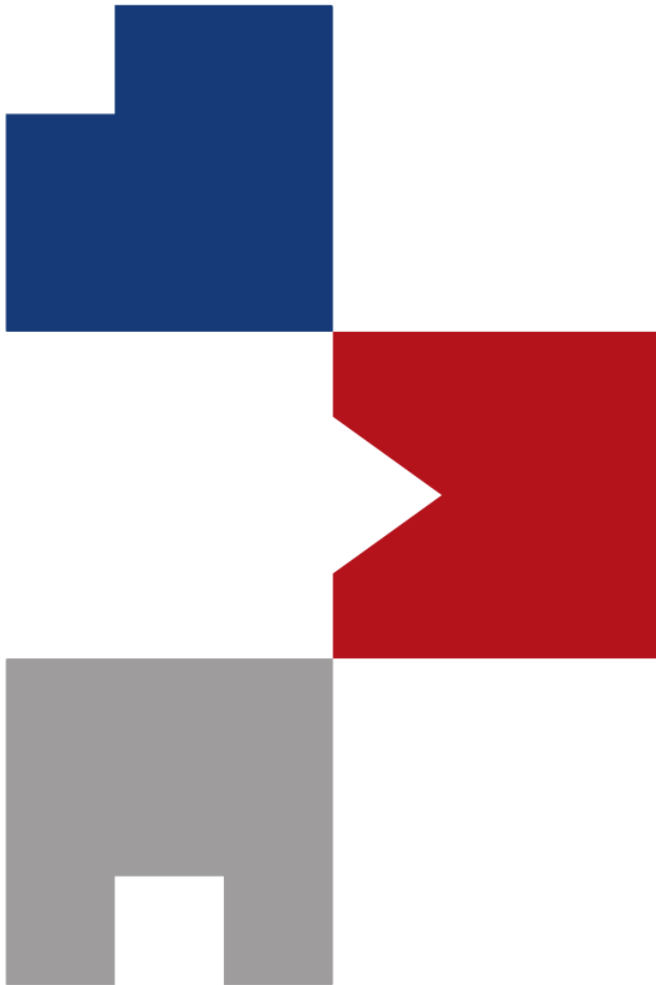


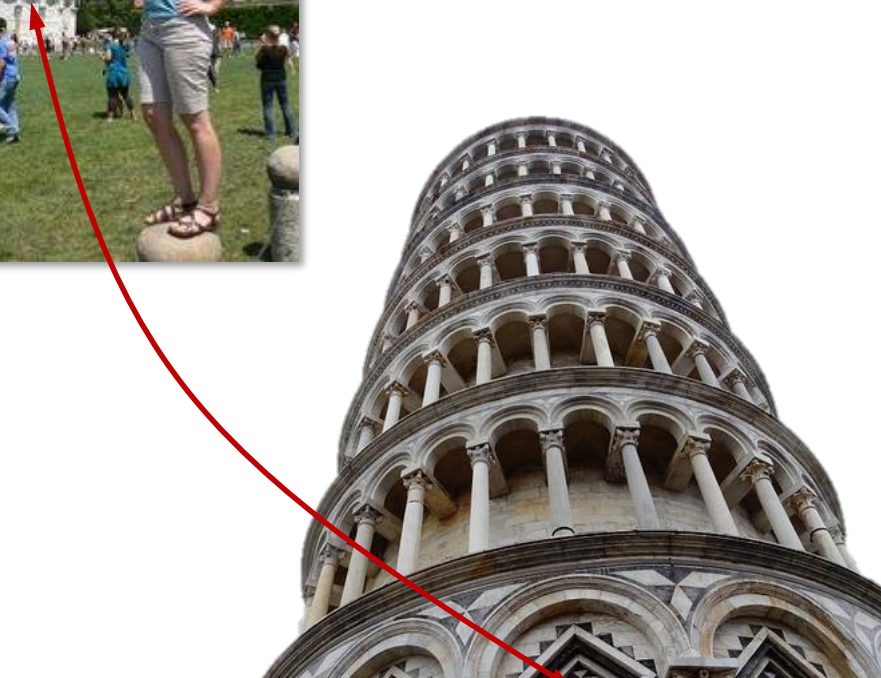
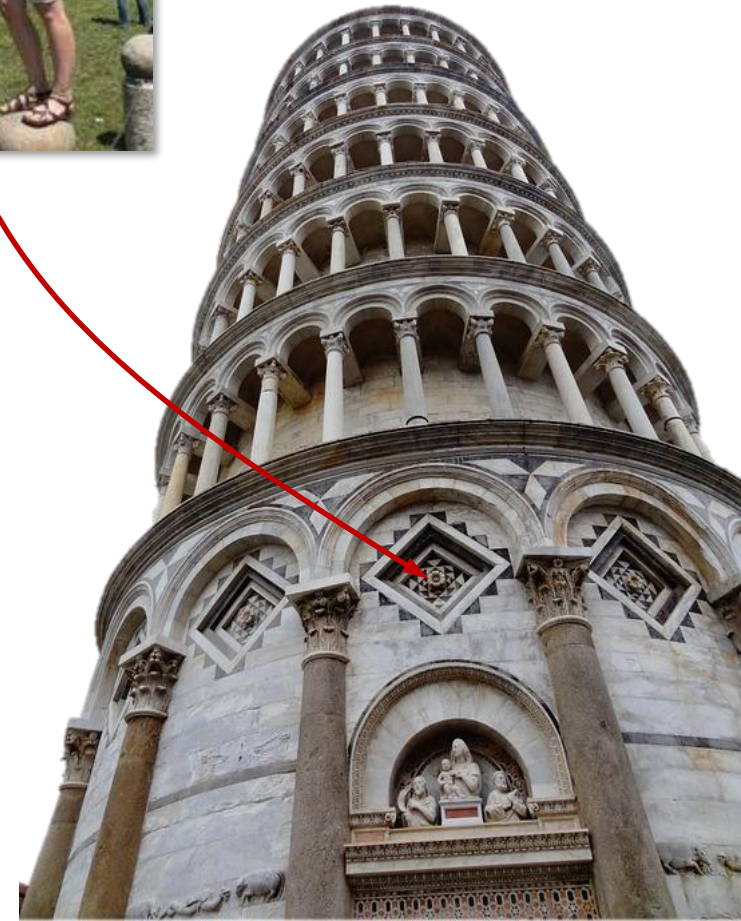
An Invitation to 3D Vision: Finding Correspondences



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Table of Contents: Finding Correspondences

- **Feature Points**
 - Q) How can we detect salient points (or parts)?
- **Feature Descriptors**
 - Q) How can we distinguish the points each other?
- **Feature Matching**
 - Q) How can we associate the points across different images?
- **Feature Tracking**
 - Q) How can we associate the points across their next image?
- **Outlier Rejection**
 - Q) How can we select correctly matched points?



Getting Started with a Quiz

- Q) What is it?



600 pixels

Getting Started with a Quiz

- Q) What is it?



Getting Started with a Quiz

- Q) What is it? **"Duck"**



600 pixels



1095 pixels



600 pixels

- Q) Why corners (junction) instead of edges or blobs?
 - Human visual systems understand objects better from corners than edges.
 - a.k.a. *feature* points, *keypoints*, *salient* points, and *interest* points

Getting Started with a Quiz

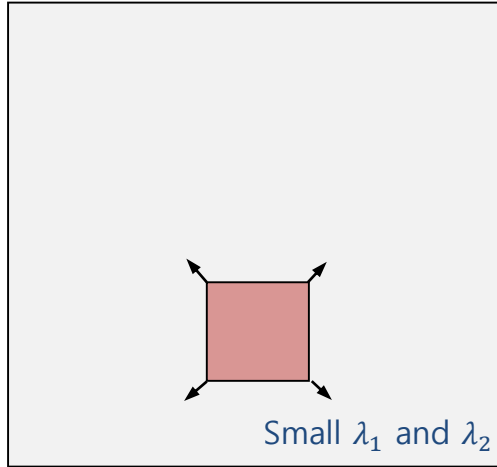
- Q) Why corners (junction) instead of edges or blobs? Where are the image patches?



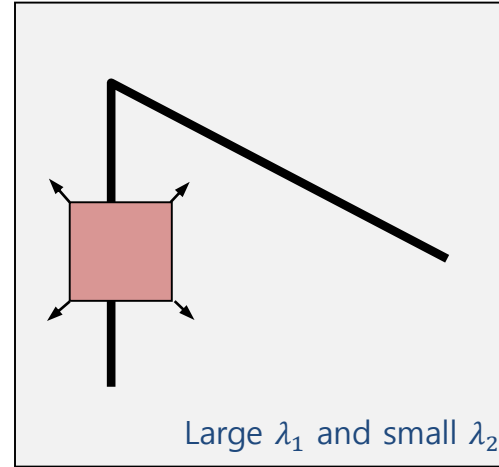
- Q) What is a good feature? (Requirements)
 - **Repeatability** (to be invariance/robustness to transformation and noise)
 - **Distinctiveness** (to be easy to distinguish or match)
 - **Locality** (due to occlusion)
 - Quantity (sufficient number), accuracy (localization), efficiency (computing time), ...

Feature Point) Harris Corner (1988)

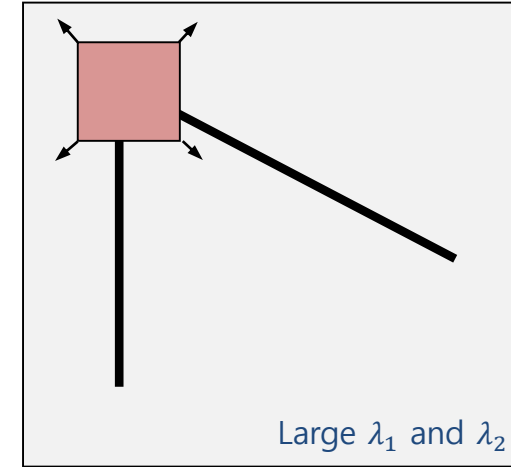
- Key idea: **Sliding window**



“flat” region:
no change
in all directions



“edge”:
no change
along the edge direction



“corner”:
significant change
in all directions

– Formulation

- $I(x + \Delta_x, y + \Delta_y) \approx I(x, y) + [I_x(x, y) \quad I_y(x, y)] \begin{bmatrix} \Delta_x \\ \Delta_y \end{bmatrix}$ where $I_x = \frac{\partial I}{\partial x}$ and $I_y = \frac{\partial I}{\partial y}$
- $D(\Delta_x, \Delta_y) = \sum_{(x,y) \in W} (I(x + \Delta_x, y + \Delta_y) - I(x, y))^2 \approx [\Delta_x \quad \Delta_y] \underbrace{\begin{bmatrix} \sum_W I_x^2 & \sum_W I_x I_y \\ \sum_W I_x I_y & \sum_W I_y^2 \end{bmatrix}}_M [\Delta_x \\ \Delta_y]$
- Two eigen values of **M**: λ_1 and λ_2

Feature Point) Harris Corner (1988)

- Key idea: **Sliding window**

- Formulation

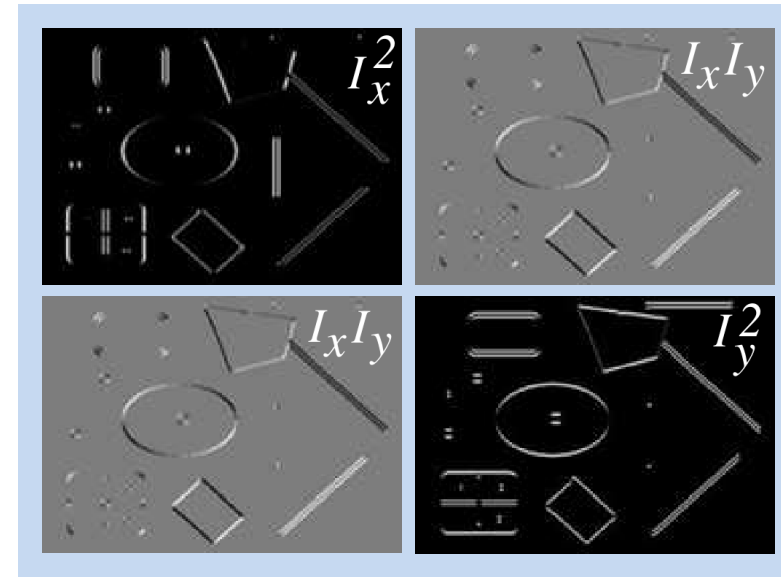
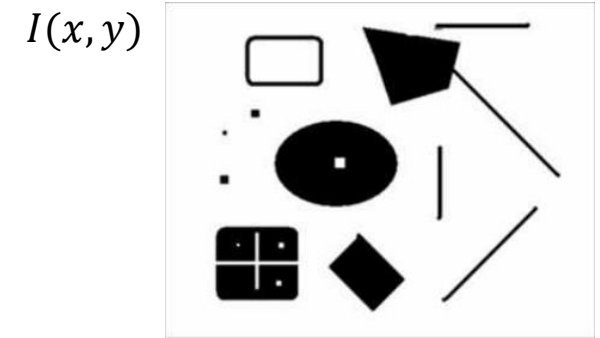
- $I(x + \Delta_x, y + \Delta_y) \approx I(x, y) + [I_x(x, y) \ I_y(x, y)] \begin{bmatrix} \Delta_x \\ \Delta_y \end{bmatrix}$ where $I_x = \frac{\partial I}{\partial x}$ and $I_y = \frac{\partial I}{\partial y}$
 - $D(\Delta_x, \Delta_y) = \sum_{(x,y) \in W} (I(x + \Delta_x, y + \Delta_y) - I(x, y))^2 \approx [\Delta_x \ \Delta_y] \underbrace{\begin{bmatrix} \sum_W I_x^2 & \sum_W I_x I_y \\ \sum_W I_x I_y & \sum_W I_y^2 \end{bmatrix}}_M [\Delta_x \ \Delta_y]$

- **Harris corner response**

- $\text{cornerness} = \det(M) - k \text{trace}(M)^2$
 - Note) $\det(M) = \lambda_1 \lambda_2$, $\text{trace}(M) = \lambda_1 + \lambda_2$, and $k \in [0.04, 0.06]$

- Note) **Good-Feature-to-Track** (a.k.a. GFTT or Shi-Tomasi corner; 1994)

- $\text{cornerness} = \min(\lambda_1, \lambda_2)$



M



$\det(M) - k \text{trace}(M)^2$

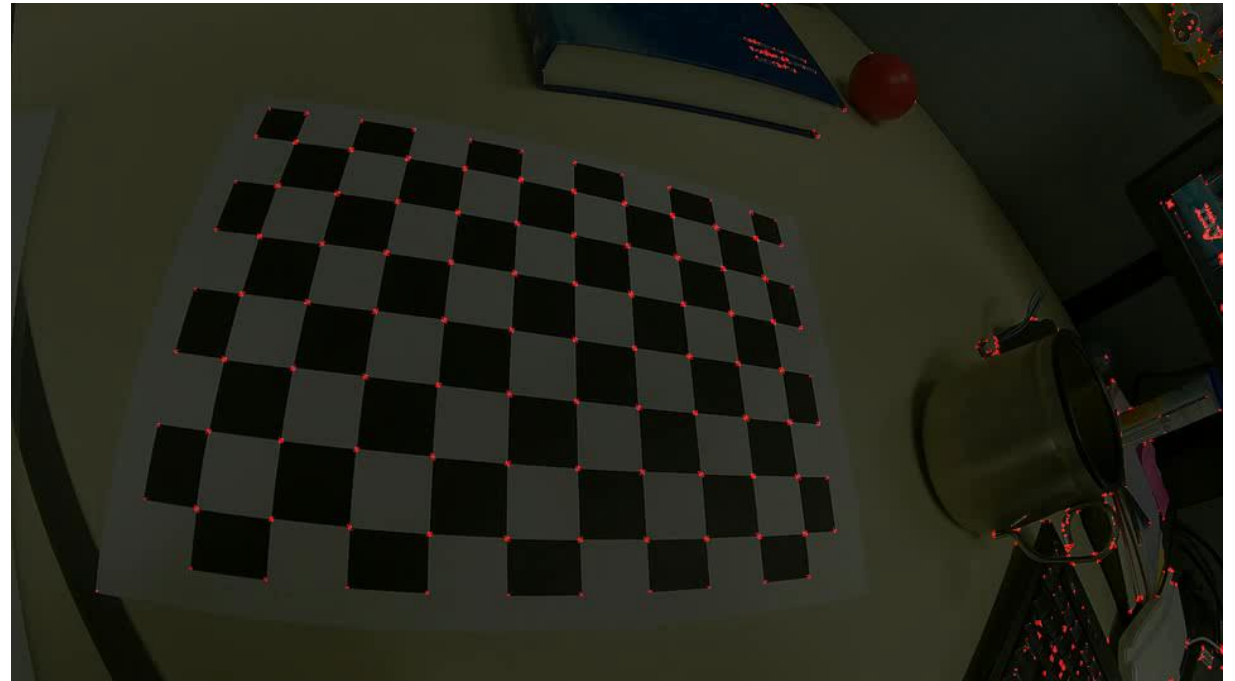
Feature Point) Harris Corner (1988)

- Example) **Harris corner implementation** [harris_corner_implement.py]

```
def cornerHarris(img, ksize=3, k=0.04):  
    # Compute gradients and M matrix  
    Ix = cv.Sobel(img, cv.CV_32F, 1, 0)  
    Iy = cv.Sobel(img, cv.CV_32F, 0, 1)  
    M11 = cv.GaussianBlur(Ix*Ix, (ksize, ksize), 0)  
    M22 = cv.GaussianBlur(Iy*Iy, (ksize, ksize), 0)  
    M12 = cv.GaussianBlur(Ix*Iy, (ksize, ksize), 0)  
  
    # Compute Harris cornerness  
    detM = M11 * M22 - M12 * M12  
    traceM = M11 + M22  
    cornerness = detM - k * traceM**2  
    return cornerness
```

$$M = \begin{bmatrix} \sum_w I_x^2 & \sum_w I_x I_y \\ \sum_w I_x I_y & \sum_w I_y^2 \end{bmatrix}$$

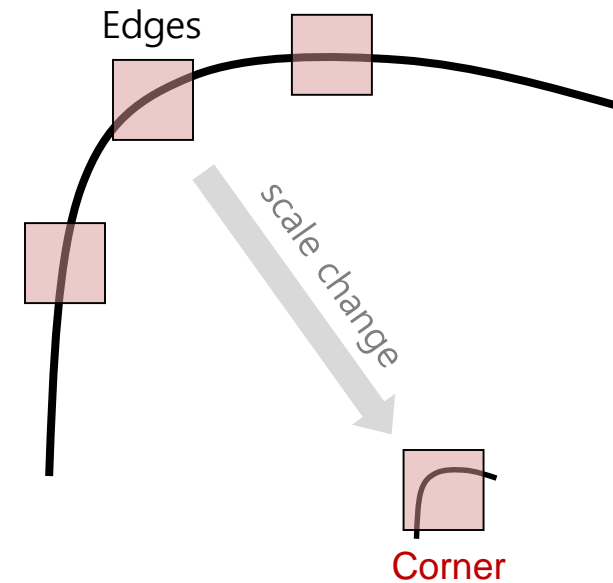
$$\text{cornerness} = \det(M) - k \text{trace}(M)^2$$



Feature Point) Harris Corner (1988)

