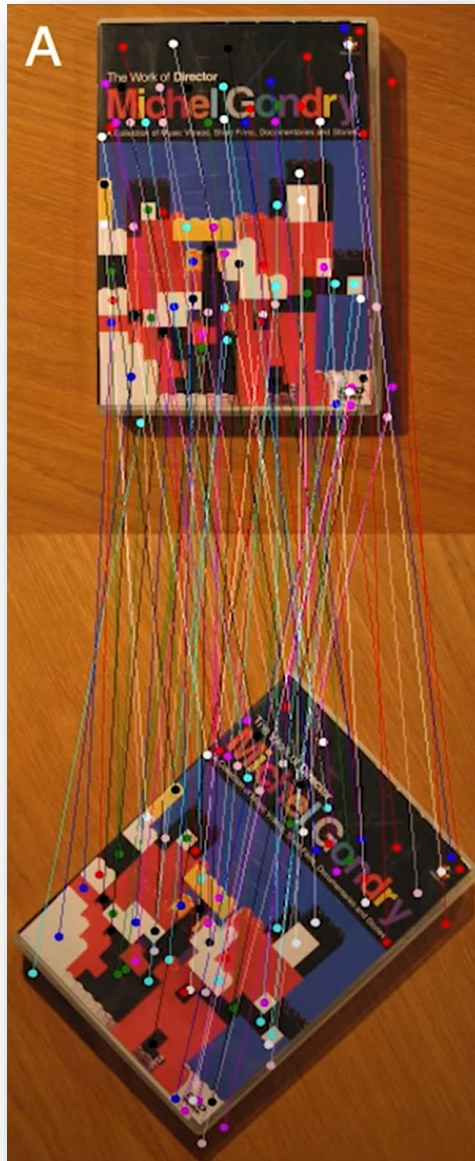
$$d(H_1, H_2) = \sum_k \min(H_1(k), H_2(k))$$

Larger the distance metric, **better** the match.

