## Image Features: <br> Local Descriptors

## Local Features

- Detection: Identify the interest points.
- Description: Extract a feature descriptor around each interest point.
- Matching: Determine correspondence between descriptors in two views.

[Source: K. Grauman]


## Invariances


[Source: T. Tuytelaars]

## Invariances


[Source: T. Tuytelaars]

## What If We Just Took Pixels?

- The simplest way is to write down the list of intensities to form a feature vector, and normalize them (i.e., mean 0 , variance 1 ).
- Why normalization?
- But this is very sensitive to even small shifts, rotations and any affine transformation.
region A



## Tones Of Better Options

- SIFT
- PCA-SIFT
- GLOH
- HOG
- SURF
- DAISY
- LBP
- Shape Contexts
- Color Histograms


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## SIFT Descriptor [Lowe 2004]

- SIFT stands for Scale Invariant Feature Transform
- Invented by David Lowe, who also did DoG scale invariant interest points
- Actually in the same paper, which you should read:


## David G. Lowe

Distinctive image features from scale-invariant keypoints
International Journal of Computer Vision, 2004
Paper: http://www.cs.ubc.ca/~lowe/papers/ijcv04.pdf

(a) image gradients

(b) keypoint descriptor

## SIFT Descriptor

(1) Our scale invariant interest point detector gives scale $\rho$ for each keypoint

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor

(2) For each keypoint, we take the Gaussian-blurred image at corresponding scale $\rho$

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor

(3) Compute the gradient magnitude and orientation in neighborhood of each keypoint

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor

(3) Compute the gradient magnitude and orientation in neighborhood of each keypoint
magnitude of gradient:

$$
|\nabla I(x, y)|=\sqrt{\left(\frac{\partial\left(I(x, y) * G_{\rho}\right)}{\partial x}\right)^{2}+\left(\frac{\partial\left(I(x, y) * G_{\rho}\right)}{\partial y}\right)^{2}}
$$

## gradient orientation:

$$
\theta(x, y)=\arctan \left(\frac{\partial I * G_{\rho}}{\partial y} / \frac{\partial I * G_{\rho}}{\partial x}\right)
$$

(in case you forgot;))

## SIFT Descriptor

(9) Compute dominant orientation of each keypoint. How?

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing Dominant Orientation

- Compute a histogram of gradient orientations, each bin covers $10^{\circ}$

$$
16 \times 16
$$

compute histograms of orientations
 by orientation increments of $10^{\circ}$

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing Dominant Orientation

- Compute a histogram of gradient orientations, each bin covers $10^{\circ}$
- Orientations closer to the keypoint center should contribute more

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## SIFT Descriptor: Computing Dominant Orientation

- Compute a histogram of gradient orientations, each bin covers $10^{\circ}$
- Orientations closer to the keypoint center should contribute more
- Orientation giving the peak in the histogram is the keypoint's orientation

[Adopted from: F. Flores-Mangas]


## SIFT Descriptor

(9) Compute dominant orientation

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor

(5) Compute a 128 dimensional descriptor: $4 \times 4$ grid, each cell is a histogram of 8 orientation bins relative to dominant orientation

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing the Feature Vector

- Compute the orientations relative to the dominant orientation


## $16 \times 16$ patch <br> centered in $\left(x_{i}, y_{i}\right)$



[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing the Feature Vector

- Compute the orientations relative to the dominant orientation

$$
\begin{gathered}
16 \times 16 \text { patch } \\
\text { centered in }\left(x_{i}, y_{i}\right)
\end{gathered}
$$


[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing the Feature Vector

- Compute the orientations relative to the dominant orientation
- Form a $4 \times 4$ grid. For each grid cell compute a histogram of orientations for 8 orientation bins spaced apart by $45^{\circ}$

$$
\begin{aligned}
& 16 \times 16 \text { patch } \\
& \text { centered in }\left(x_{i}, y_{i}\right)
\end{aligned}
$$

## SIFT descriptor


compute histogram of orientations this time 8 bins spaced by $45^{\circ}$

[Adopted from: F. Flores-Mangas]

## SIFT Descriptor: Computing the Feature Vector

- Compute the orientations relative to the dominant orientation
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## $16 \times 16$ patch centered in $\left(x_{i}, y_{i}\right)$ <br> SIFT descriptor





## SIFT Descriptor: Computing the Feature Vector

- Compute the orientations relative to the dominant orientation
- Form a $4 \times 4$ grid. For each grid cell compute a histogram of orientations for 8 orientation bins spaced apart by $45^{\circ}$
- Form the 128 dimensional feature vector

$$
\begin{aligned}
& 16 \times 16 \text { patch } \\
& \text { centered in }\left(x_{i}, y_{i}\right)
\end{aligned}
$$

SIFT descriptor


## SIFT Descriptor: Post-processing

- The resulting 128 non-negative values form a raw version of the SIFT descriptor vector.
- To reduce the effects of contrast or gain (additive variations are already removed by the gradient), the 128-D vector is normalized to unit length: $f_{i}=f_{i} /\left\|f_{i}\right\|$


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- Great engineering effort!
- What is SIFT invariant to?