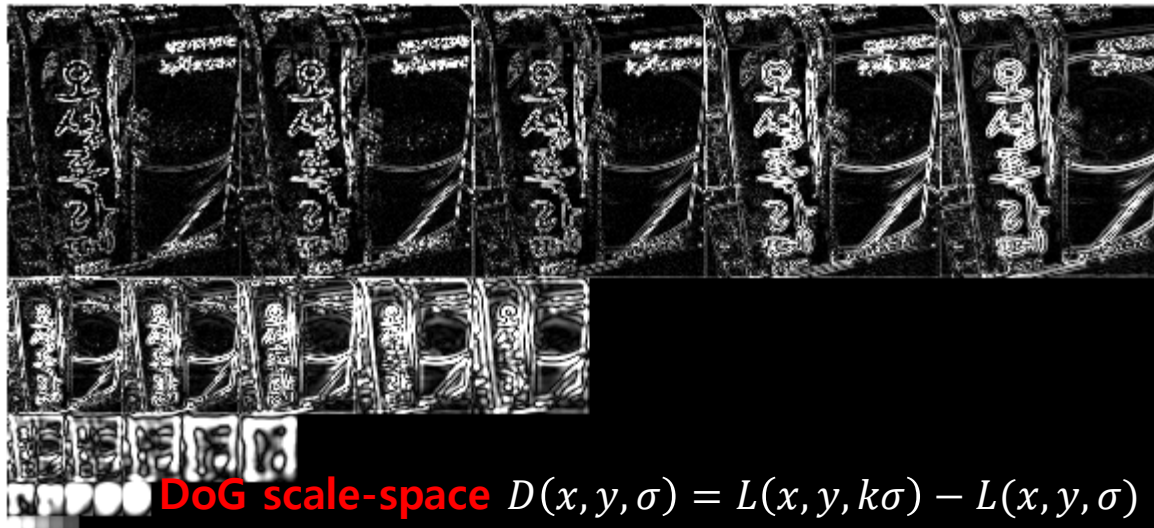
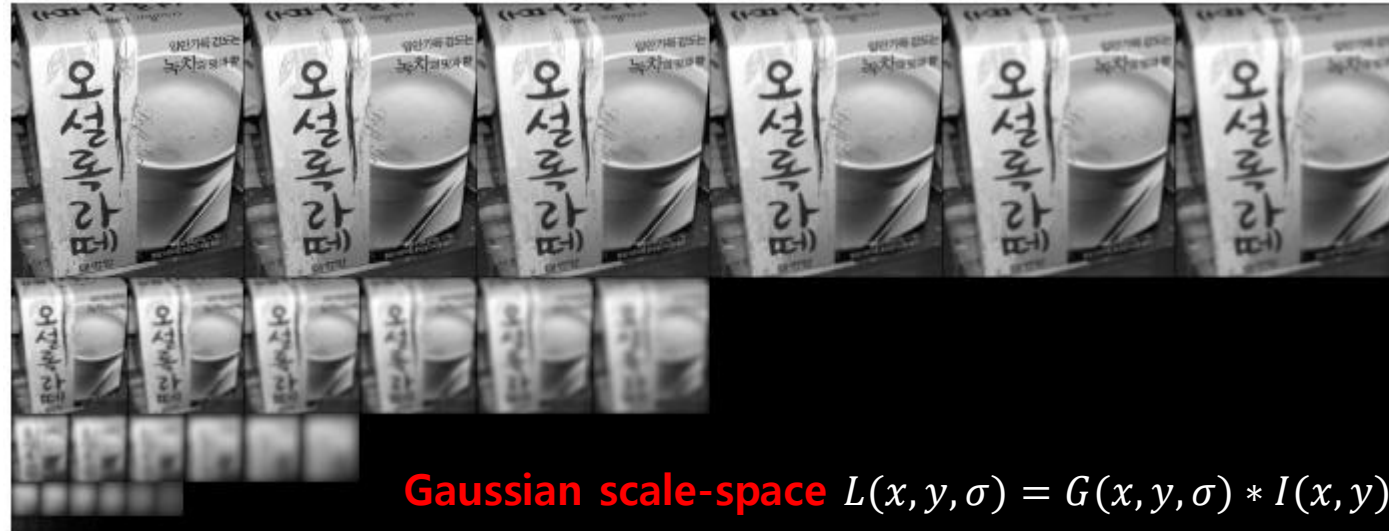
- Properties

- Invariant to translation, rotation, and intensity shift ($I \rightarrow I + b$) ~~intensity scaling ($I \rightarrow aI$)~~
- Variant to **image scaling**



Feature Point) SIFT (Scale-Invariant Feature Transform; 1999)

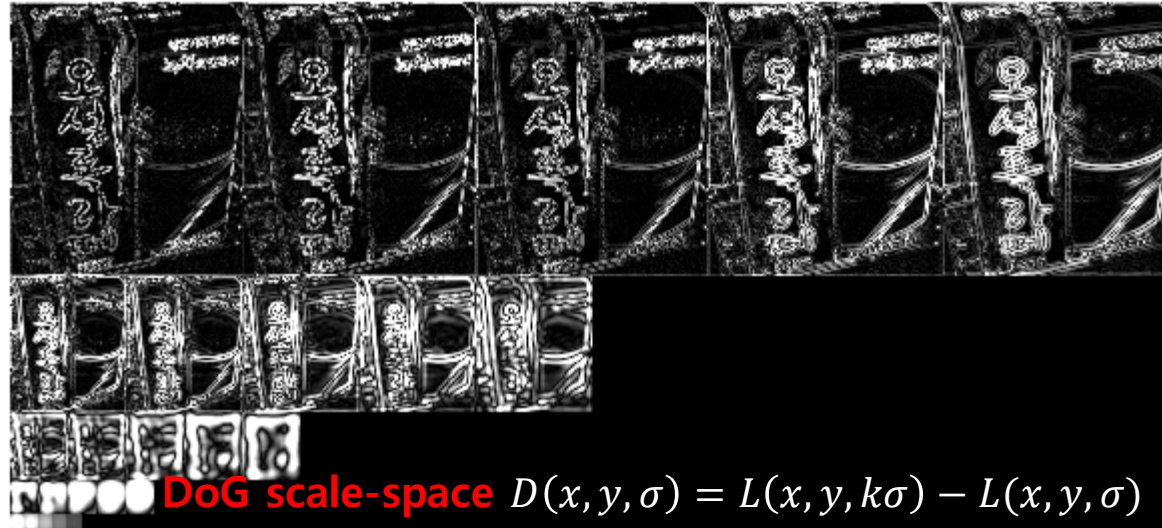
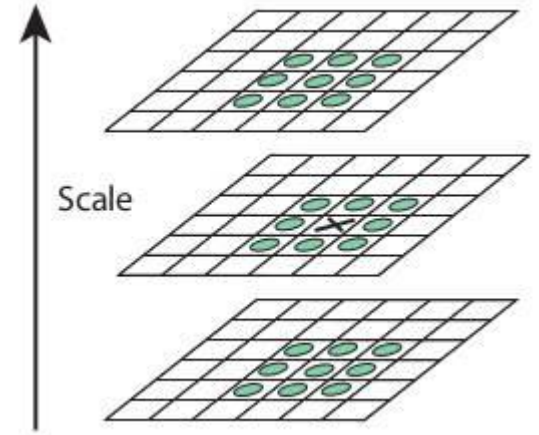
- Key idea: **Scale-space** (~ [image pyramid](#)) and **DoG** ([difference of Gaussian](#))



Feature Point) SIFT (Scale-Invariant Feature Transform; 1999)

- Key idea: **Scale-space** (~ [image pyramid](#)) and **DoG** ([difference of Gaussian](#))
- Part #1) **Feature point detection**
 1. Find **local extrema** (minima and maxima) in DoG scale-space
 2. Localize their position accurately (sub-pixel level) using 3D quadratic function
 3. Eliminate **low contrast candidates**, $|D(\mathbf{x})| < \tau$
 4. Eliminate **candidates on edges**,

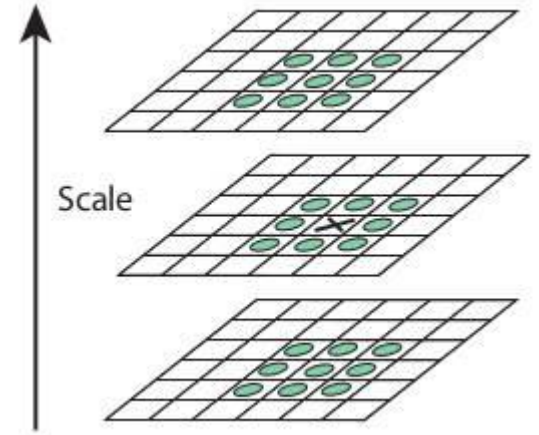
$$\frac{\text{trace}(H)^2}{\det(H)} < \frac{(r+1)^2}{r} \quad \text{where} \quad H = \begin{bmatrix} D_{xx} & D_{xy} \\ D_{xy} & D_{yy} \end{bmatrix}$$



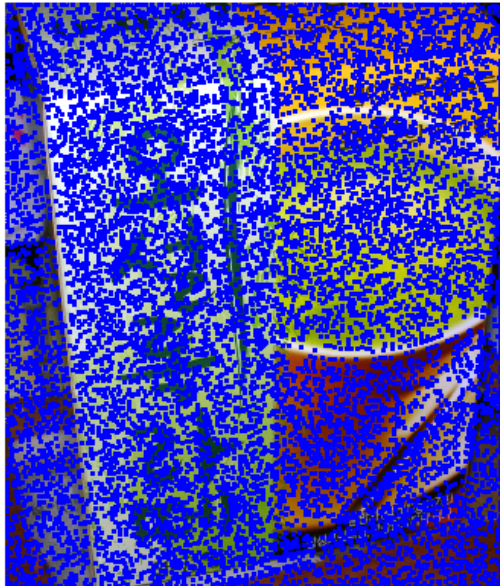
DoG scale-space $D(x, y, \sigma) = L(x, y, k\sigma) - L(x, y, \sigma)$

Feature Point) SIFT (Scale-Invariant Feature Transform; 1999)

- Key idea: **Scale-space** (~ [image pyramid](#)) and **DoG** ([difference of Gaussian](#))
- Part #1) **Feature point detection**
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$$\frac{\text{trace}(H)^2}{\det(H)} < \frac{(r+1)^2}{r} \quad \text{where} \quad H = \begin{bmatrix} D_{xx} & D_{xy} \\ D_{xy} & D_{yy} \end{bmatrix}$$



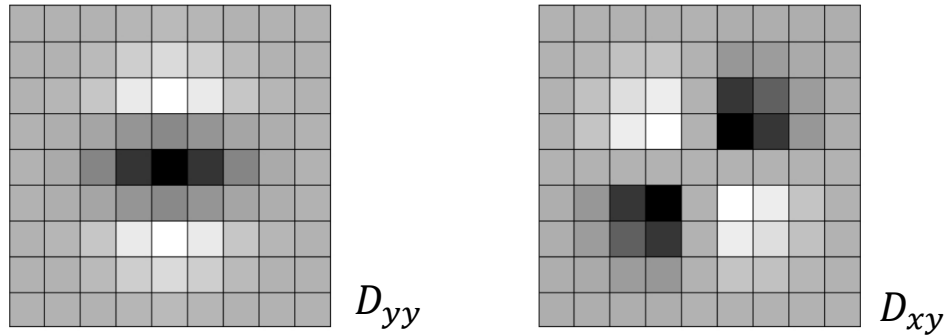
Local extrema (N: 11479)



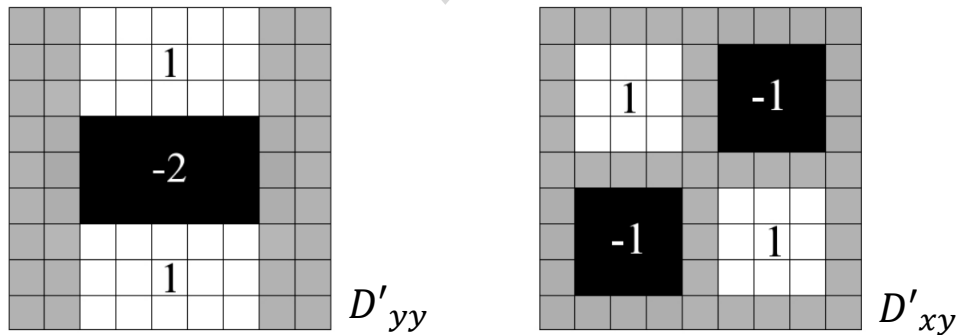
Feature points (N: 971)

Feature Point) SURF (Speeded Up Robust Features; 2006)

- Key idea: **Approximation of SIFT**
 - e.g. DoG approximation Haar-like features and **integral image**



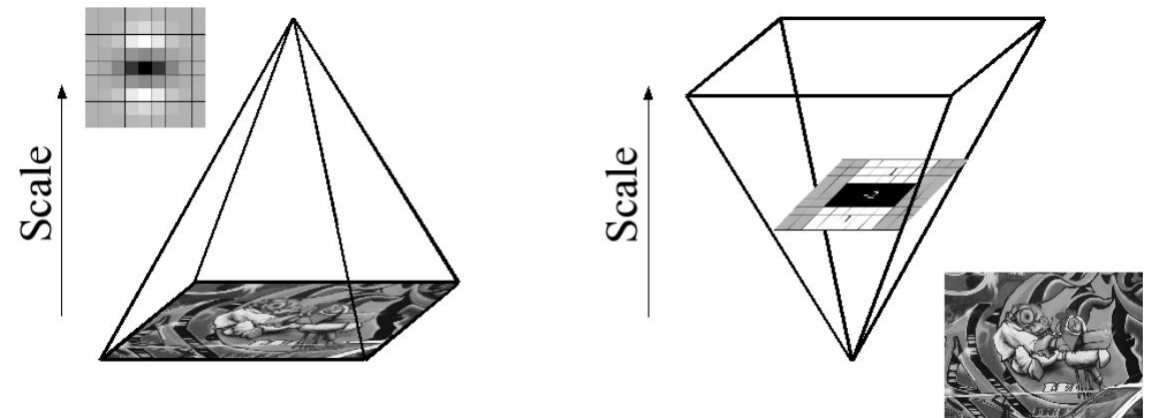
DoG approximation



O

$$S(x, y) = \sum_{i=0}^x \sum_{j=0}^y I(i, j)$$

$\Sigma = A - B - C + D$



Feature Descriptor) SIFT (Scale-Invariant Feature Transform; 1999)

■ Part #2) Orientation assignment

1. Derive magnitude and orientation of gradient of each patch

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

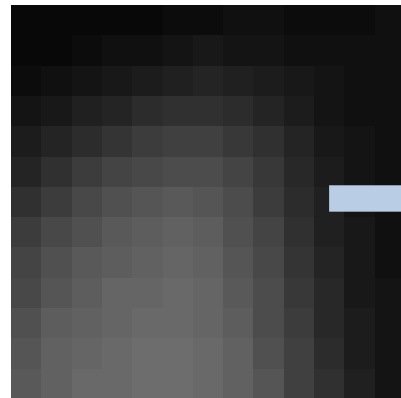
$$\theta(x, y) = \tan^{-1} \frac{L(x, y+1) - L(x, y-1)}{L(x+1, y) - L(x-1, y)}$$

2. Find **the strongest orientation**

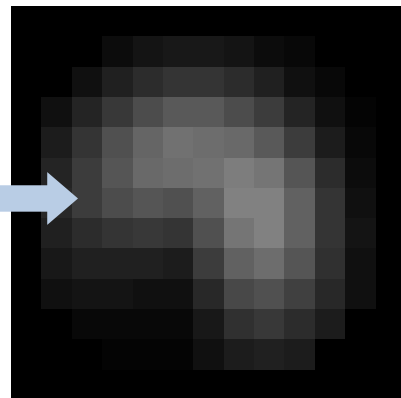
- Histogram voting (36 bins) with Gaussian-weighted magnitude



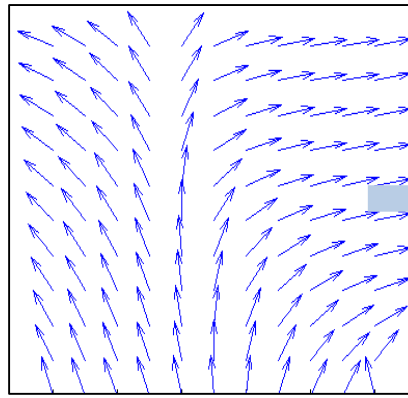
Feature scales and orientations



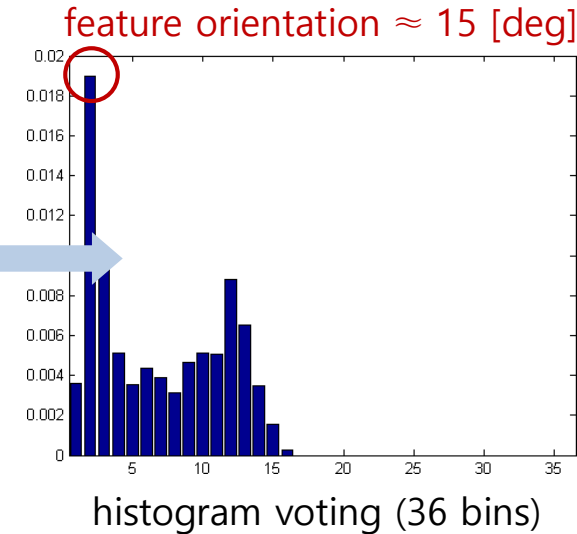
feature patch



gradient magnitude
 $m(x, y)$



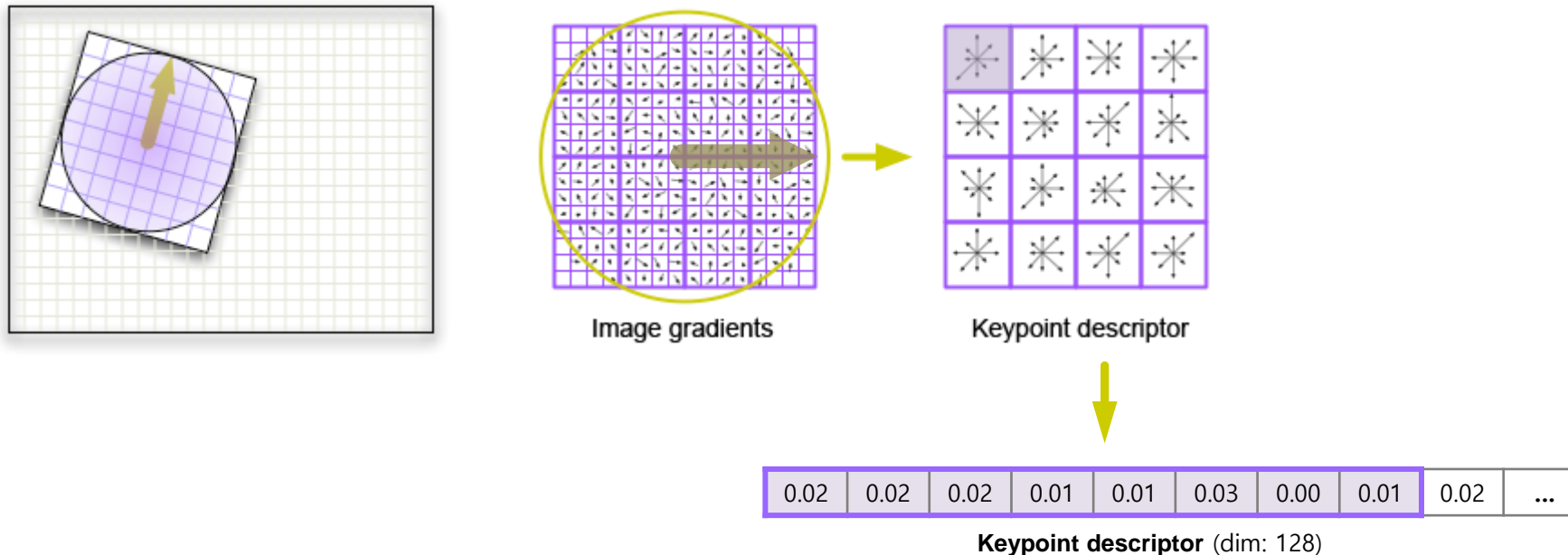
gradient orientation
 $\theta(x, y)$



Feature Descriptor) SIFT (Scale-Invariant Feature Transform; 1999)

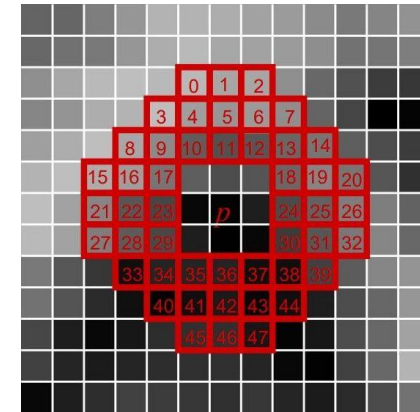
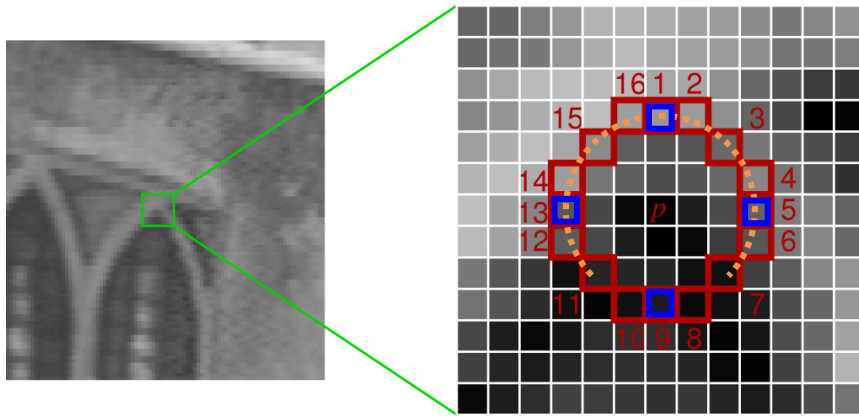
■ Part #3) Feature descriptor extraction

1. Build a 4x4 gradient histogram (8 bins) from each patch (16x16 pixels)
 - Use Gaussian-weighted magnitude again
 - Use relative angles w.r.t. the assigned feature orientation
2. Encode the histogram into a 128-dimensional vector
 - Apply normalization to be an unit vector



Feature Point) FAST (Features from Accelerated Segment Test; 2006)