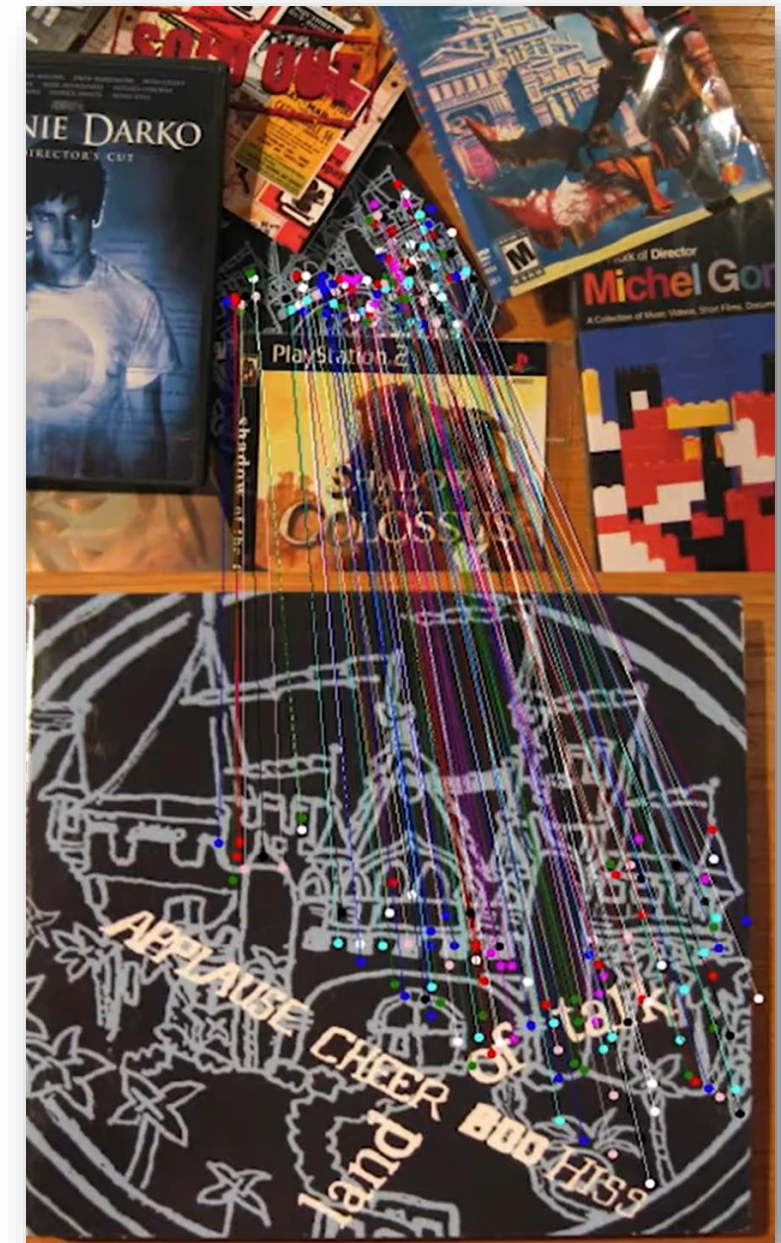
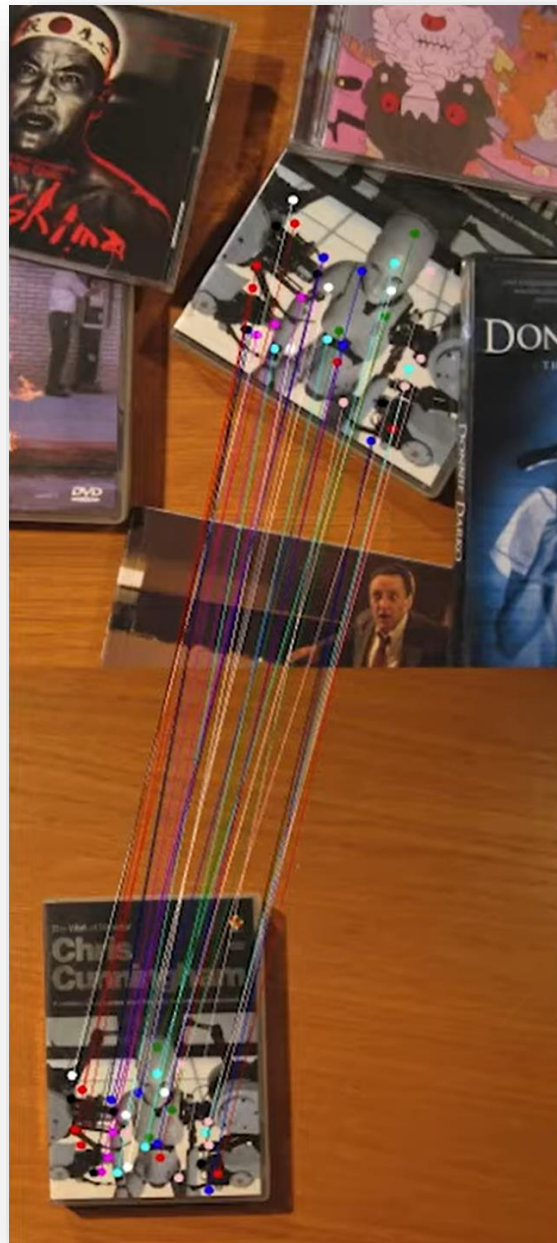
SIFT Results: Scale Invariance