## Properties of SIFT

Invariant to:

- Scale
- Rotation

Partially invariant to:

- Illumination changes (sometimes even day vs. night)
- Camera viewpoint (up to about 60 degrees of out-of-plane rotation)
- Occlusion, clutter (why?)

Also important:

- Fast and efficient - can run in real time
- Lots of code available


## Examples



Figure: Matching in day / night under viewpoint change
[Source: S. Seitz]

## Example



Figure: NASA Mars Rover images with SIFT feature matches
[Source: N. Snavely]

- The dimensionality of SIFT is pretty high, i.e., 128D for each keypoint
- Reduce the dimensionality using linear dimensionality reduction
- In this case, principal component analysis (PCA)
- Use 10D or so descriptor
[Source: R. Urtasun]


## Gradient location-orientation histogram (GLOH)

- Developed by Mikolajczyk and Schmid (2005): variant of SIFT that uses a log-polar binning structure instead of the four quadrants.
- The spatial bins are 11 , and 15 , with eight angular bins (except for the central region), for a total of 17 spatial bins and 16 orientation bins.
- The 272D histogram is then projected onto a 128D descriptor using PCA trained on a large database.

(a) image gradients

(b) keypoint descriptor
[Source: R. Szeliski]


## Other Descriptors

- SURF
- DAISY
- LBP
- HOG
- Shape Contexts
- Color Histograms


## Local Features

- Detection: Identify the interest points.
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[Source: K. Grauman]


# Image Features: Matching the Local Descriptors 

## Matching the Local Descriptors

Once we have extracted keypoints and their descriptors, we want to match the features between pairs of images.

- Ideally a match is a correspondence between a local part of the object on one image to the same local part of the object in another image
- How should we compute a match?


Figure: Images from K. Grauman

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Figure: Images from K. Grauman

## Matching the Local Descriptors

- Simple: Compare them all, compute Euclidean distance

$f_{x} \square 11 \cdots 11$



## Matching the Local Descriptors

- Simple: Compare them all, compute Euclidean distance



## Matching the Local Descriptors

- Find closest match (min distance). How do we know if match is reliable?



## Matching the Local Descriptors

- Find also the second closest match. Match reliable if first distance "much" smallar then carend dictonea



## Matching the Local Descriptors

- Compute the ratio:

$$
\phi_{i}=\frac{\left\|f_{i}-f_{i}^{\prime *}\right\|}{\left\|f_{i}-f_{i}^{\prime * *}\right\|}
$$

where $f_{i}^{\prime *}$ is the closest and $f_{i}^{\prime * *}$ second closest match to $f_{i}$.


## Which Threshold to Use?

- Setting the threshold too high results in too many false positives, i.e., incorrect matches being returned.
- Setting the threshold too low results in too many false negatives, i.e., too many correct matches being missed


Figure: Images from R. Szeliski

## Which Threshold to Use?

- Threshold ratio of nearest to 2 nd nearest descriptor
- Typically: $\phi_{i}<0.8$


Figure: Images from D. Lowe
[Source: K. Grauman]

## Applications of Local Invariant Features

- Wide baseline stereo
- Motion tracking
- Panorama stitching
- Mobile robot navigation
- 3D reconstruction
- Recognition
- Retrieval
[Source: K. Grauman]


## Wide Baseline Stereo


[Source: T. Tuytelaars]

## Recognizing the Same Object



Schmid and Mohr 1997


Rothganger et al. 2003


Sivic and Zisserman, 2003


Lowe 2002
[Source: K. Grauman]

## Motion Tracking



Figure: Images from J. Pilet

## Now What

- Now we know how to extract scale and rotation invariant features
- We even know how to match features across images
- Can we use this to find Waldo in an even more sneaky scenario?


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template

Waldo on the road

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template

He comes closer... We know how to solve this

## Now What

- Now we know how to extract scale and rotation invariant features
- We even know how to match features across images
- Can we use this to find Waldo in an even more sneaky scenario?


Someone takes a (weird) picture of him!

## Find My DVD!

- More interesting: If we have DVD covers (e.g., from Amazon), can we match them to DVDs in real scenes?