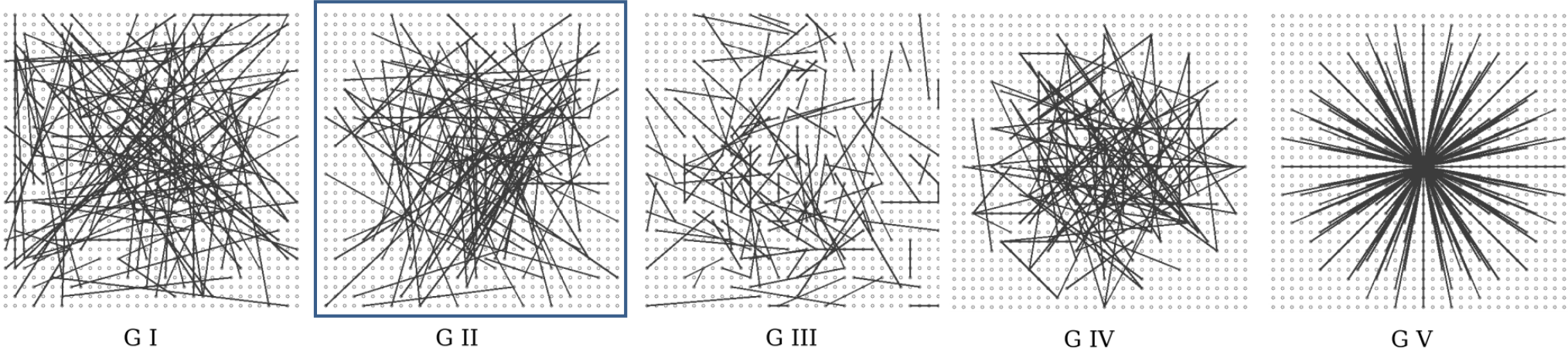
- Key idea: **Intensity check of continuous arc of n pixels**
 - Is this point p a corner? (I_p : intensity at p , t : the intensity threshold)
 - Is a segment of n continuous pixels brighter than $I_p + t$? (OR) Is the segment darker than $I_p - t$?
 - Note) High-speed non-corner rejection: Checking intensity values at 1, 9, 5, and 13 (n : 12)



- Too many corners! → Non-maximum suppression
- Versions
 - FAST-9 (n : 9; `cv.FastFeatureDetector_TYPE_9_16`), FAST-12 (n : 12), ...
 - FAST-ER: Training a decision tree to enhance repeatability with more pixels

Feature Descriptor) BRIEF (Binary Robust Independent Elementary Features; 2010)

- Key idea: **Intensity comparison of a sequence of random pairs (binary test)**
 - Path size: 31 x 31 pixels (Note: Applying smoothing for stability and repeatability)



- Descriptor size: 128 tests (128 bits) → 16 bytes
 - Note) SIFT: 128-dimensional vector → 512 bytes
- Versions: The number of tests
 - BRIEF-32, BRIEF-64, BRIEF-128 (16 bytes), BRIEF-256 (32 bytes), BRIEF-512 (64 bytes), ...
- Combination examples
 - SIFT feature points + BRIEF descriptors
 - FAST feature points + BRIEF descriptors

Feature Point and Descriptor) **ORB (Oriented FAST and rotated BRIEF, 2011)**

- Key idea: **Adding rotation invariance to BRIEF**

- **Oriented FAST**

- Generate scale pyramid for scale invariance
 - Detect *FAST-9* points (filtering with Harris corner response)
 - Calculate feature orientation by *intensity centroid* $C = (\frac{m_{10}}{m_{00}}, \frac{m_{01}}{m_{00}})$

$$\theta = \tan^{-1} \frac{m_{01}}{m_{10}} \quad \text{where} \quad m_{pq} = \sum_{x,y} x^p y^q I(x,y)$$

- **Rotation-aware BRIEF**

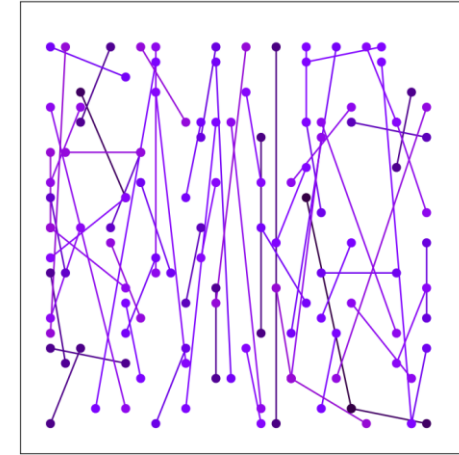
- Extract BRIEF descriptors w.r.t. the known orientation
 - Use better comparison pairs trained by greedy search

- Combination (default): **ORB**

- FAST-9 detector (with orientation) + BRIEF-256 descriptor (with trained pairs)

- Computing time

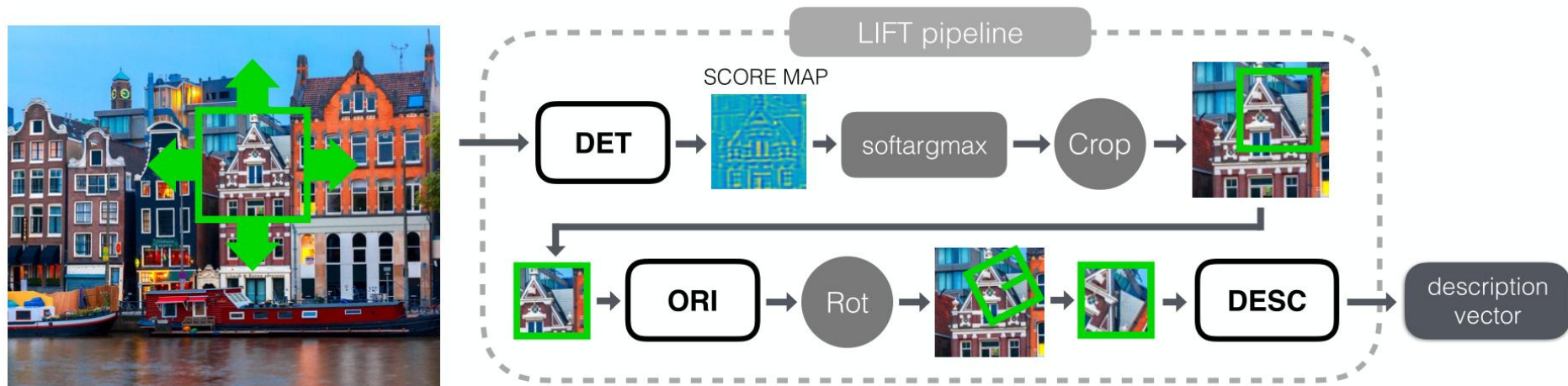
- ORB: **15.3 [msec]** / SURF: 217.3 [msec] / SIFT: 5228.7 [msec] @ 24 images (640x480) in Pascal dataset



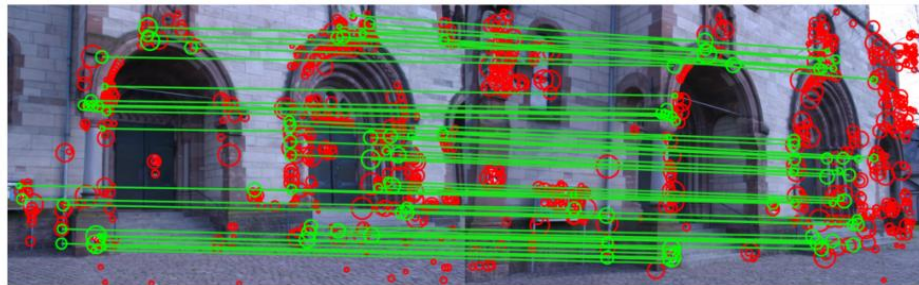
Feature Point and Descriptor) **LIFT (Learned Invariant Feature Transform; 2016)**

- Key idea: **Deep neural network**

- CNN network: **DET** (feature detector) + **ORI** (orientation estimator) + **DESC** (feature descriptor)
- Training data: Photo Tourism dataset with [VisualSFM](#) (SIFT)



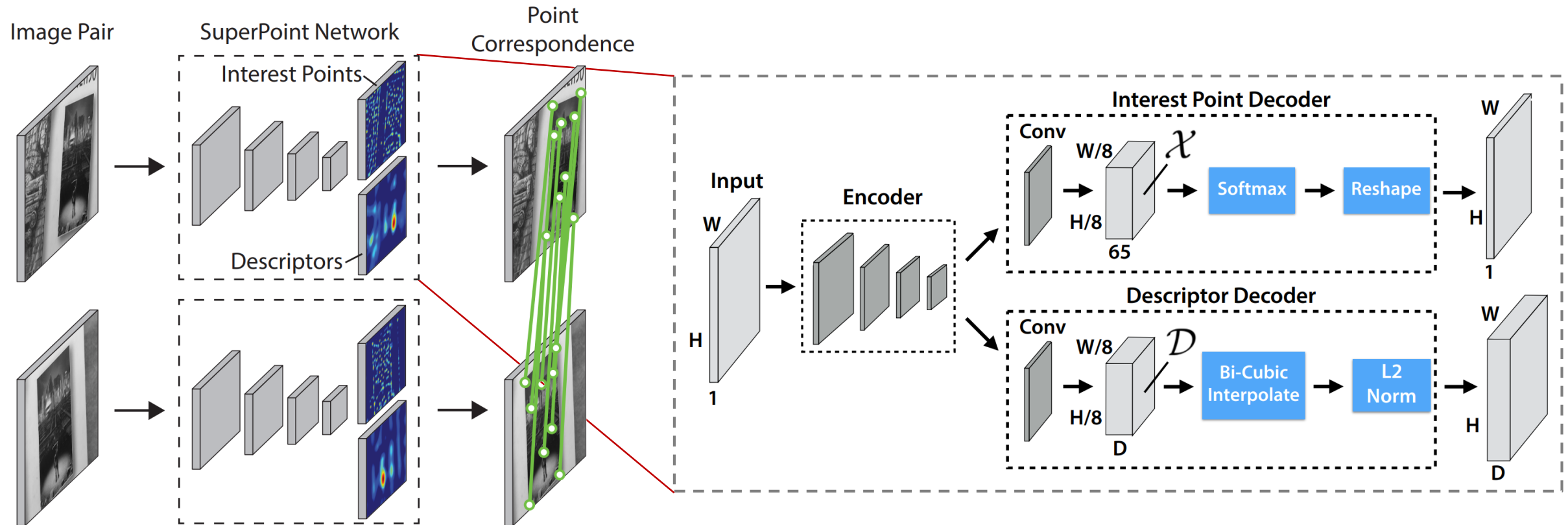
SIFT



LIFT

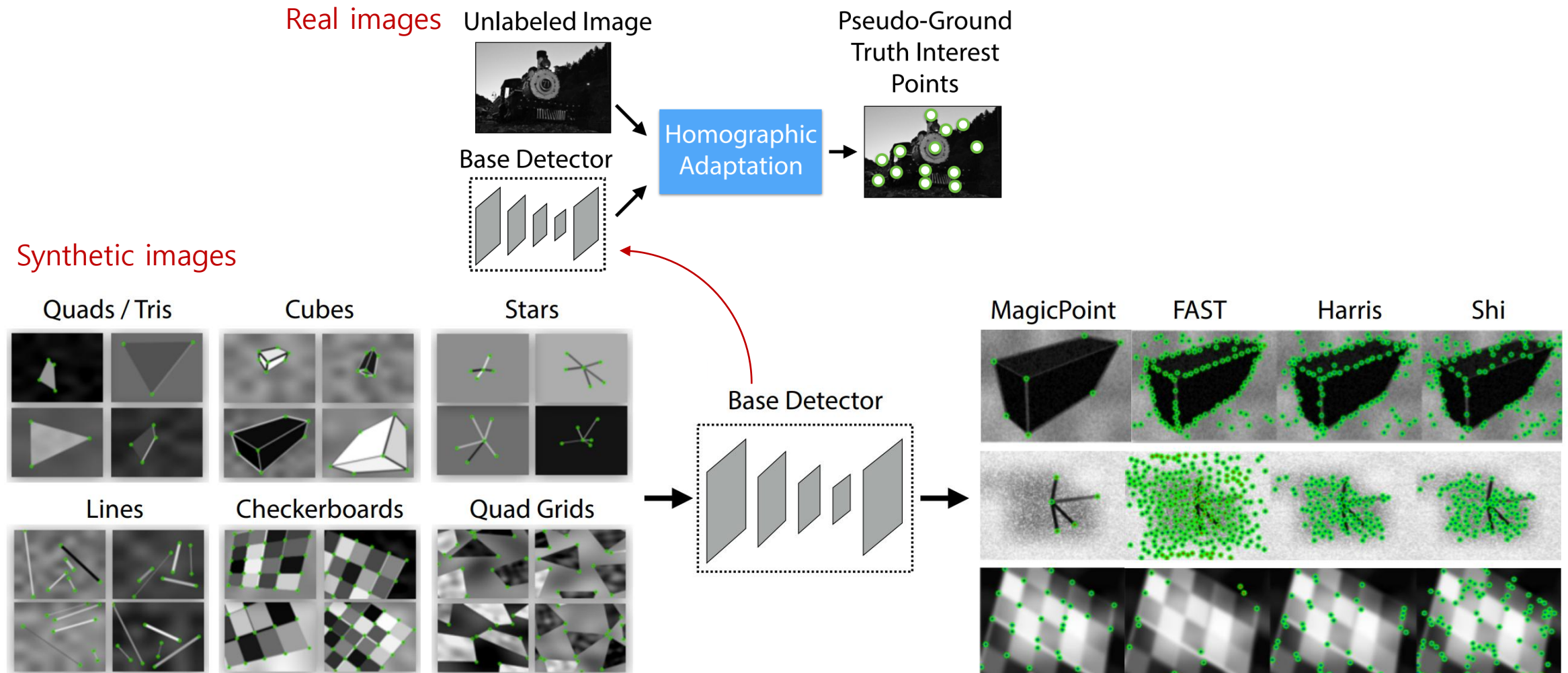
Feature Point and Descriptor) SuperPoint (2017)

- Key idea: **Self-supervised training with homography transformation**
 - CNN network: Encoder (~ VGG) + Decoders (for point and descriptor)



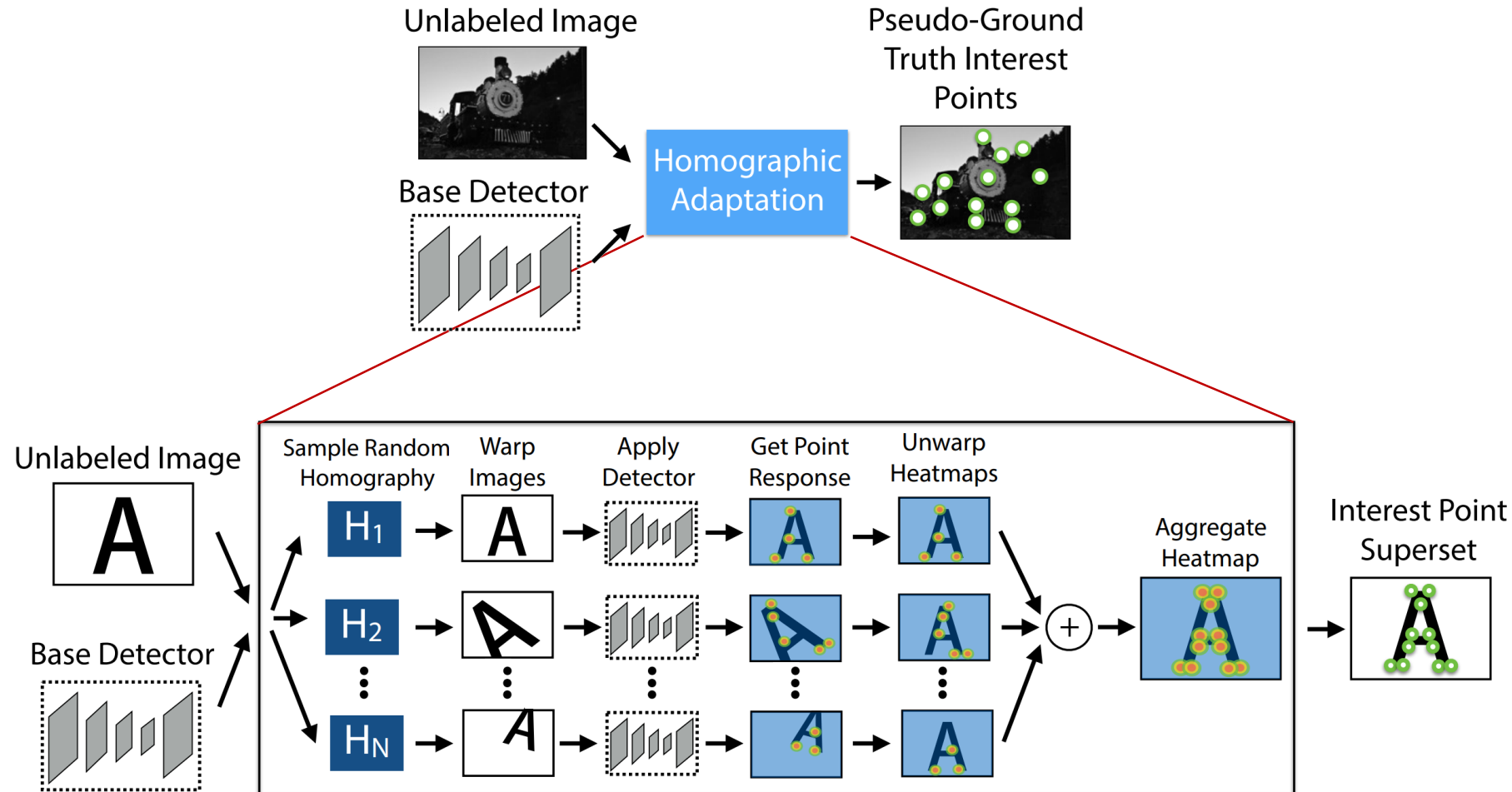
Feature Point and Descriptor) **SuperPoint (2017)**

- Key idea: **Self-supervised training with homography transformation**
 - Training data generation: The base detector, *MagicPoint* → Ground truth interest points