SIFT Results: Rotation Invariance



SIFT Results: Robust to Clutter

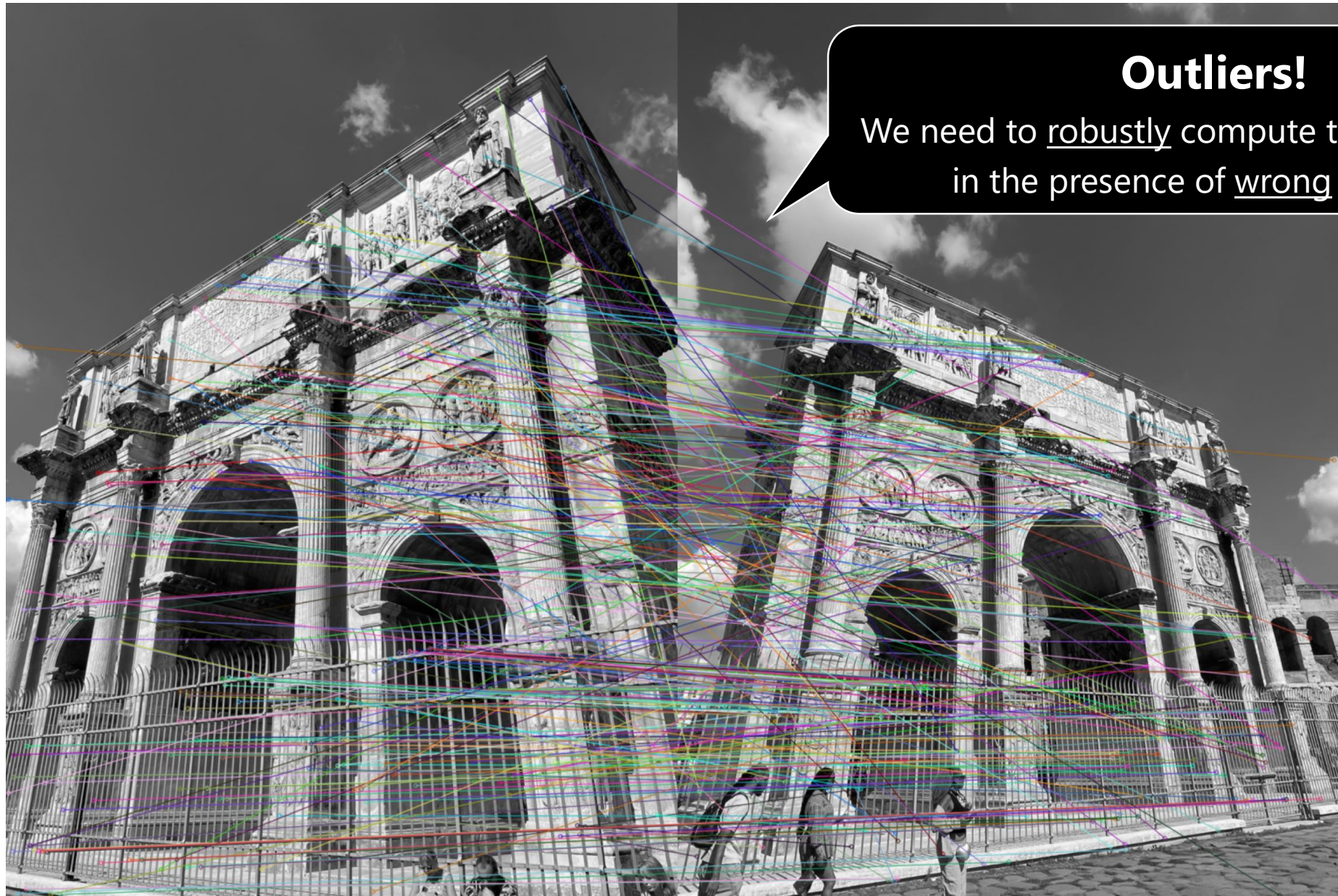


Experiment 1: SIFT Matching with Different Motions and Image Transformations

<https://view.kentech.ac.kr/f088fa7f-874e-44bc-bd6d-6084b42dfdf7>



How to Handle Outliers?



Outliers!

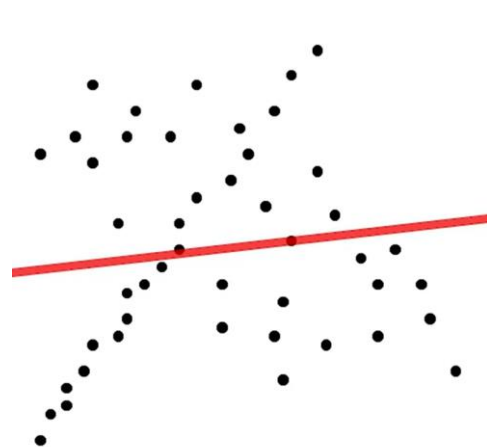
We need to robustly compute transformations in the presence of wrong matches.

RANdom SAmple Consensus

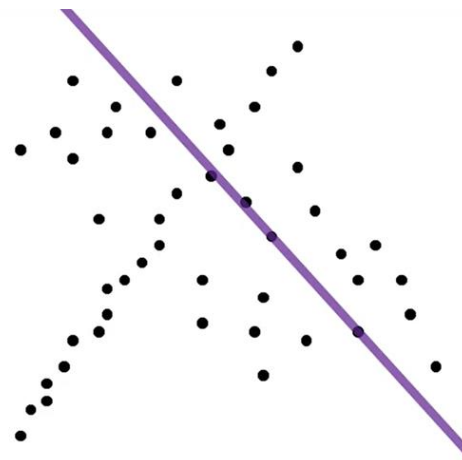
General RANSAC Algorithm :

1. Randomly choose s samples. Typically s is the minimum samples to fit a model.
2. Fit the model to the randomly chosen samples.
3. Count the number M of data points (inliers) that fit the model within a measure of error ϵ .
4. Repeat Steps 1-3 N times.
5. Choose the model that has the largest number M of inliers.