## Matching Planar Objects In New Viewpoints

## What Kind of Transformation Happened To My DVD?



## What Kind of Transformation Happened To My DVD?

- Rectangle goes to a parallelogram (almost but not really, but let's believe that for now)



## All 2D Linear Transformations

Linear transformations are combinations of É

- Scale,
- Rotation
- Shear
- Mirror

$$
\left[\begin{array}{l}
x^{\prime} \\
y^{\prime}
\end{array}\right]=\left[\begin{array}{ll}
a & b \\
c & d
\end{array}\right]\left[\begin{array}{l}
x \\
y
\end{array}\right]
$$

[Source: N. Snavely]

## All 2D Linear Transformations

Properties of linear transformations:

- Origin maps to origin
- Lines map to lines


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x^{\prime} \\
y^{\prime}
\end{array}\right]=\left[\begin{array}{ll}
a & b \\
c & d
\end{array}\right]\left[\begin{array}{ll}
e & f \\
g & h
\end{array}\right]\left[\begin{array}{ll}
i & j \\
k & l
\end{array}\right]\left[\begin{array}{l}
x \\
y
\end{array}\right]
$$

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y
\end{array}\right]
$$

## What about the translation?

[Source: N. Snavely]

## All 2D Linear Transformations

Properties of linear transformations:

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x \\
y
\end{array}\right]
$$

What about the translation?
[Source: N. Snavely]

## Affine Transformations

Affine transformations are combinations of:

- Linear transformations, and
- Translations

$$
\left[\begin{array}{l}
x^{\prime} \\
y^{\prime}
\end{array}\right]=\left[\begin{array}{ll}
a & b \\
c & d
\end{array}\right]\left[\begin{array}{l}
x \\
y
\end{array}\right]+\left[\begin{array}{l}
e \\
f
\end{array}\right]
$$

same as:

$$
\left[\begin{array}{l}
x^{\prime} \\
y^{\prime}
\end{array}\right]=\left[\begin{array}{lll}
a & b & e \\
c & d & f
\end{array}\right]\left[\begin{array}{l}
x \\
y \\
1
\end{array}\right]
$$

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Affine transformations are combinations of:

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$$
\left[\begin{array}{l}
x^{\prime} \\
y^{\prime}
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a & b \\
c & d
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x \\
y
\end{array}\right]+\left[\begin{array}{l}
e \\
f
\end{array}\right]
$$

same as:

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\left[\begin{array}{l}
x^{\prime} \\
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a & b & e \\
c & d & f
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x \\
y \\
1
\end{array}\right]
$$

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Affine transformations are combinations of:

- Linear transformations, and
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\left[\begin{array}{l}
x^{\prime} \\
y^{\prime}
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a & b & e \\
c & d & f
\end{array}\right]\left[\begin{array}{l}
x \\
y \\
1
\end{array}\right]
$$

Properties of affine transformations:

- Origin does not necessarily map to origin
- Lines map to lines
- Parallel lines remain parallel
- Ratios are preserved
- Closed under composition
- Rectangles go to parallelograms
[Source: N. Snavely]


## 2D Image Tranformations



- These transformations are a nested set of groups
- Closed under composition and inverse is a member


## What Transformation Happened to My DVD?

- Affine transformation approximates viewpoint changes for roughly planar objects and roughly orthographic cameras (more about these later in class)
- DVD went affine!



## Computing the (Affine) Transformation

Given a set of matches between images I and J

- How can we compute the affine transformation A from I to J?
- Find transform A that best "agrees" with the matches

[Source: N. Snavely]


## Computing the (Affine) Transformation

Given a set of matches between images I and J

- How can we compute the affine transformation A from I to J ?
- Find transform A that best "agrees" with the matches

[Source: N. Snavely]


## Computing the Affine Transformation



- Let $\left(x_{i}, y_{i}\right)$ be a point on the reference (model) image, and $\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$ its match in the test image
- An affine transformation $A$ maps $\left(x_{i}, y_{i}\right)$ to $\left(x_{i}^{\prime}, y_{i}^{\prime}\right)$ :

$$
\left[\begin{array}{l}
x_{i}^{\prime} \\
y_{i}^{\prime}
\end{array}\right]=\left[\begin{array}{lll}
a & b & e \\
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x_{i} \\
y_{i} \\
1
\end{array}\right]
$$

- We can rewrite this into a simple linear system:

$$
\left[\begin{array}{cccccc}
x_{i} & y_{i} & 0 & 0 & 1 & 0 \\
0 & 0 & x_{i} & y_{i} & 0 & 1
\end{array}\right]\left[\begin{array}{l}
a \\
b \\
c \\
d \\
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f
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## Computing the Affine Transformation

- But we have many matches:

- For each match we have two more equations
- How many matches do we need to compute A?
- 6 parameters $\rightarrow 3$ matches
- But the more, the better (more reliable)
- How do we compute A?