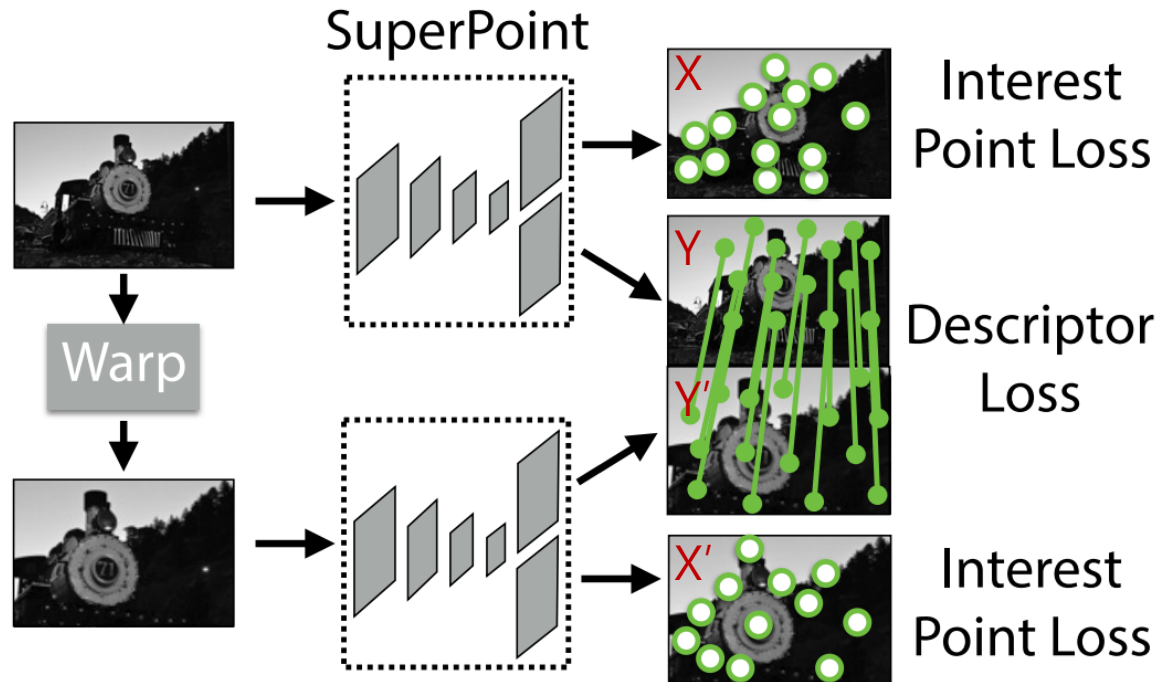
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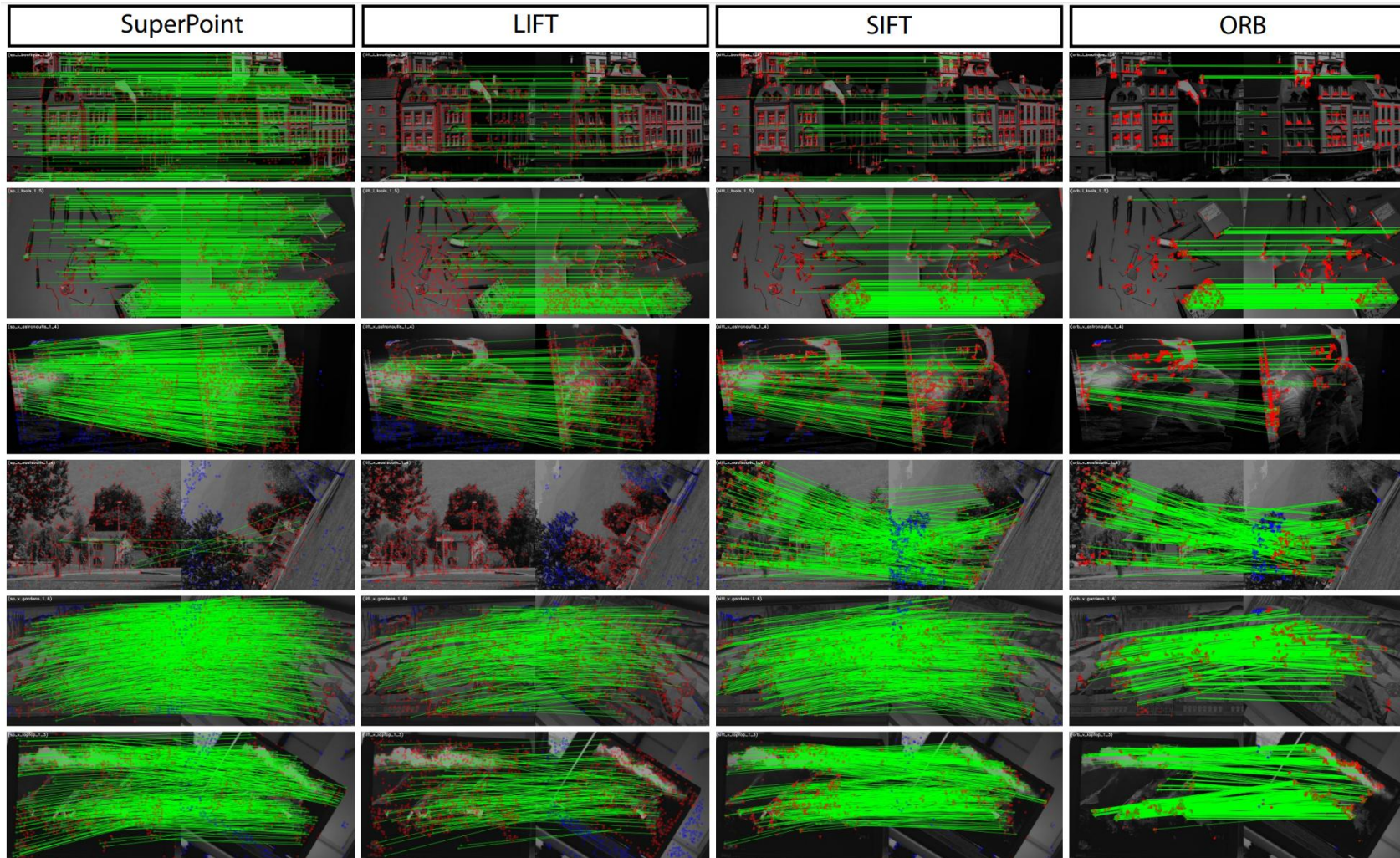
Feature Point and Descriptor) **SuperPoint (2017)**

- Key idea: **Self-supervised training with homography transformation**
 - Training data augmentation: Random homography transformation
 - Loss functions: Interest Point Loss (X, Y) + Interest Point Loss (X', Y') + **Descriptor Loss (Y, Y')**



Feature Point and Descriptor) SuperPoint (2017)

- Key idea: **Self-supervised training with homography transformation**
 - Real-time performance: **70 FPS** (13 msec) on 640x480 images with NVIDIA Titan X GPU



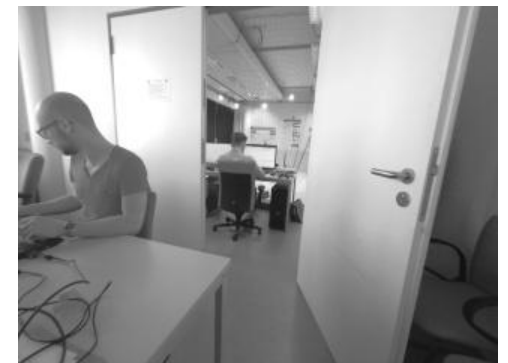
Freiburg RGBD:



Microsoft 7 Scenes:



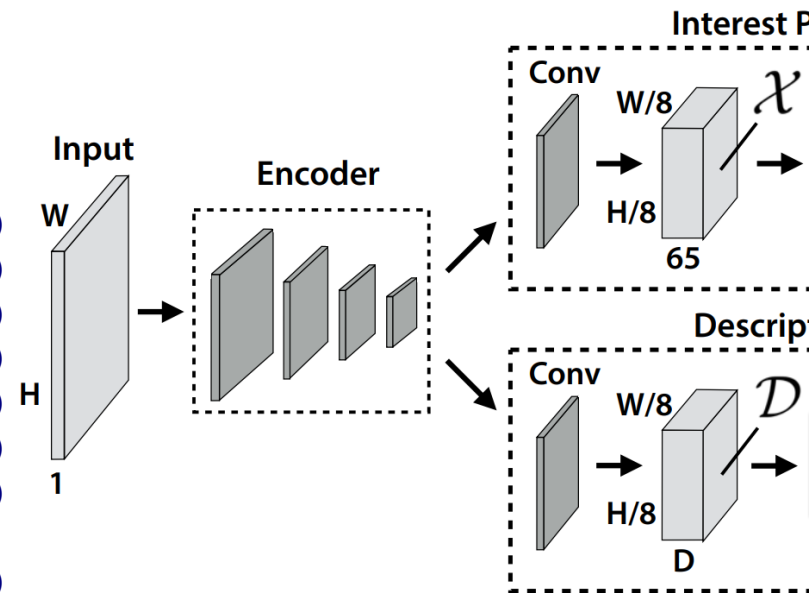
MonoVO:



Feature Point and Descriptor) SuperPoint (2017)

- Example) **SuperPoint implementation** [demo_superpoint.py] @ [Github \(authors\)](#)

```
class SuperPointNet(torch.nn.Module):
    """ Pytorch definition of SuperPoint Network. """
    def __init__(self):
        super(SuperPointNet, self).__init__()
        self.relu = torch.nn.ReLU(inplace=True)
        self.pool = torch.nn.MaxPool2d(kernel_size=2, stride=2)
        c1, c2, c3, c4, c5, d1 = 64, 64, 128, 128, 256, 256
        # Shared Encoder.
        self.conv1a = torch.nn.Conv2d(1, c1, kernel_size=3, stride=1, padding=1)
        self.conv1b = torch.nn.Conv2d(c1, c1, kernel_size=3, stride=1, padding=1)
        self.conv2a = torch.nn.Conv2d(c1, c2, kernel_size=3, stride=1, padding=1)
        self.conv2b = torch.nn.Conv2d(c2, c2, kernel_size=3, stride=1, padding=1)
        self.conv3a = torch.nn.Conv2d(c2, c3, kernel_size=3, stride=1, padding=1)
        self.conv3b = torch.nn.Conv2d(c3, c3, kernel_size=3, stride=1, padding=1)
        self.conv4a = torch.nn.Conv2d(c3, c4, kernel_size=3, stride=1, padding=1)
        self.conv4b = torch.nn.Conv2d(c4, c4, kernel_size=3, stride=1, padding=1)
        # Detector Head.
        self.convPa = torch.nn.Conv2d(c4, c5, kernel_size=3, stride=1, padding=1)
        self.convPb = torch.nn.Conv2d(c5, 65, kernel_size=1, stride=1, padding=0)
        # Descriptor Head.
        self.convDa = torch.nn.Conv2d(c4, c5, kernel_size=3, stride=1, padding=1)
        self.convDb = torch.nn.Conv2d(c5, d1, kernel_size=1, stride=1, padding=0)
```



```
class SuperPointNet(torch.nn.Module):
```

```
...
```

```
def forward(self, x):
```

```
    """ Forward pass that jointly computes unprocessed point and descriptor
    tensors.
```

```
    Input
```

```
    x: Image pytorch tensor shaped N x 1 x H x W.
```

```
    Output
```

```
    semi: Output point pytorch tensor shaped N x 65 x H/8 x W/8. (Note: 65 channels includes a dustbin.)
```

```
    desc: Output descriptor pytorch tensor shaped N x 256 x H/8 x W/8.
```

```
    """
```

```
# Shared Encoder.
```

```
x = self.relu(self.conv1a(x))
```

```
x = self.relu(self.conv1b(x))
```

```
x = self.pool(x)
```

```
x = self.relu(self.conv2a(x))
```

```
x = self.relu(self.conv2b(x))
```

```
x = self.pool(x)
```

```
x = self.relu(self.conv3a(x))
```

```
x = self.relu(self.conv3b(x))
```

```
x = self.pool(x)
```

```
x = self.relu(self.conv4a(x))
```

```
x = self.relu(self.conv4b(x))
```

```
# Detector Head.
```

```
cPa = self.relu(self.convPa(x))
```

```
semi = self.convPb(cPa)
```

```
# Descriptor Head.
```

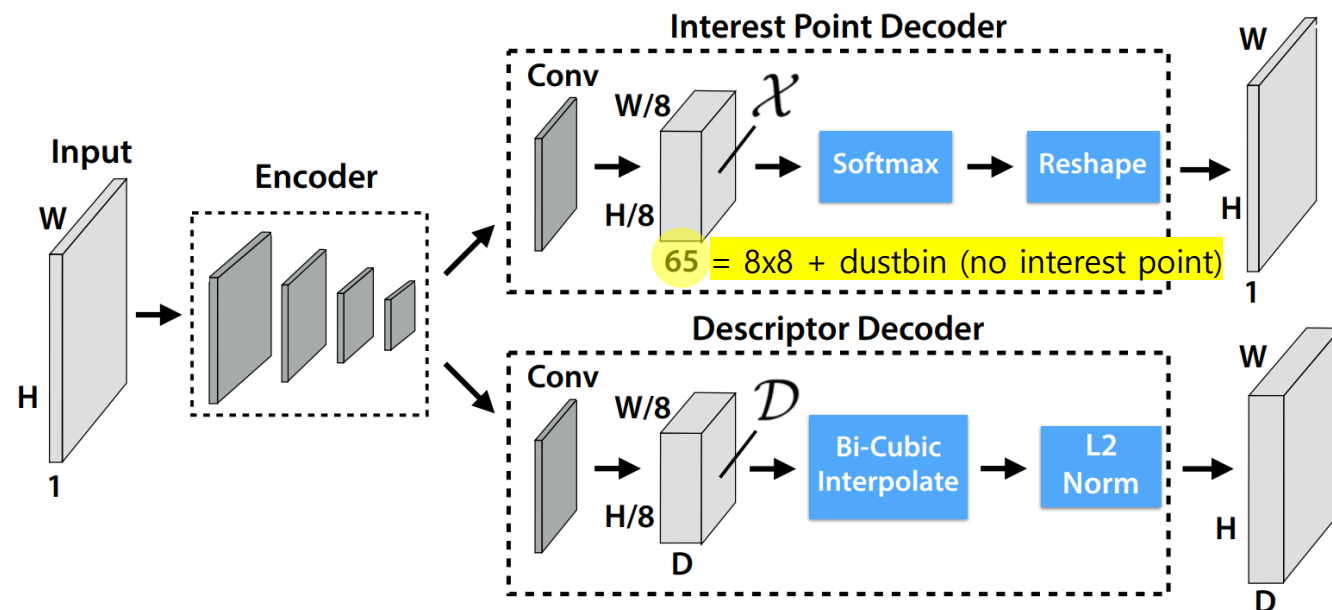
```
cDa = self.relu(self.convDa(x))
```

```
desc = self.convDb(cDa)
```

```
dn = torch.norm(desc, p=2, dim=1) # Compute the norm.
```

```
desc = desc.div(torch.unsqueeze(dn, 1)) # Divide by norm to normalize.
```

```
return semi, desc
```




```
class SuperPointNet(torch.nn.Module):
```

```
...
```

```
def run(self, img):
```

```
    """ Process a numpy image to extract points and descriptors.
```

```
    Input
```

```
    img - HxW numpy float32 input image in range [0,1].
```

```
    Output
```

```
    corners - 3xN numpy array with corners [x_i, y_i, confidence_i]^T.
```

```
    desc - 256xN numpy array of corresponding unit normalized descriptors.
```

```
    heatmap - HxW numpy heatmap in range [0,1] of point confidences.
```

```
    """
```

```
...
```

```
# Forward pass of network.
```

```
outs = self.net.forward(inp)
```

```
semi, coarse_desc = outs[0], outs[1]
```

```
# Convert pytorch -> numpy.
```

```
semi = semi.data.cpu().numpy().squeeze()
```

```
# --- Process points.
```

```
dense = np.exp(semi) # Softmax.
```

```
dense = dense / (np.sum(dense, axis=0)+.00001) # Should sum to 1.
```

```
# Remove dustbin.
```

```
nodust = dense[:-1, :, :]
```

```
# Reshape to get full resolution heatmap.
```

```
Hc = int(H / self.cell)
```

```
Wc = int(W / self.cell)
```

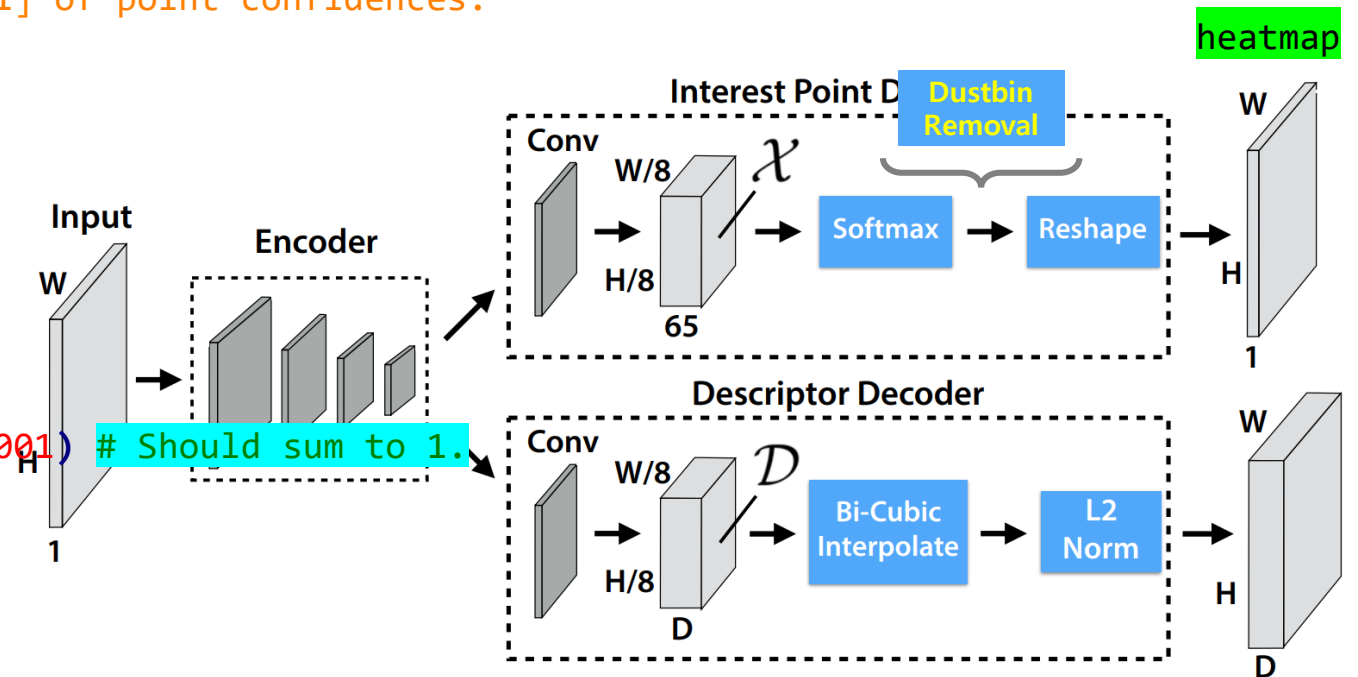
```
nodust = nodust.transpose(1, 2, 0)
```

```
heatmap = np.reshape(nodust, [Hc, Wc, self.cell, self.cell])
```

```
heatmap = np.transpose(heatmap, [0, 2, 1, 3])
```

```
heatmap = np.reshape(heatmap, [Hc*self.cell, Wc*self.cell])
```

```
...
```



```
class SuperPointNet(torch.nn.Module):
```

```
...
```

```
def run(self, img):
```

```
    """ Process a numpy image to extract points and descriptors.
```

```
    Input
```

```
    img - HxW numpy float32 input image in range [0,1].
```

```
    Output
```

```
    corners - 3xN numpy array with corners [x_i, y_i, confidence_i]^T.
```

```
    desc - 256xN numpy array of corresponding unit normalized descriptors.
```

```
    heatmap - HxW numpy heatmap in range [0,1] of point confidences.
```

```
    """
```

```
...
```

```
# --- Process descriptor.
```

```
D = coarse_desc.shape[1]
```

```
if pts.shape[1] == 0:
```

```
    desc = np.zeros((D, 0))
```

```
else:
```

```
    # Interpolate into descriptor map using 2D point locations.
```

```
    samp_pts = torch.from_numpy(pts[:,2, :].copy())
```

```
    samp_pts[0, :] = (samp_pts[0, :] / (float(W)/2.)) - 1.
```

```
    samp_pts[1, :] = (samp_pts[1, :] / (float(H)/2.)) - 1.
```

```
    samp_pts = samp_pts.transpose(0, 1).contiguous()
```

```
    samp_pts = samp_pts.view(1, 1, -1, 2)
```

```
    samp_pts = samp_pts.float()
```

```
    if self.cuda:
```

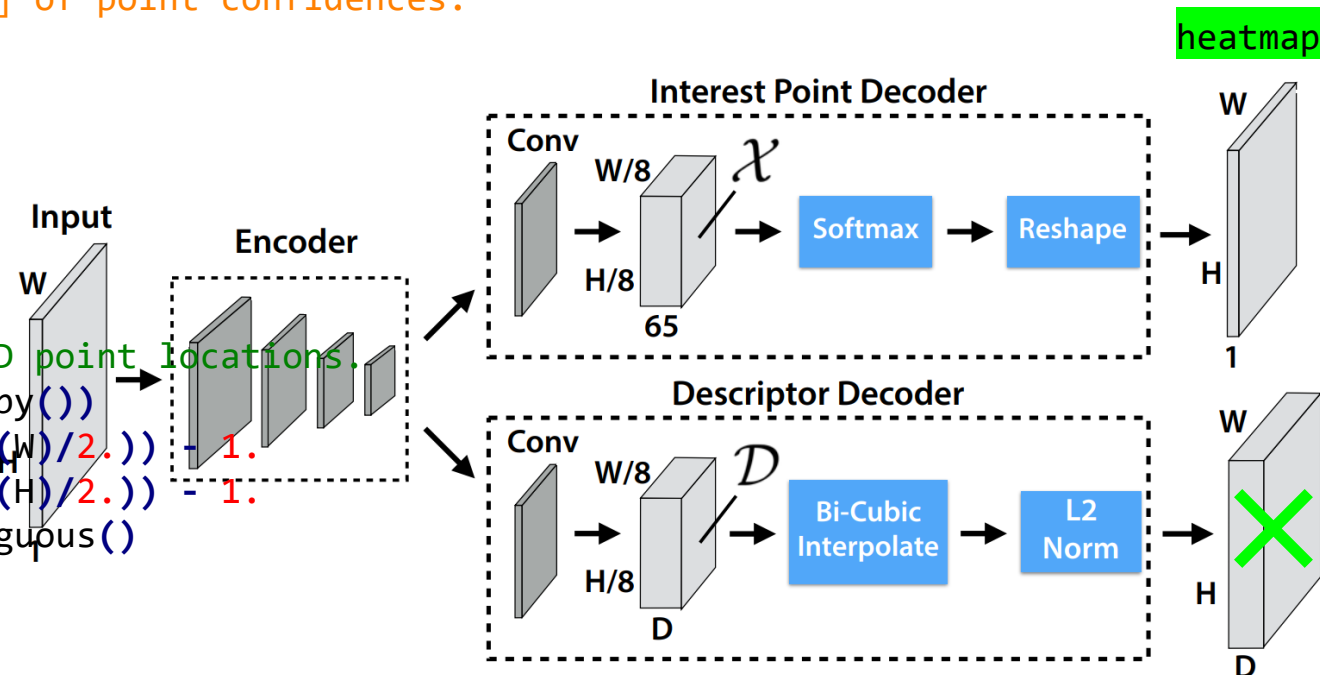
```
        samp_pts = samp_pts.cuda()
```

```
    desc = torch.nn.functional.grid_sample(coarse_desc, samp_pts)
```

```
    desc = desc.data.cpu().numpy().reshape(D, -1)
```

```
    desc /= np.linalg.norm(desc, axis=0)[np.newaxis, :]
```

```
return pts, desc, heatmap
```