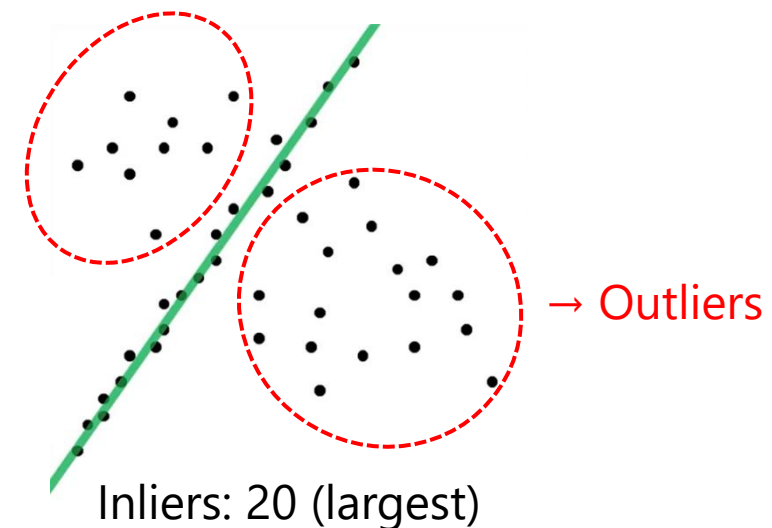
ex) Line fitting :



Inliers: 2



Inliers: 4



Inliers: 20 (largest)

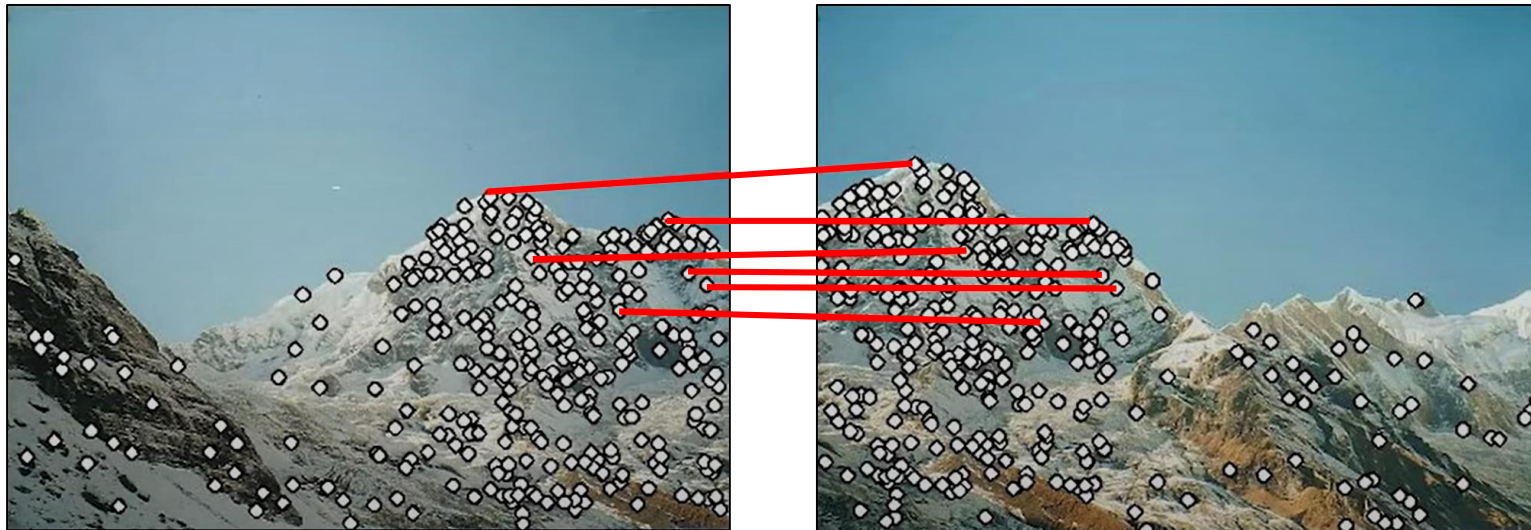
Experiment 2: Panorama Stitching using SIFT



Image 1



Image 2



Match SIFT interest points.

Experiment 2: Panorama Stitching using SIFT

Warp and combine images to create a larger image.