## Computing the Affine Transformation



- If we have 3 matches, then computing $A$ is really easy:

$$
\mathbf{a}=\mathbf{P}^{-1} \mathbf{P}^{\prime}
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\min _{a, b, \cdots, f}\left\|\mathbf{P a}-\mathbf{P}^{\prime}\right\|_{2}^{2}
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- Which has a closed form solution:

$$
\mathbf{a}=\left(\mathbf{P}^{T} \mathbf{P}\right)^{-1} \mathbf{P}^{T} \mathbf{P}^{\prime}
$$

## Image Alignment Algorithm: Affine Case

Given images $/$ and $J$
(1) Compute image features for $I$ and $J$
(2) Match features between $I$ and $J$
(3) Compute affine transformation $A$ between $I$ and $J$ using least squares on the set of matches

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Is there a problem with this?
[Source: N. Snavely]

## Robustness


[Source: N. Snavely]

## Simple Case

(This example is unrelated to the object matching example, but it nicely shows how to robustify estimation)

- Let's consider a simpler example ... Fit a line to the points below!


Problem: Fit a line to these datapoints


Least squares fit

## Simple Case

(This example is unrelated to the object matching example, but it nicely shows how to robustify estimation)

- Let's consider a simpler example ... Fit a line to the points below!


Problem: Fit a line to these datapoints


Least squares fit

- How can we fix this?
[Source: N. Snavely]


## Simple Idea: RANSAC

- Take the minimal number of points to compute what we want. In the line example, two points (in our affine example, three matches)


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- Count the number of all points that "agree" with the line: We call the agreeing points inliers
- "Agree" $=$ within a small distance of the line
- Repeat this many times, remember the number of inliers for each trial
- Among several trials, select the one with the largest number of inliers

This procedure is called RAndom SAmple Consensus

## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points

[Source: R. Raguram]

## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points
(2) Hypothesize a model

[Source: R. Raguram]

## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points
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(3) Compute error function

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## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points
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(9) Select points consistent with model

[Source: R. Raguram]

## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points
(2) Hypothesize a model
(3) Compute error function
(9) Select points consistent with model
(6) Repeat hypothesize and verify loop

[Source: R. Raguram]

## RANSAC for Line Fitting Example

(1) Randomly select minimal subset of points
(2) Hypothesize a model
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4 Select points consistent with model
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[Source: R. Raguram]

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(2) Hypothesize a model
(3) Compute error function
(4) Select points consistent with model
(5) Repeat hypothesize and verify loop
(6) Choose model with
 largest set of inliers
[Source: R. Raguram]

## Translations


[Source: N. Snavely]

## RAndom SAmple Consensus


[Source: N. Snavely]

## RAndom SAmple Consensus


[Source: N. Snavely]

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## RANSAC

- All the inliers will agree with each other on the translation vector; the (hopefully small) number of outliers will (hopefully) disagree with each other


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## RANSAC

- All the inliers will agree with each other on the translation vector; the (hopefully small) number of outliers will (hopefully) disagree with each other
- RANSAC only has guarantees if there are $<50 \%$ outliers
- "All good matches are alike; every bad match is bad in its own way." [Tolstoy via Alyosha Efros]
[Source: N. Snavely]


## Affine Transformation

How?

## Affine Transformation

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- Find matches across images $I$ and $J$. This gives us points $X_{I}$ in image $I$ and $X_{J}$ in $J$, where we know that the point $X_{l}^{k}$ is a match with $X_{J}^{k}$
- Iterate:
- Choose 3 pairs of matches randomly
- Compute the affine transformation
- Project all matched points $X_{I}$ from $I$ to $J$ via the computed transformation. This gives us $\hat{X}_{l}$
- Find how many matches are inliers, i.e., $\left\|\hat{X}_{I}^{k}-X_{J}^{k}\right\|<$ thresh.
- Choose the transformation with the most inliers


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- Suppose there are $20 \%$ outliers, and we want to find the correct answer with 99\% probability
- How many rounds do we need?
[Source: R. Urtasun]


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- Can fail for extremely low inlier ratios
- We can often do better than brute-force sampling
[Source: N. Snavely, slide credit: R. Urtasun]


## Ransac Verification


[Source: K. Grauman, slide credit: R. Urtasun]

## Summary - Stuff You Need To Know

To match image $I$ and $J$ under affine transformation:

- Compute scale and rotation invariant keypoints in both images
- Compute a (rotation invariant) feature vector in each keypoint (e.g., SIFT)
- Match all features in $I$ to all features in $J$
- For each feature in reference image $/$ find closest match in $J$
- If ratio between closest and second closest match is $<0.8$, keep match
- Do RANSAC to compute affine transformation $A$ :
- Select 3 matches at random
- Compute A
- Compute the number of inliers
- Repeat
- Find A that gave the most inliers