Summary) Feature Points and Descriptors

Feature Points	Gradient-based <ul style="list-style-type: none">▪ Harris▪ GFTT (a.k.a. Shi-Tomasi)▪ SIFT▪ SURF	Intensity-based <ul style="list-style-type: none">▪ FAST	DL-based <ul style="list-style-type: none">▪ LIFT▪ SuperPoint
Feature Descriptor	Real-valued <ul style="list-style-type: none">▪ SIFT▪ SURF	Binary-valued <ul style="list-style-type: none">▪ BRIEF▪ ORB (FAST+BRIEF)	Real-valued (DL-based) <ul style="list-style-type: none">▪ LIFT▪ SuperPoint
Advantages Disadvantages	(+) Accurate (-) Slow	(+) Fast (-) Inaccurate (+) Less storage	(+) Accurate (+) Fast (-) GPU requirement

Table of Contents

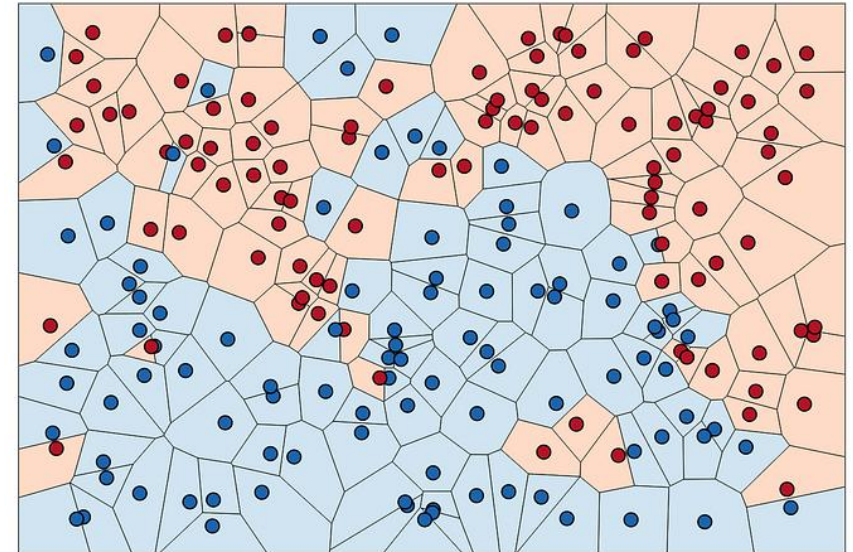
- **Feature Points**
 - Gradient-based: Harris corner, GFTT corner, SIFT, SURF
 - Intensity-based: FAST
 - DL-based: LIFT, SuperPoint
- **Feature Descriptors**
 - Real-valued: SIFT, SURF (DL-based: LIFT, SuperPoint)
 - Binary-valued: BRIEF, ORB
- **Feature Matching**
 - Q) How can we associate the points across different images?
- **Feature Tracking**
 - Q) How can we associate the points across their next image?
- **Outlier Rejection**
 - Q) How can we select correctly matched points?



Feature Matching) For Real-valued Descriptors

▪ Real-valued descriptors

- Distance measures
 - [Euclidean distance](#): $l_2(\mathbf{d}, \mathbf{d}') = \|\mathbf{d} - \mathbf{d}'\|_2$ (↓ : similar)
 - [Cosine similarity](#): $s_c(\mathbf{d}, \mathbf{d}') = \frac{\mathbf{d} \cdot \mathbf{d}'}{\|\mathbf{d}\| \|\mathbf{d}'\|}$ (↑ : similar)
 - Note) Matching measures can be combined or advanced.
 - e.g. The ratio of the best and second best similarity > threshold
 - It may select more distinguishable feature matching.
- Matching algorithms
 - [Brute-force search](#)
 - Time complexity: $O(N)$ for N descriptors
 - Approximated [nearest neighborhood search](#) (ANN search)
 - Time complexity: $O(\log N)$ or less for N descriptors
 - Note) [Big-ANN Competition](#) (recent: NeurIPS 2023)



Feature Matching) For Binary-valued Descriptors

- **Binary-valued descriptors**

- Distance measures

- [Hamming distance](#): $l_h(\mathbf{d}, \mathbf{d}') = \sum_i (d_i \neq d'_i)$

- e.g. 4-bit descriptors

- $l_h(0110, \textcolor{red}{1}110) = 1$ vs. (6, 14)

- $l_h(0110, 011\textcolor{red}{1}) = 1$ vs. (6, 7)

- $l_h(0110, 01\textcolor{red}{01}) = 2$ vs. (6, 5)

- Note) Hamming distance is the L_1 -norm with binary-valued descriptors.

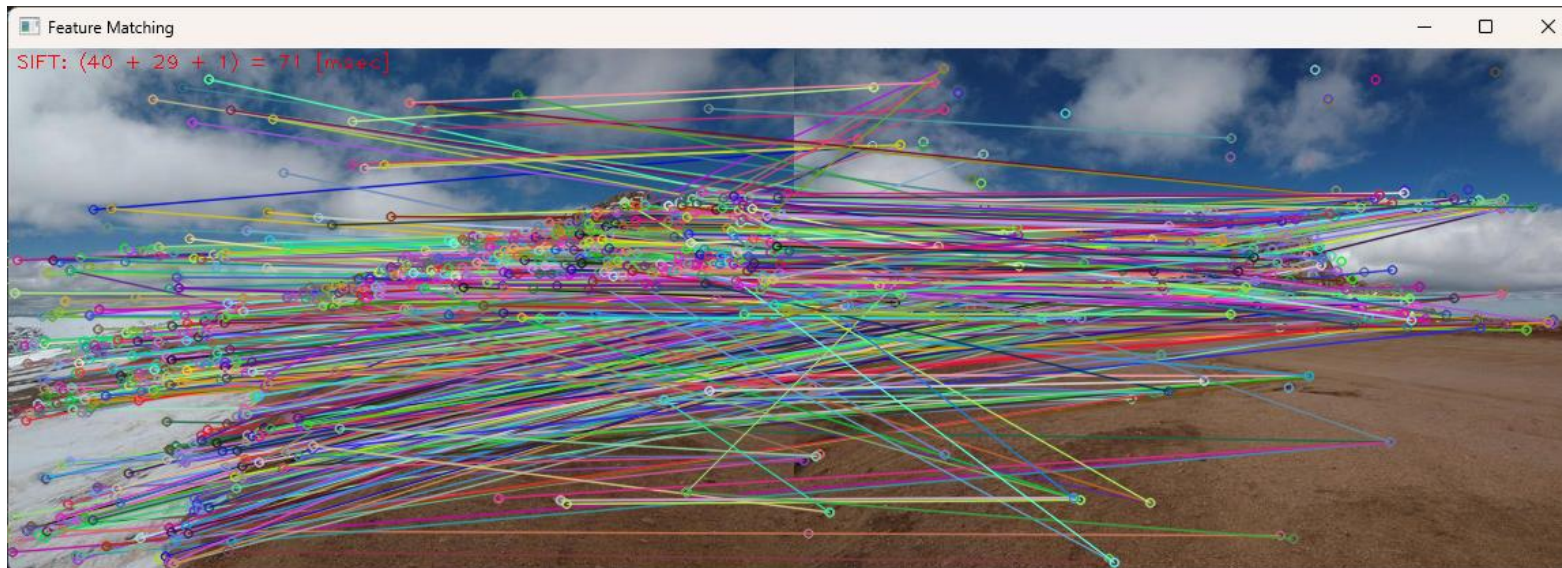
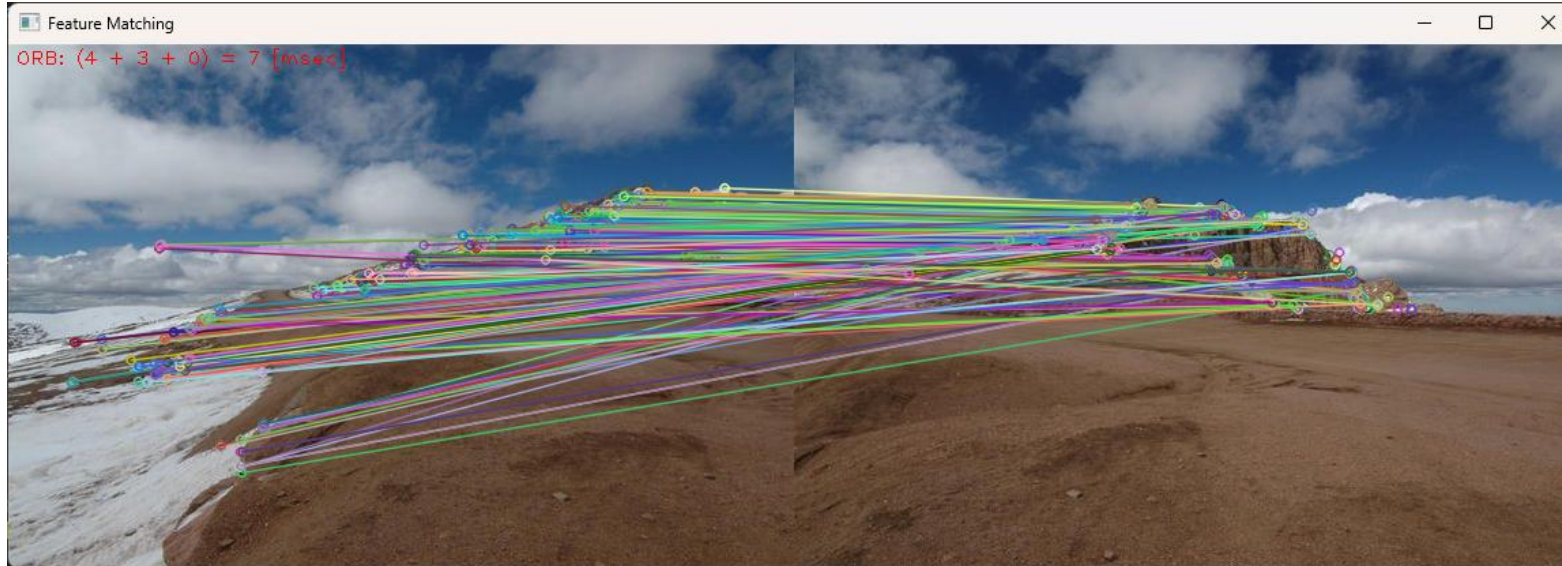
- Matching algorithms

- [Brute-force search](#)

- Time complexity: $O(N)$ for N descriptors

Feature Matching

- Example) **Feature matching comparison** [feature_matching.py]



Feature Matching

- Example) **Feature matching comparison** [feature_matching.py]

```
# Load two images
```

```
img1 = cv.imread('../data/hill01.jpg')
```

```
img2 = cv.imread('../data/hill02.jpg')
```

```
# Instantiate feature detectors and matchers
```

```
# Note) You can specify options for each detector in its creation.
```

```
features = [
```

```
    ...
```

```
    {'name': 'FAST',      'detector': cv.FastFeatureDetector_create(),  
     'matcher' : None}, # No descriptor
```

```
    ...
```

```
    {'name': 'ORB',      'detector': cv.ORB_create(),  
     'matcher' : cv.DescriptorMatcher_create('BruteForce-Hamming')},
```

```
    {'name': 'SIFT',     'detector': cv.SIFT_create(),  
     'matcher' : cv.DescriptorMatcher_create('BruteForce')},
```

```
]
```

Feature Matching

- Example) **Feature matching comparison** [feature_matching.py]

```
# Detect feature points
keypoints1 = features[f_select]['detector'].detect(img1)
keypoints2 = features[f_select]['detector'].detect(img2)

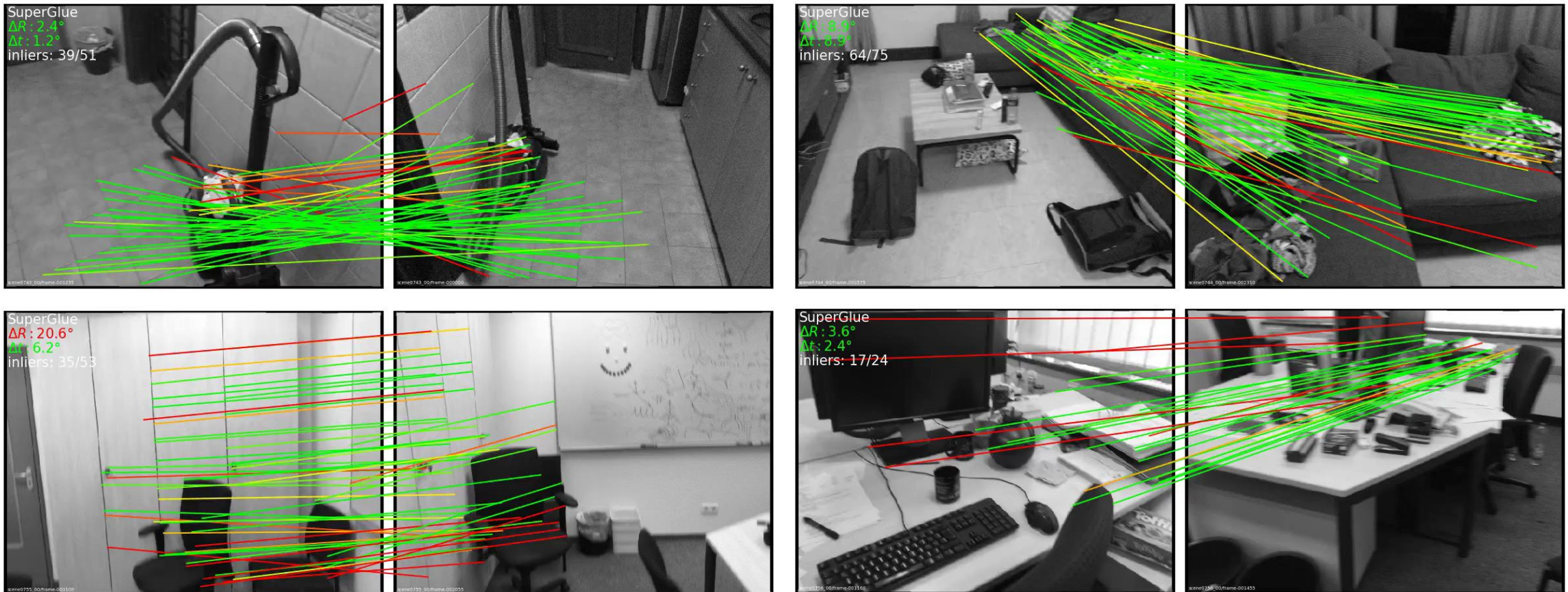
if features[f_select]['matcher'] is not None:
    # Extract feature descriptors
    keypoints1, descriptors1 = features[f_select]['detector'].compute(img1, keypoints1)
    keypoints2, descriptors2 = features[f_select]['detector'].compute(img2, keypoints2)

    # Match the feature descriptors
    match = features[f_select]['matcher'].match(descriptors1, descriptors2)

# Show the matched image
if features[f_select]['matcher'] is not None:
    img_merged = cv.drawMatches(img1, keypoints1, img2, keypoints2, match, None)
else:
    img1_keypts = cv.drawKeypoints(img1, keypoints1, None)
    img2_keypts = cv.drawKeypoints(img2, keypoints2, None)
    img_merged = np.hstack((img1_keypts, img2_keypts))
info = features[f_select]['name'] + ...
cv.putText(img_merged, info, (5, 15), cv.FONT_HERSHEY_PLAIN, 1, (0, 0, 255))
cv.imshow('Feature Matching', img_merged)
```

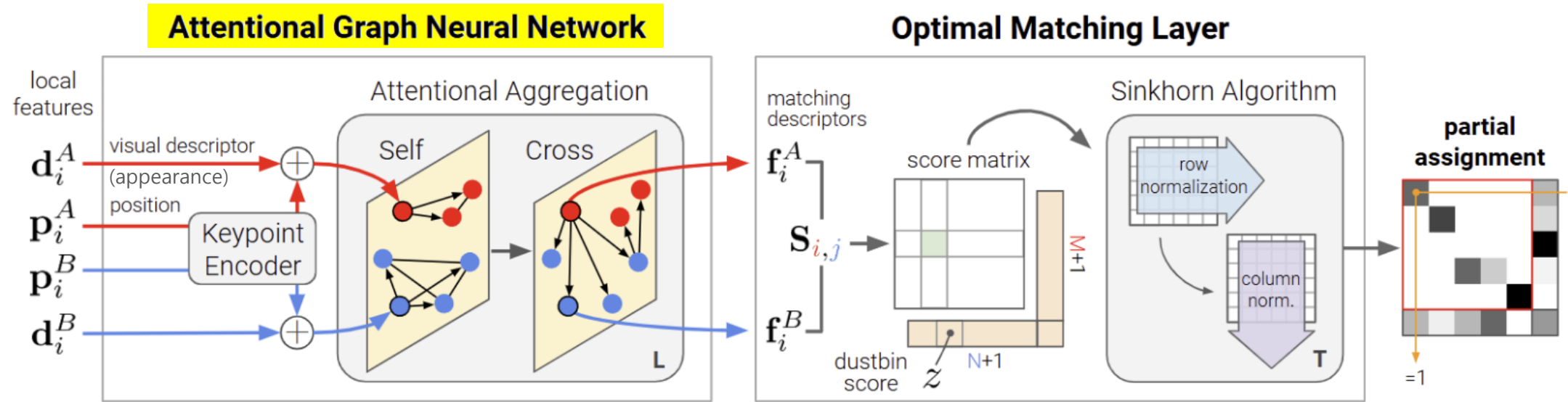

Feature Matching) SuperGlue (2020) → LightGlue (2023)

- Key idea: **Encoding feature points and descriptors using a graph neural network with attention**
 - For extreme wide-baseline image pairs in real-time on GPU
 - e.g. Relative pose estimation on the [ScanNet dataset](#) (more **correct matches** and less **mismatches**)

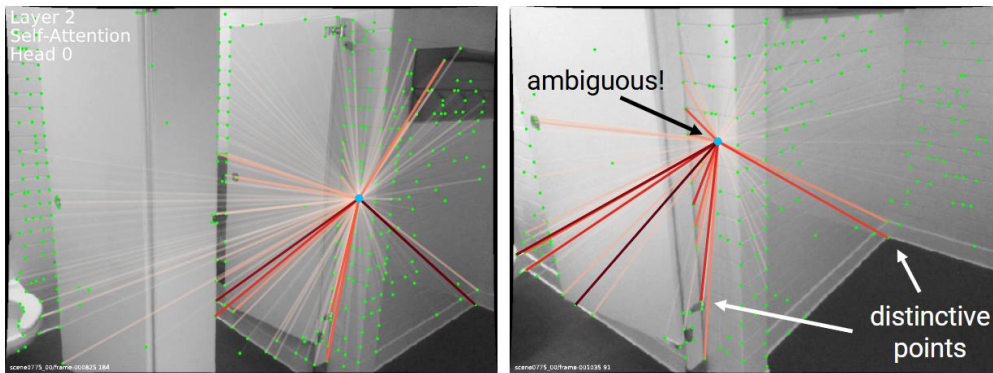


Feature Matching) SuperGlue (2020)

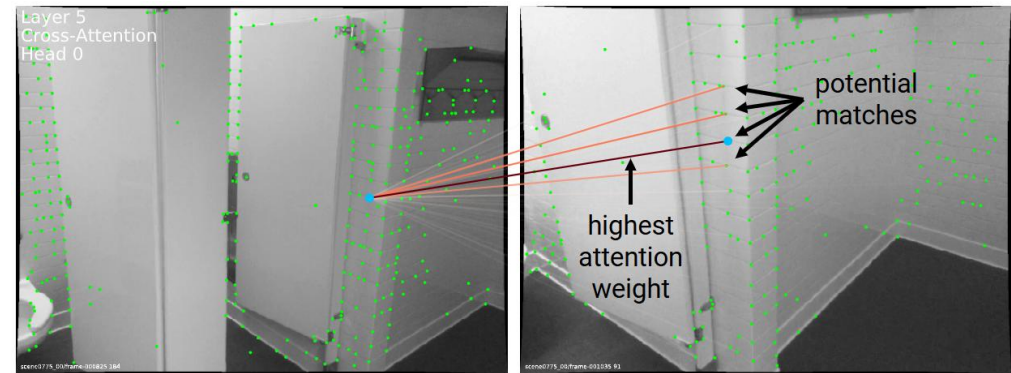
- Key idea: **Encoding feature points and descriptors using a graph neural network with attention**



Self-attention: Intra-image information flow

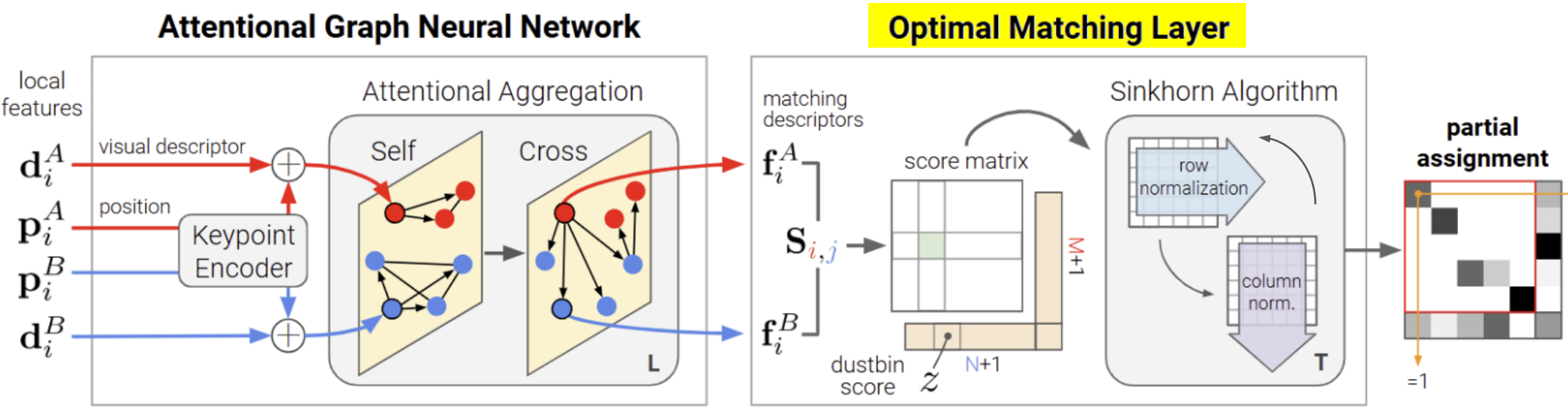


Cross-attention: Inter-image communication



Feature Matching) SuperGlue (2020)

- Key idea: **Encoding feature points and descriptors using a graph neural network with attention**



- Matching score $S_{i,j}$: Inner product of f_i^A and f_j^B
- Sinkhorn algorithm: Differentiable & soft [Hungarian algorithm](#)

e.g. Optimal assignment problem

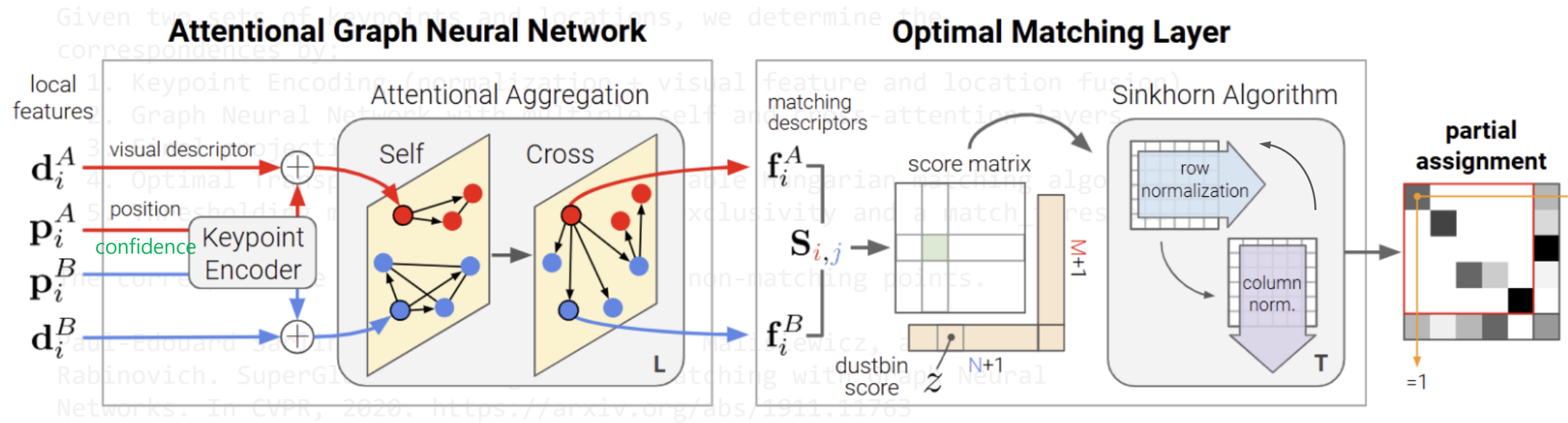
Task \ Worker	Clean bathroom	Sweep floors	Wash windows
Alice	\$8	\$4	\$7
Bob	\$5	\$2	\$3
Carol	\$9	\$4	\$8

Feature Matching) SuperGlue (2020)

- Example) **SuperGlue** implementation [models/superglue.py] @ [Github \(authors\)](#)

```
class SuperGlue(nn.Module):
```

```
    """SuperGlue feature matching middle-end
```



```
    """
```

```
default_config = {
    'descriptor_dim': 256,
    'weights': 'indoor',
    'keypoint_encoder': [32, 64, 128, 256], # 4-layer CNN: [3, 32, 128, 256, 256] channels (kernel size = 1)
    'GNN_layers': ['self', 'cross'] * 9, # 18-layer attention network + 1-layer CNN (final projection)
    'sinkhorn_iterations': 100, # T = 100
    'match_threshold': 0.2,
}
```


Feature Extraction and Matching

- Example) Feature matching comparison: [Image Matching WebUI](#)

SpacesRealcat/image-matching-webuilike86Running

AppFilesCommunity

Image MatchingStructure from Motion(under-dev)




Image Matching WebUI

This Space demonstrates [Image Matching WebUI](#) by vincent qin. Feel free to play with it, or duplicate to run image matching without a queue!

For more details about supported local features and matchers, please refer to <https://github.com/Vincentqyw/image-matching-webui>

All algorithms run on CPU for inference, causing slow speeds and high latency. For faster inference, please download the [source code](#) for local deployment.

Your feedback is valuable to me. Please do not hesitate to report any bugs [here](#).

Matching Model

superpoint+superglue

Image Source

☒ upload☐ webcam☐ clipboard

Image 0


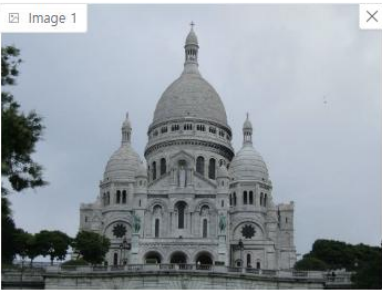


Image 1





Reset

Run Match

Advanced Setting

Open for More: Examples

Examples (click one of the images below to Run Match). Thx: WxBS

Image 0	Image 1	Match threshold	Max features	Keypoint threshold	Matching Mod
		0.1	2000	0.01	disk

Open for More: Keypoints

Open for More: Raw Matches (Green for good matches, Red for bad)

Open for More: Ransac Matches (Green for good matches, Red for bad)

Ransac Matches

Ransac matched keypoints

#Matches: 155





Image 1 - Ransac matched keypoints



Open for More: Matches Statistics


Open for More: Warped Image

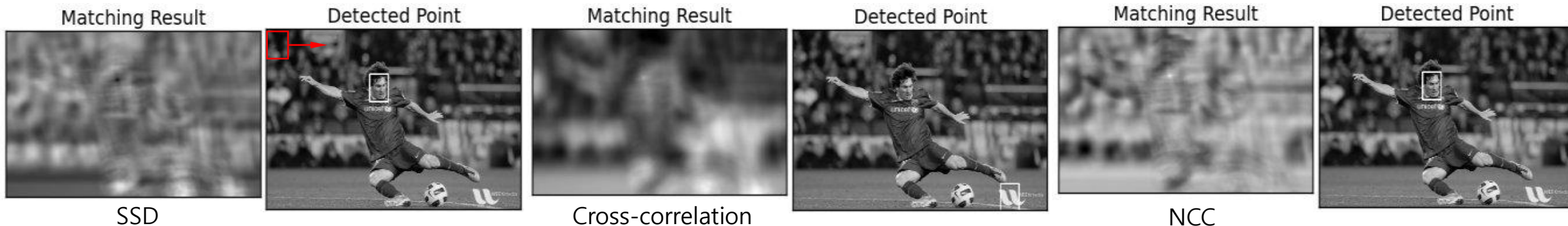
Wrapped Pair

Image 0

Image 1 - warped

Feature **Template** Matching) For Image Patches

- **Raw image patches** (or **histogram**) can be used as descriptors.
 - Distance measures
 - SAD ([sum of absolute difference](#)): $l_1(\mathbf{d}, \mathbf{d}') = \|\mathbf{d} - \mathbf{d}'\|_1 = \sum_i |d_i - d'_i|$
 - SSD (sum of squared difference): $l_2^2(\mathbf{d}, \mathbf{d}') = \|\mathbf{d} - \mathbf{d}'\|_2^2 = \sum_i (d_i - d'_i)^2$
 - [Cross-correlation](#): $s_{cc}(\mathbf{d}, \mathbf{d}') = \mathbf{d} \cdot \mathbf{d}' = \sum_i d_i d'_i$
 - NCC (normalized cross-correlation): $s_{NCC}(\mathbf{d}, \mathbf{d}') = \frac{\mathbf{d} \cdot \mathbf{d}'}{\|\mathbf{d}\| \|\mathbf{d}'\|}$ (\sim cosine similarity)
 - ZNCC (zero-mean normalized cross-correlation): $s_{ZNCC}(\mathbf{d}, \mathbf{d}') = \frac{\bar{\mathbf{d}} \cdot \bar{\mathbf{d}'}}{\|\bar{\mathbf{d}}\| \|\bar{\mathbf{d}'}\|}$ where $\bar{\mathbf{d}} = \mathbf{d} - E(\mathbf{d})$ and $\bar{\mathbf{d}'} = \mathbf{d}' - E(\mathbf{d}')$
 - Matching algorithms
 - Sliding **window** (\mathbf{d} : )



Feature Tracking) Lukas-Kanade Optical Flow (1981)

- Key idea: **Finding movement of a patch whose pixel values are same**

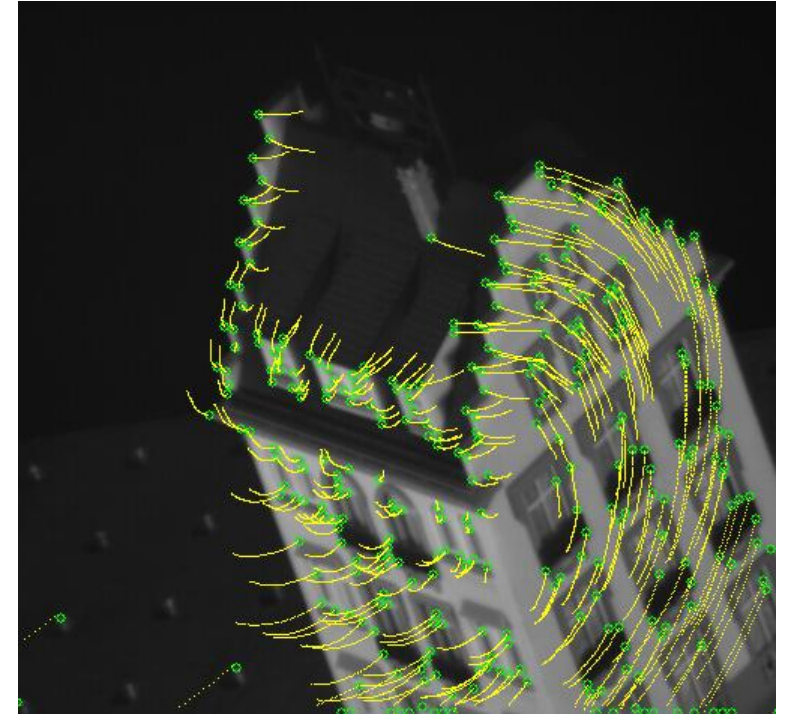
- Brightness constancy constraint: $I(x, y, t) = I(x + \Delta_x, y + \Delta_y, t + \Delta_t)$

$$I_x \frac{\Delta_x}{\Delta_t} + I_y \frac{\Delta_y}{\Delta_t} + I_t = 0 \quad \leftarrow \quad I(x + \Delta_x, y + \Delta_y, t + \Delta_t) \approx I(x, y, t) + I_x \Delta_x + I_y \Delta_y + I_t \Delta_t$$

$$A\mathbf{x} = \mathbf{b} \quad \text{where} \quad A = [I_x \quad I_y], \quad \mathbf{x} = [\Delta_x, \Delta_y]^T, \quad \text{and} \quad \mathbf{b} = [-I_t] \quad (\Delta_t = 1)$$

$$\therefore \mathbf{x} = A^+ \mathbf{b}$$

- Combination: **KLT tracker**
 - GFTT (a.k.a. Shi-Tomasi) detector + **Lukas-Kanade** optical flow
- Advantages and disadvantages (feature tracking vs. matching)
 - (+) No descriptor required (\rightarrow fast and compact)
 - (-) Continuous feature tracking causes drift errors.
 - (-) Not working in wide-baseline cases
 - (+) Able to control matching range



Feature Tracking) Lukas-Kanade Optical Flow (1981)

- Example) **Feature matching comparison** [feature_tracking_klt.py]



Feature Tracking) Lukas-Kanade Optical Flow (1981)

- Example) **Feature matching comparison** [feature_tracking_klt.py]

```
# Open a video and get an initial image
video = cv.VideoCapture(video_file)
assert video.isOpened()

_, gray_prev = video.read()
if gray_prev.ndim >= 3 and gray_prev.shape[2] > 1:
    gray_prev = cv.cvtColor(gray_prev, cv.COLOR_BGR2GRAY)

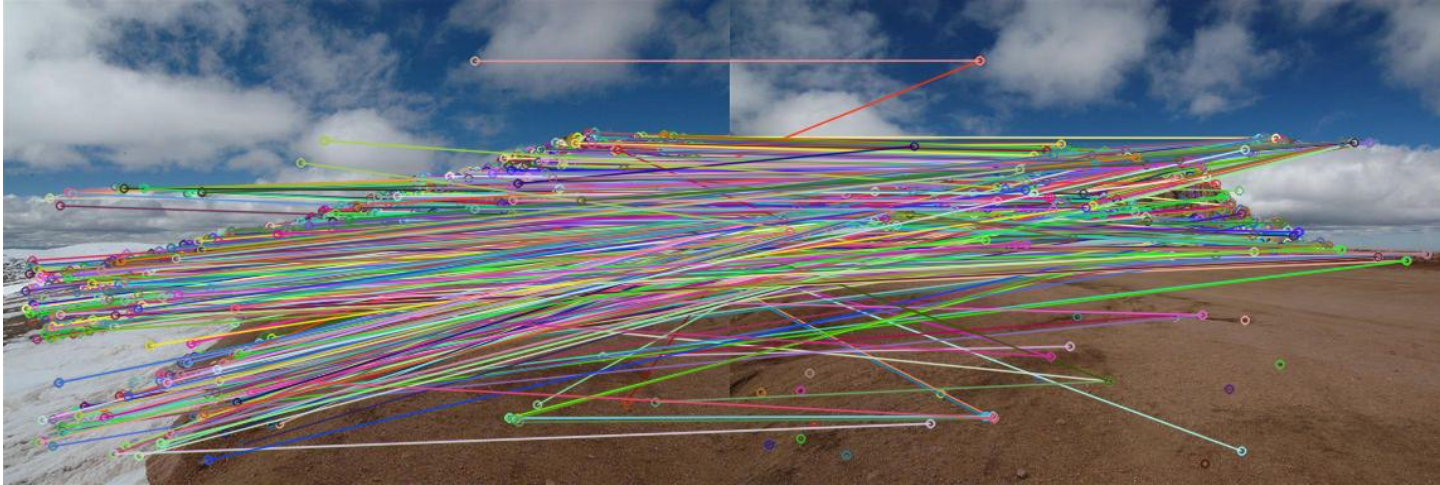
# Run the KLT feature tracker
while True:
    # Grab an image from the video
    valid, img = video.read()
    if not valid:
        break
    if img.ndim >= 3 and img.shape[2] > 1:
        gray = cv.cvtColor(img, cv.COLOR_BGR2GRAY)
    else:
        gray = img.copy()

    # Extract optical flow
    pts_prev = cv.goodFeaturesToTrack(gray_prev, 2000, 0.01, 10)
    pts, status, error = cv.calcOpticalFlowPyrLK(gray_prev, gray, pts_prev, None)
    gray_prev = gray

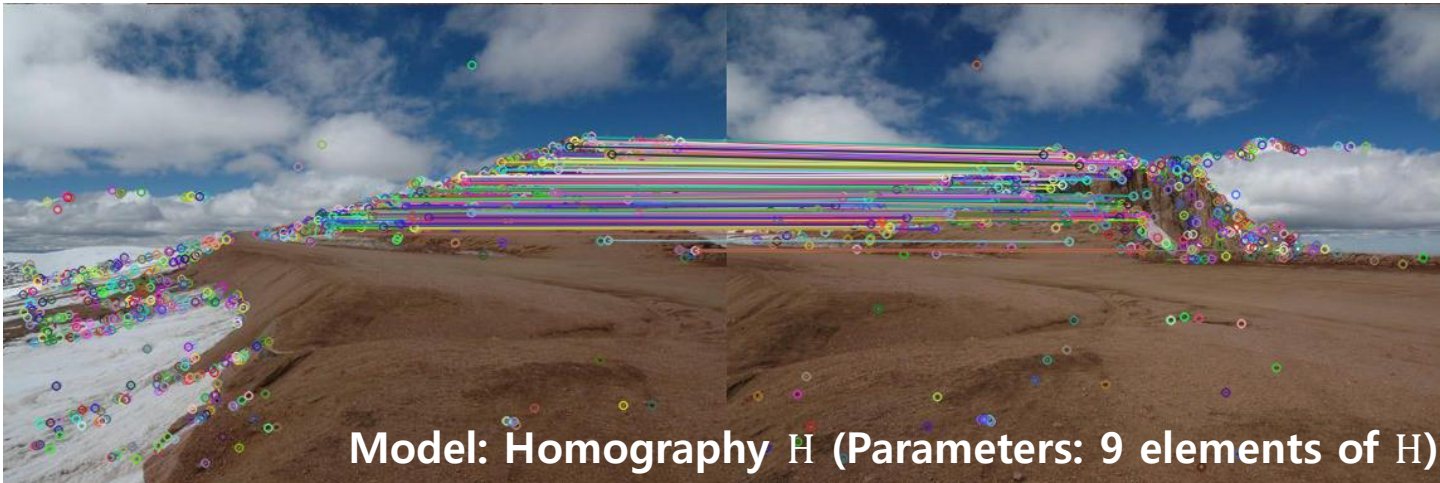
    # Show the optical flow on the image
    if img.ndim < 3 or img.shape[2] < 3:
```


Why Outliers?

Putative feature matches (inliers + outliers)



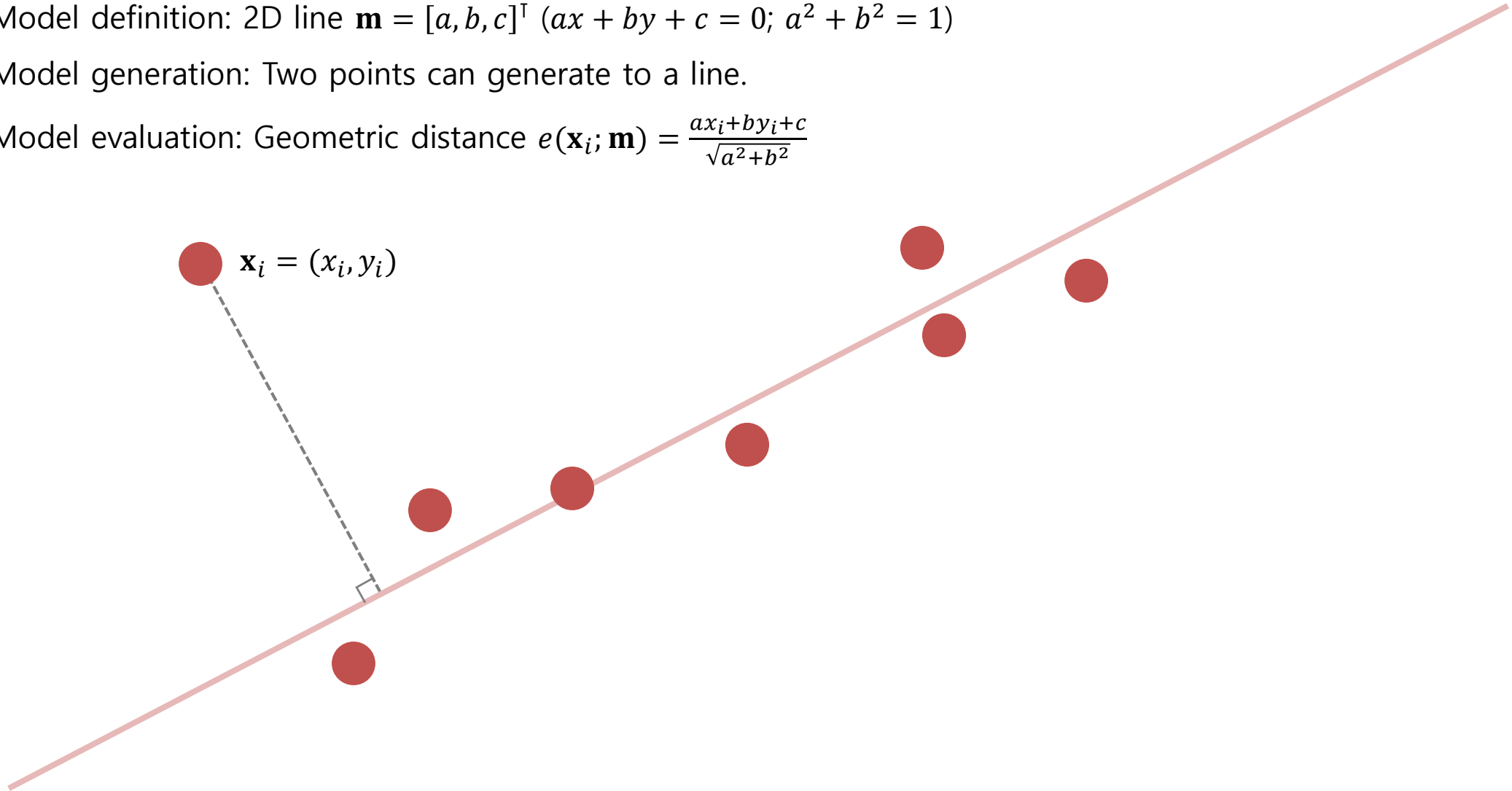
After applying RANSAC (inliers)



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- Example) **Line fitting with [RANSAC](#)**

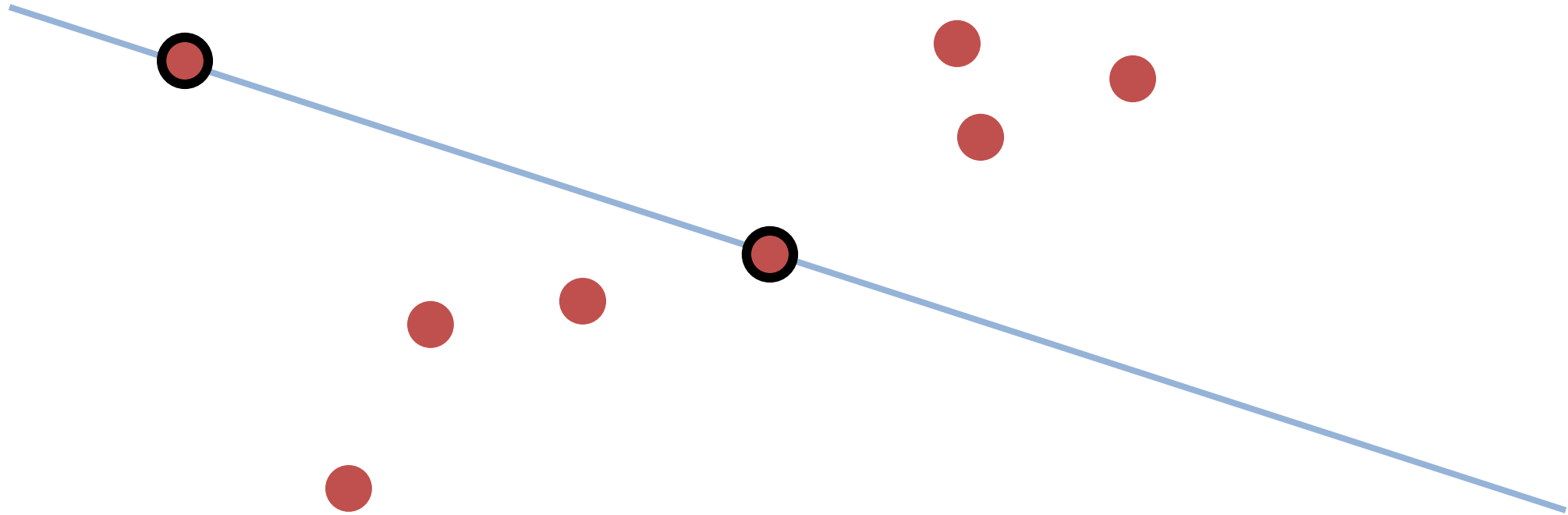
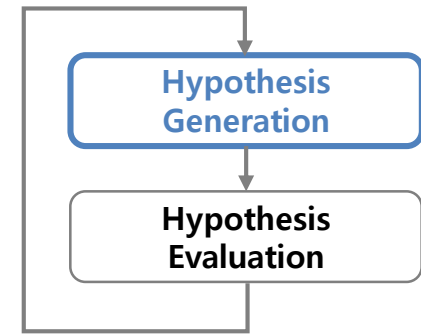
- Model definition: 2D line $\mathbf{m} = [a, b, c]^T$ ($ax + by + c = 0$; $a^2 + b^2 = 1$)
- Model generation: Two points can generate to a line.
- Model evaluation: Geometric distance $e(\mathbf{x}_i; \mathbf{m}) = \frac{ax_i + by_i + c}{\sqrt{a^2 + b^2}}$



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- RANSAC

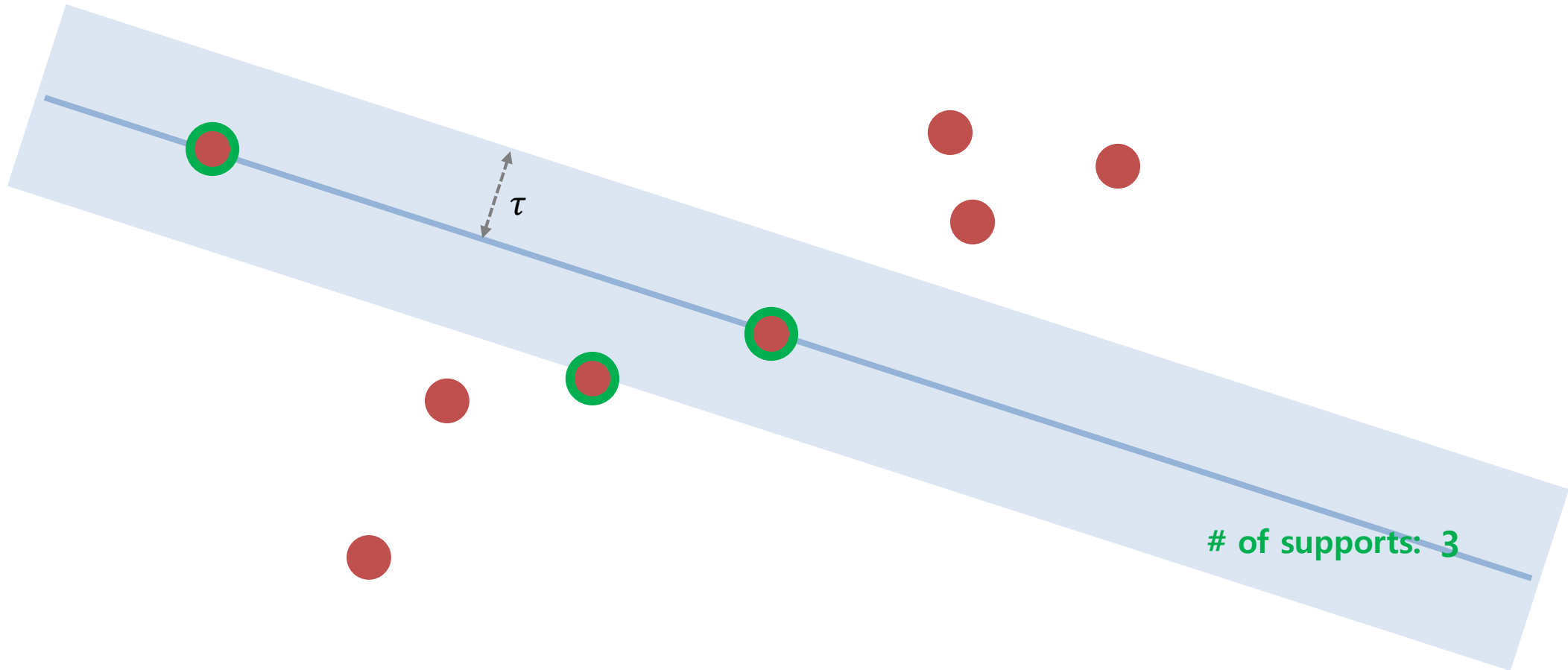
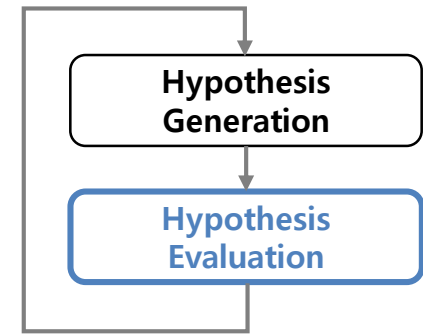
- Iterations of 1) hypothesis generation and 2) its evaluation (~ trial and error)
 - e.g. In this example, an hypothesis is a 2D line from two points.



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- RANSAC

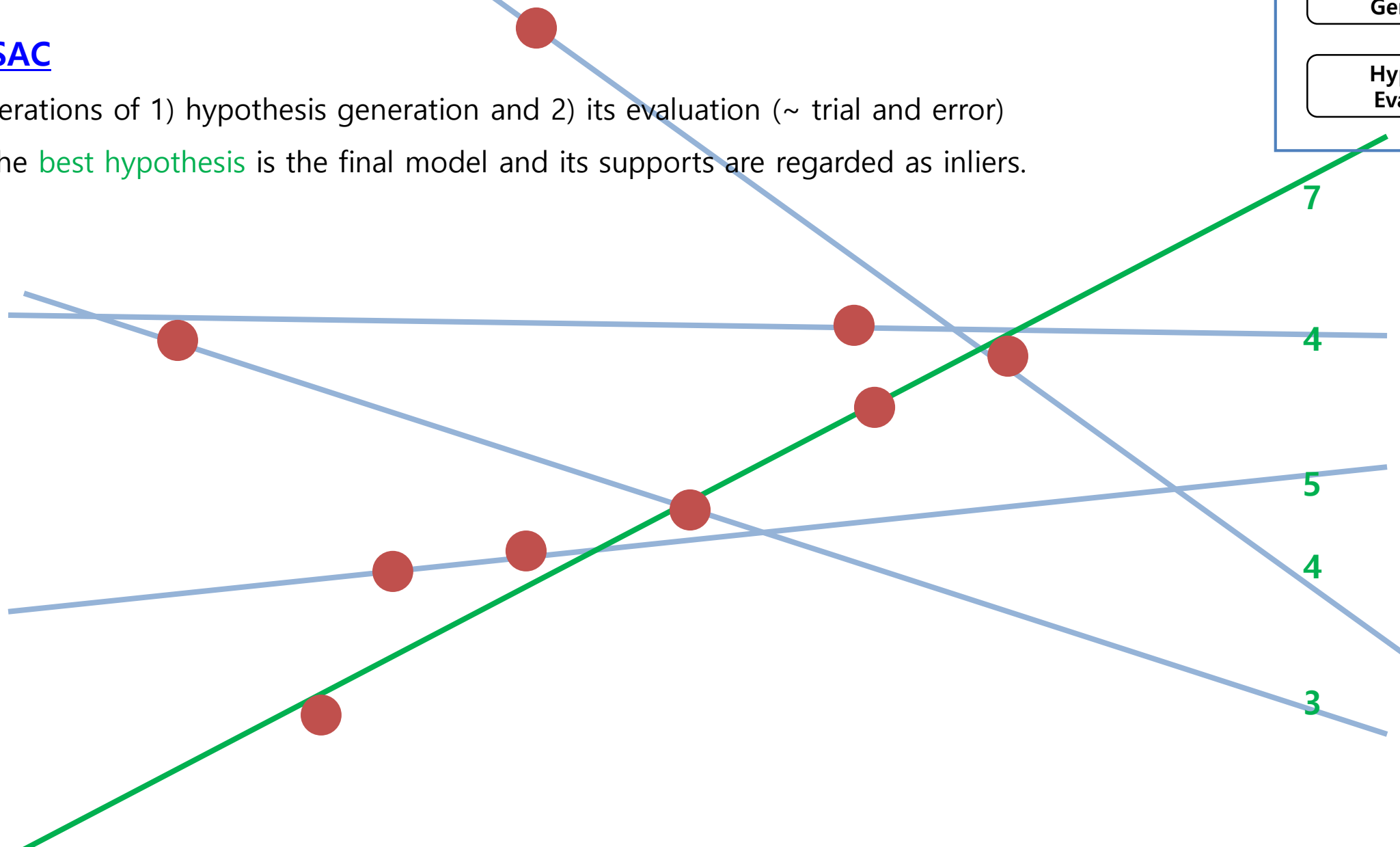
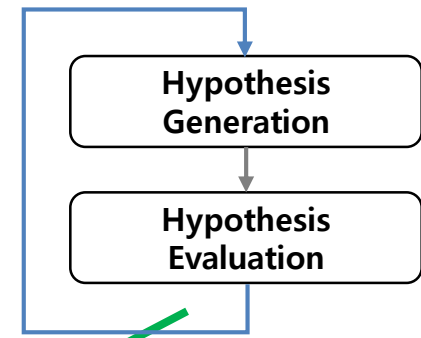
- Iterations of 1) hypothesis generation and 2) its evaluation (~ trial and error)
 - e.g. In this example, the 2D line is supported by points within the given threshold τ .



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- RANSAC

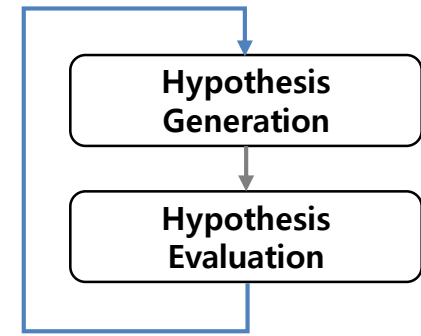
- Iterations of 1) hypothesis generation and 2) its evaluation (~ trial and error)
- The **best hypothesis** is the final model and its supports are regarded as inliers.



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

▪ RANSAC

- Iterations of 1) hypothesis generation and 2) its evaluation (~ trial and error)
- **Q) How many iterations t are required?**
 - Parameters and assumptions
 - s : Success probability (confidence level)
 - d : The number of samples for model generation
 - γ : Inlier ratio
 - Success criteria: **Selecting d samples from inliers within t iterations**
 - Failure sampling: $1 - \gamma^d$
 - Failure probability: $1 - s = (1 - \gamma^d)^t$
 - **The minimum number of iteration for success: $t = \frac{\log(1-s)}{\log(1-\gamma^d)}$**



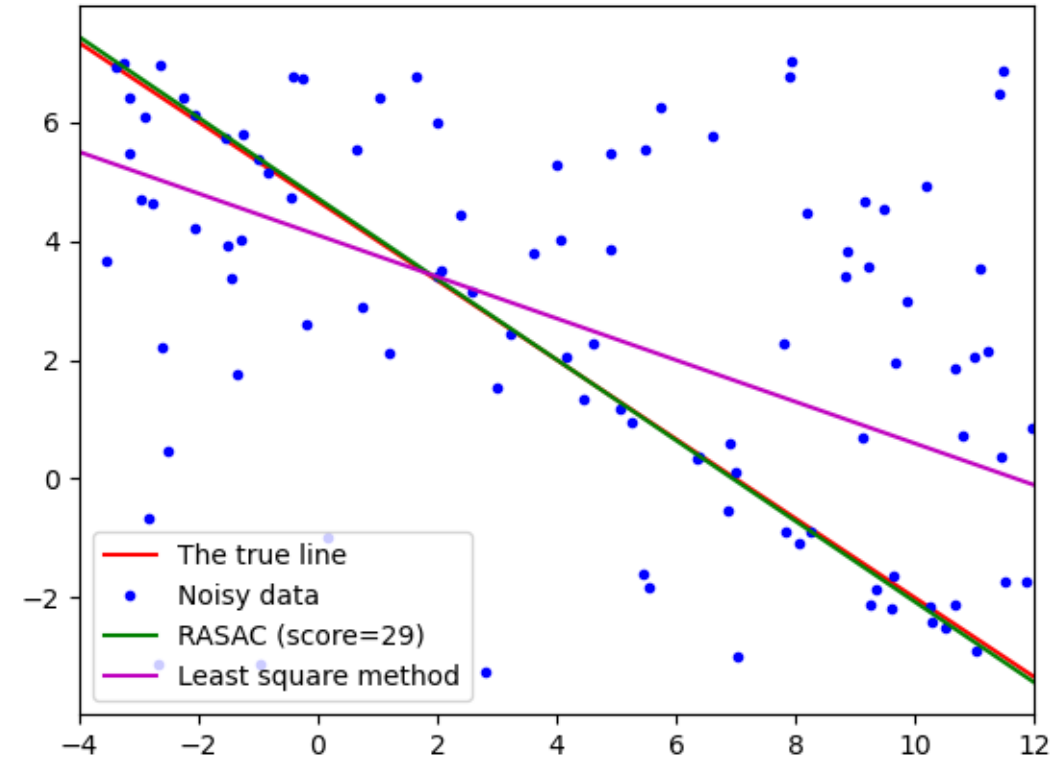
Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

▪ Example) **Line fitting with RANSAC** [line_fitting_ransac.py]

- Model definition: 2D line $\mathbf{m} = [a, b, c]^T$ ($ax + by + c = 0$; $a^2 + b^2 = 1$)
- Model generation: Two points can generate to a line.
- Model evaluation: Geometric distance $e(\mathbf{x}_i; \mathbf{m}) = \frac{ax_i + by_i + c}{\sqrt{a^2 + b^2}}$

```
def generate_line(pts):  
    # Line model:  $y = ax + b$   
    a = (pts[1][1] - pts[0][1]) / (pts[1][0] - pts[0][0])  
    b = pts[0][1] - a * pts[0][0]  
  
    # Line model:  $ax + by + c = 0$  ( $a^2 + b^2 = 1$ )  
    line = np.array([a, -1, b])  
    return line / np.linalg.norm(line[:2])
```

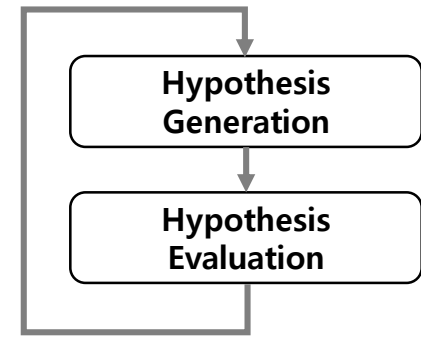
```
def evaluate_line(line, p):  
    a, b, c = line  
    x, y = p  
    return np.fabs(a*x + b*y + c)
```



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- Example) **Line fitting with RANSAC** [line_fitting_ransac.py]

```
def generate_line(pts):  
    ...  
  
def evaluate_line(line, p):  
    ...  
  
def fit_line_ransac(data, n_sample, ransac_trial, ransac_threshold):  
    best_score = -1  
    best_model = None  
    for _ in range(ransac_trial):  
        # Step 1: Hypothesis generation  
        sample = random.choices(data, k=n_sample)  
        model = generate_line(sample)  
  
        # Step 2: Hypothesis evaluation  
        score = 0  
        for p in data:  
            error = evaluate_line(model, p)  
            if error < ransac_threshold:  
                score += 1  
        if score > best_score:  
            best_score = score  
            best_model = model  
  
    return best_model, best_score
```



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- Example) **Line fitting with RANSAC** [line_fitting_ransac.py]

```
if __name__ == '__main__':
    true_line = np.array([2, 3, -14]) / np.sqrt(2*2 + 3*3) # The line model: a*x + b*y + c = 0 (a^2 + b^2 = 1)
    data_range = np.array([-4, 12])
    data_num = 100
    noise_std = 0.2
    outlier_ratio = 0.7

    # Generate noisy points with outliers
    line2y = lambda line, x: (line[0] * x + line[2]) / -line[1] # ax + by + c = 0 -> y = (ax + c) / -b
    y_range = sorted(line2y(true_line, data_range))
    data = []
    for _ in range(data_num):
        x = np.random.uniform(*data_range)
        if np.random.rand() < outlier_ratio:
            y = np.random.uniform(*y_range) # Generate an outlier
        else:
            y = line2y(true_line, x) # Generate an inlier
            x += np.random.normal(scale=noise_std) # Add Gaussian noise
            y += np.random.normal(scale=noise_std)
        data.append((x, y))
    data = np.array(data)

    # Estimate a line using RANSAC
    best_line, best_score = fit_line_ransac(data, 2, 100, 0.3) # log(1 - 0.999) / log(1 - 0.3^2) = 73

    # Estimate a line using OpenCV (for reference)
```

$$t > \frac{\log(1-s)}{\log(1-\gamma^d)} = \frac{\log(1-0.999)}{\log(1-0.3^2)} = 73$$

Review) Planar Homography

- Example) **Planar image stitching** [image_stitching.py]

```
# Load two images
img1 = cv.imread('../data/hill01.jpg')
img2 = cv.imread('../data/hill02.jpg')

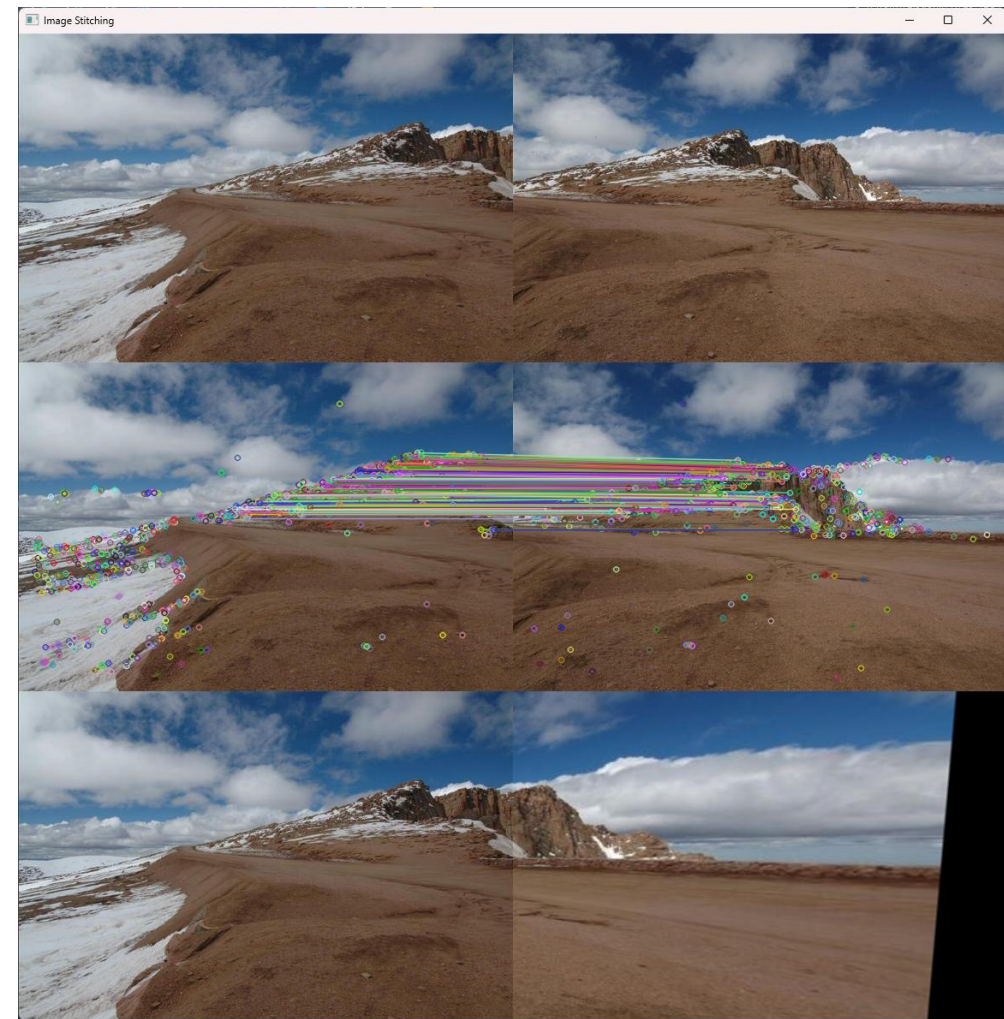
# Retrieve matching points
fdetector = cv.BRISK_create()
keypoints1, descriptors1 = fdetector.detectAndCompute(img1, None)
keypoints2, descriptors2 = fdetector.detectAndCompute(img2, None)

fmatcher = cv.DescriptorMatcher_create('BruteForce-Hamming')
match = fmatcher.match(descriptors1, descriptors2)

# Calculate planar homography and merge them
pts1, pts2 = [], []
for i in range(len(match)):
    pts1.append(keypoints1[match[i].queryIdx].pt)
    pts2.append(keypoints2[match[i].trainIdx].pt)
pts1 = np.array(pts1, dtype=np.float32)
pts2 = np.array(pts2, dtype=np.float32)

H, inlier_mask = cv.findHomography(pts2, pts1, cv.RANSAC)
img_merged = cv.warpPerspective(img2, H, (img1.shape[1]*2, img1.shape[0]))
img_merged[:, :img1.shape[1]] = img1 # Copy

# Show the merged image
img_matched = cv.drawMatches(img1, keypoints1, img2, keypoints2, match, None, None, None,
```



Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- Example) **Planar Homography estimation with RANSAC** [image_stitching_implement.py]

```
from homography_estimation_implement import getPerspectiveTransform
from image_warping_implement import warpPerspective2

def findHomography(src, dst, n_sample, ransac_trial, ransac_threshold):
    ...

if __name__ == '__main__':
    # Load two images
    ...

    # Retrieve matching points
    ...

    # Calculate planar homography and merge them
    ...

    H, inlier_mask = findHomography(pts2, pts1, 4, 1000, 2) #  $\log(1 - 0.999) / \log(1 - 0.3^4) = 849$ 
    img_merged = warpPerspective2(img2, H, (img1.shape[1]*2, img1.shape[0]))
    img_merged[:, :img1.shape[1]] = img1 # Copy

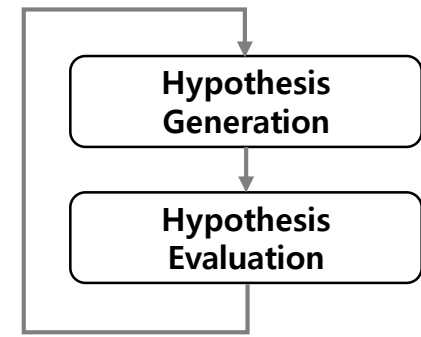
    # Show the merged image
    img_matched = cv.drawMatches(img1, keypoints1, img2, keypoints2, match, None, None, None,
                                matchesMask=inlier_mask) # Remove `matchesMask` if you want to show all ...
    merge = np.vstack((np.hstack((img1, img2)), img_matched, img_merged))
    cv.imshow(f'Planar Image Stitching with My RANSAC (score={sum(inlier_mask)}), merge)
    cv.waitKey(0)
```

Outlier Rejection) RANSAC (Random Sample Consensus; 1981)

- Example) **Homography estimation with RANSAC** [image_stitching_implement.py]

```
def evaluate_homography(H, p, q):  
    p2q = H @ np.array([[p[0]], [p[1]], [1]])  
    p2q /= p2q[-1]  
    return np.linalg.norm(p2q[:2].flatten() - q)    Note)  $e(\mathbf{x}, \mathbf{x}') = \|\mathbf{H}\mathbf{x} - \mathbf{x}'\|_2$ 
```

```
def findHomography(src, dst, n_sample, ransac_trial, ransac_threshold):  
    best_score = -1  
    best_model = None  
    for _ in range(ransac_trial):  
        # Step 1: Hypothesis generation  
        sample_idx = random.choices(range(len(src)), k=n_sample)  
        model = getPerspectiveTransform(src[sample_idx], dst[sample_idx])  
  
        # Step 2: Hypothesis evaluation  
        score = 0  
        for (p, q) in zip(src, dst):  
            error = evaluate_homography(model, p, q)  
            if error < ransac_threshold:  
                score += 1  
        if score > best_score:  
            best_score = score  
            best_model = model  
  
    # Generate the best inlier mask  
    best_inlier_mask = np.zeros(len(src), dtype=np.uint8)  
    for idx, (p, q) in enumerate(zip(src, dst)):
```



Outlier Rejection) Least Squares Method vs. RANSAC

- **Least Squares Method**

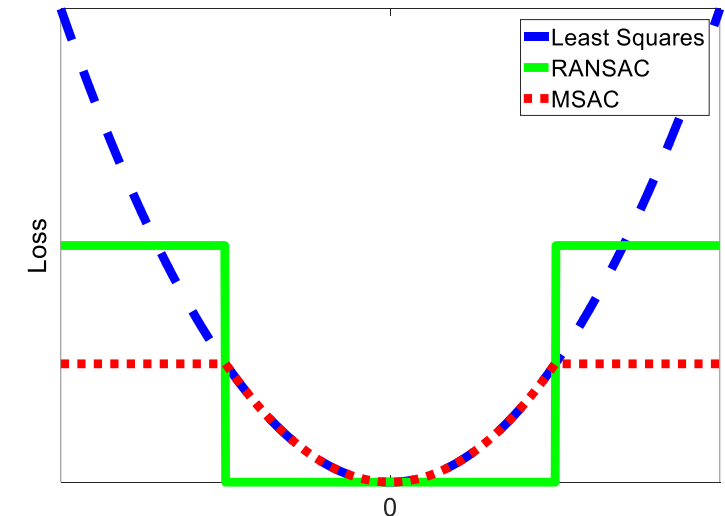
- Find a model while minimizing sum of squared errors, $\operatorname{argmin}_{\mathbf{m}} \sum_i e(\mathbf{x}_i; \mathbf{m})^2$

- **RANSAC**

- Find a model while maximizing the number of supports (\sim inlier candidates)
~ minimizing the number of outlier candidates

- **Q) Why RANSAC was robust to outliers?**

- Problem formulation with a **loss function** ρ : $\operatorname{argmin}_{\mathbf{m}} \sum_i \rho(e(\mathbf{x}_i; \mathbf{m}))$
- Loss function of least squares method: $\rho(x) = x^2$
- Loss function of RANSAC: $\rho(x) = \begin{cases} 0 & \text{if } |x| < \tau \\ 1 & \text{otherwise} \end{cases}$



Outlier Rejection) M-estimator and MSAC

▪ Robust regression

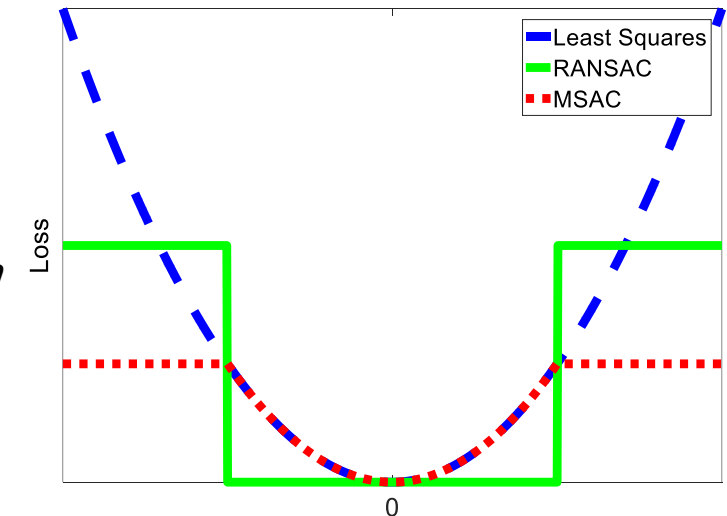
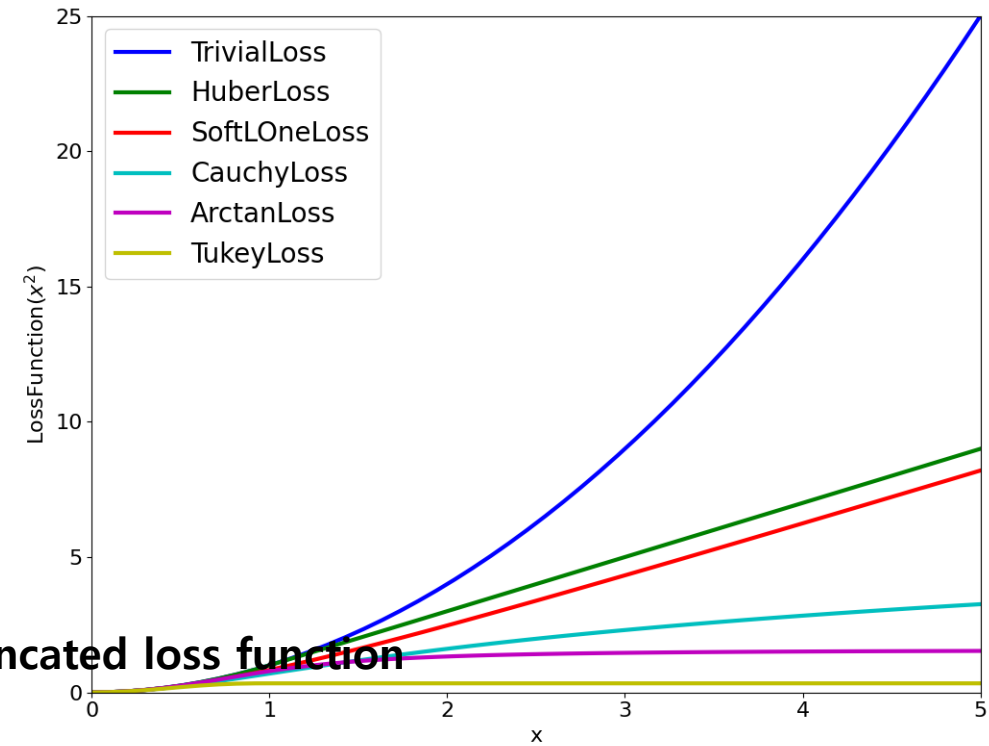
- Problem formulation with a **loss function** ρ : $\operatorname{argmin}_{\mathbf{m}} \sum_i \rho(e(\mathbf{x}_i; \mathbf{m}))$
- Loss function of least squares method: $\rho(x) = x^2$
- Loss function of RANSAC: $\rho(x) = \begin{cases} 0 & \text{if } |x| < \tau \\ 1 & \text{otherwise} \end{cases}$

▪ M-estimator (~ weighted least squares)

- Find a model while minimizing sum of (squared) errors **with a truncated loss function**
- Huber loss function: $\rho(x) = \begin{cases} x^2 & \text{if } |x| < \tau \\ \tau(2|x| - \tau) & \text{otherwise} \end{cases}$
- Cauchy loss function: $\rho(x) = \ln(1 + \frac{x^2}{\tau^2})$

▪ MSAC (RANSAC implementation in OpenCV)

- Consider inliers *with their quality* and regard the outliers *equally with limitation*
- Loss function of MSAC: $\rho(x) = \begin{cases} x^2 & \text{if } |x| < \tau \\ \tau^2 & \text{otherwise} \end{cases}$



Review) Solving Nonlinear Equation using Nonlinear Optimization

- **Example) Line fitting from more than two points such as (1, 4), (4, 2), and (7, 1), using [SciPy](#)**
 - Unknown: Line parameters a , b , and c (line representation: $ax + by + c = 0$)
 - Cost function: $f(a, b, c) = \sum_i \left(\frac{ax_i + by_i + c}{\sqrt{a^2 + b^2}} \right)^2$
 - Optimizer: [Gauss-Newton method](#) (least squares)

```
import numpy as np
from scipy.optimize import least_squares

def geometric_error(line, pts):
    a, b, c = line
    err = [(a*x + b*y + c) / np.sqrt(a*a + b*b) for (x, y) in pts]
    return err

pts = [(1, 4), (4, 2), (7, 1)]
line_init = [1, 1, 0]
result = least_squares(geometric_error, line_init, args=(pts,))
line = result['x'] / -result['x'][1] # [-0.50372575, -1., 4.34823633]
```


Outlier Rejection) M-estimator

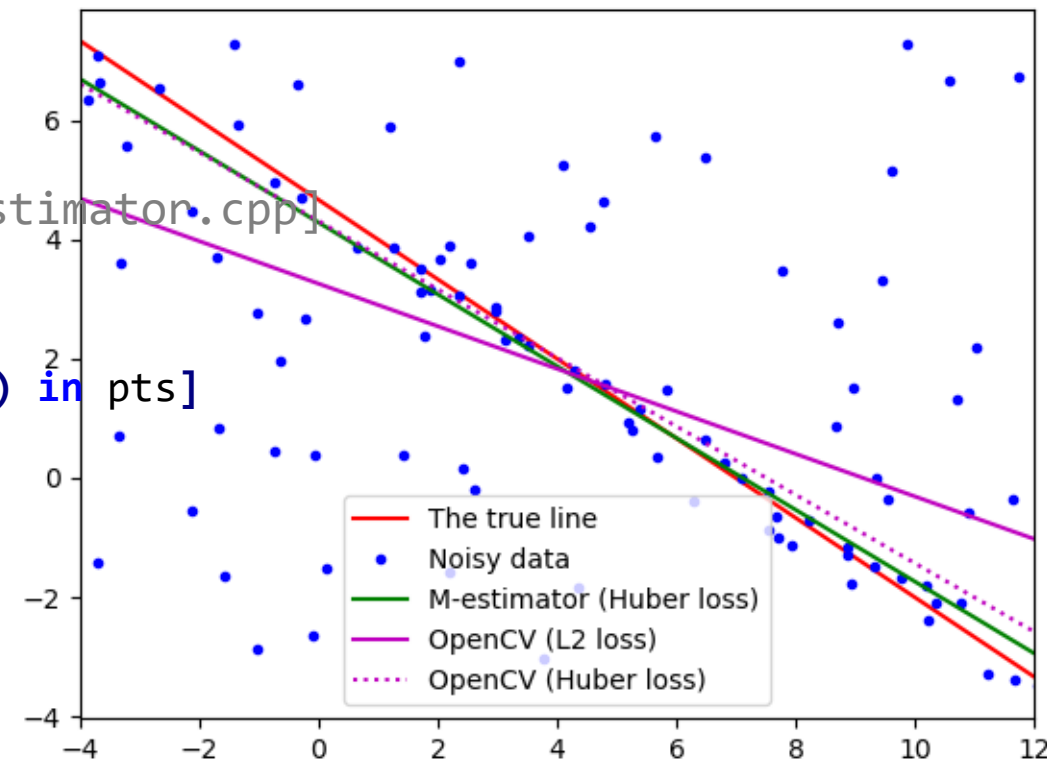
- Example) **Line fitting with M-estimator** [line_fitting_m_estimator.cpp]

```
def geometric_error(line, pts):
    a, b, c = line
    err = [(a*x + b*y + c) / np.sqrt(a*a + b*b) for (x, y) in pts]
    return err

if __name__ == '__main__':
    # Generate noisy points with outliers
    ...

    # Estimate a line using least squares with a robust kernel
    init_line = [1, 1, 0]
    result = least_squares(geometric_error, init_line, args=(data,), loss='huber', f_scale=0.3)
    mest_line = result['x'] / np.linalg.norm(result['x'][:2])

    # Estimate a line using OpenCV (for reference)
    # Note) OpenCV line model: n_y * (x - x_0) = n_x * (y - y_0)
    nnxy = cv.fitLine(data, cv.DIST_L2, 0, 0.01, 0.01).flatten()
    lsqr_line = np.array([nnxy[1], -nnxy[0], -nnxy[1]*nnxy[2] + nnxy[0]*nnxy[3]])
    nnxy = cv.fitLine(data, cv.DIST_HUBER, 0, 0.01, 0.01).flatten()
    huber_line = np.array([nnxy[1], -nnxy[0], -nnxy[1]*nnxy[2] + nnxy[0]*nnxy[3]])
```



Outlier Rejection) Applications (Model Size)

- **Bottom-up approaches (~ voting)**

- ~~Hough transform~~ (not suitable for more than three model parameters)

- **A single datum** votes multiple model candidates.
 - Note) The parameter space is maintained as a multi-dimensional histogram (discretization).
 - Score: The number of hits by data

- **RANSAC family**

- **A sample of data** votes a single model candidate.
 - Score: The number of supports (whose error is within threshold)

- Note) Application examples (**small-size model**): Line fitting, homography estimation, relative pose estimation, PnP

- **Top-down approaches**

- **Optimization with truncated loss functions (e.g. M-estimator)**

- The initial model parameter moves along with the gradient of the given cost function with **whole data**.
 - Score: A cost function with a truncated loss function.

- Note) Application examples (**large-size model**): Camera calibration, visual SLAM, SfM

Summary) Finding Correspondences

▪ Feature Points

- Gradient-based: Harris corner, GFTT corner, SIFT, SURF, ...
- Intensity-based: FAST, ...
- DL-based: LIFT, SuperPoint, ...

▪ Feature Descriptors

- Real-valued: SIFT, SURF, ... (DL-based: LIFT, SuperPoint, ...)
- Binary-valued: BRIEF, ORB, ...

▪ Feature Matching

- Distance measures: Euclidean distance, cosine similarity, Hamming distance, ...
- Matching methods: Brute-force search, ANN search, ...
- DL-based: SuperGlue, ...

▪ Feature Tracking

- KLT tracker = GFTT detector + LK optical flow

▪ Outlier Rejection (**Robust Regression**)

- **RANSAC** → **MSAC** (inlier quality)
- Least squares method → **M-estimator** (robust kernels)

