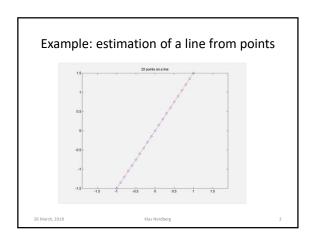
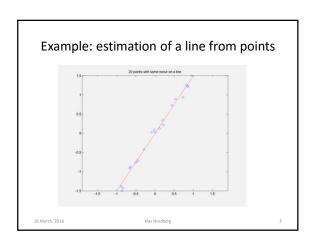
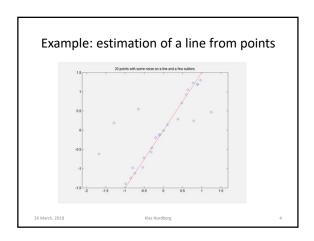
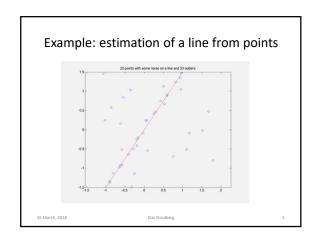
Solving the Correspondence Problem with RANSAC





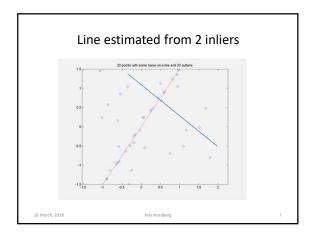




Observations

- We need (in this case!) a minimum of 2 points to determine a line
- Given such a line I, we can determine how well any other point ${\boldsymbol{y}}$ fits the line ${\boldsymbol{l}}$
- For example: distance between ${f y}$ and ${f l}$
- If we pick 2 *arbitrary* points from the dataset:
 - $-% \frac{1}{2}\left(-\frac{1}{2}\right) =0$ We can easily determine a line l

 - We can easily determine a line I
 1 is the correct line with some probability p_{IINE}
 p_{IINE} is related to the chance of picking only inliers
 p_{IINE} is larger the fewer points that are used to determine I
 In general: if I is the correct line there are more additional points that can be fitted to the line than if I is an incorrect line



Basic iteration

- 1. Pick 2 random points
- 2. Fit a line I to the points
- 3. Determine how many other points in the dataset that can be fitted to ${\bf l}$ with some minimal error ϵ .
 - This forms the consensus set C
- 4. If *C* is sufficiently large, then the fitted line is probably OK. Keep it

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Basic algorithm

- Iterate r times
 - 1. Pick 2 random points
 - 2. Fit a line I to the points
 - 3. Form the consensus set C, together with
 - Number of points in C
 - Matching error ϵ_C of the set C relative to the line
 - 4. If the consensus set is sufficiently large, then the fitted line is OK. In particular if N and/or $\epsilon_{\rm C}$ is better than the last line that was OK. Then keep it.
- For each iteration, we increase p_{SUCCESS} = the probability that the correct line has been determined
 - $-\$ We need to iterate sufficiently many time to raise $p_{\rm SUCCESS}$ to a useful level

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RANSAC

- This algorithm is called RANSAC
 - RANdom SAmple Consensus
- Published by Fischler & Bolles in 1981
 - "Random Sample Consensus: A Paradigm for Model Fitting with Applications to Image Analysis and Automated Cartography". Comm. of the ACM 24: 381–395.
- Several extensions / variations in the literature
 - Preemptive RANSAC
 - PROSAC
 - **–** ...

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RANSAC

- After r iterations, RANSAC finds a reasonable estimate of the line (i.e. from only inliers) with a probability of p
- 1 p = P(pick at least one outlier in each iteration) = $(1-w^2)^r$
- $p = 1 (1-w^2)^r$
- If w is known, we can choose r to make p as large as we want (but not = 1!)
- Example: w = 0.5
 - p = 0.94 when r = 10, p = 0.99 when r = 20

The correspondence problem

- Given a set of interest points in two images, we want to determine correspondences, i.e., pairs of points that correspond to the same 3D point
- If there is a small relative baseline:
 - Use tracking (Lucas-Kanade, etc)
 - Track POIs in image 1 to their corresponding positions in image 2
 - Can be applied to parts an image sequence
 A POL to reign the disappears of the south like in a
 - A POI typically disappears after a while in a longer sequence
 - Track-retrack
 - Remove all POIs that cannot be tracked forward and backward in time over several images

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The correspondence problem

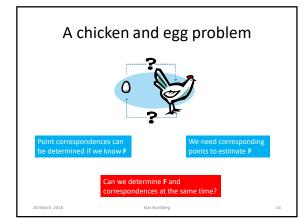
- If there are large baseline between the two images, tracking performance degrades
 - Another approach is needed





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Solving chicken and egg problem?



- Let there be two views with P₁
 points in one view and P₂ points
 in the other view
- We don't know which points in the first view that correspond to which points in the other view
- There is a set D of P₁ × P₂ possible correspondences, or tentative correspondences

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Chicken and egg revisited

 The correct correspondences can be fitted to F, i.e., they satisfy the epipolar constraint for some F that only depends on which two views are used



- They are the inliers
- The incorrect correspondences are *outliers*

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Use RANSAC

- Pick 8 random points from D
- We don't know if they really correspond, but this can be tested:
 - 1. Use the 8-point algorithm to estimate **F**
 - 2. Check how well **F** matches each pair in D
 - 3. Collect those that fit well into the consensus set C
 - 4. If C is sufficiently large: **F** is OK: keep **F** and C
- Iterate *r* times

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Probabilities

- Let w be the fraction of inliers in D
- In each iteration we pick N points that are all inliers with probability w^N (approximately)
- The probability of not all N points are inliers is then given by $1 w^N$
- The probability of not all N points are inliers in r iterations is $(1 w^N)^r$
- The probability that in K iteration, at least once, all N points are inliers: $p = 1 (1 w^N)^r$
- Solve for *r*:

 $r = \frac{\log(1-p)}{\log(1-w^N)}$

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General observation

 The expected number of iterations, r, to reach a certain probability p is

$$r = \frac{\log(1-p)}{\log(1-w^N)}$$

- For fixed p, r is reduced if w is made larger
- For fixed p, r is reduced if N is made smaller

The odds are against us

- From the outset, the set of all tentative correspondences between two images can be VERY large (= P₁ × P₂)
- VERY few of these are inliers: w is VERY small
- Here N = 8
- This means that r must be VERY⁸ large in order to make p close to 1

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The correspondence problem

- The correspondence problem is often addressed by finding two sets of points that we want to bring into correspondence
 - Typically: interest points in images (POI)
 - Typically: different number of points in the sets
- · Without any outer information:
 - Any point in set 1 can correspond to any point in set 2
 - In practice, often not a feasible approach!
 - Too many outliers (w too small)

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Visual appearance and RANSAC

- The set of correspondences in D has m possible correspondences and only m₀ of them are correct (m - m₀ are incorrect)
- Probability of picking a correct correspondence w = m₀/m
- If we can reduce the number incorrect correspondences, without removing correct ones, m will decrease while m₀ is constant
 - \Rightarrow w increases \Rightarrow r decreases for fixed p

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Matching matrix

- Given P₁ points in image 1 and P₂ points in image 2
 - Form a $P_1 \times P_2$ matching matrix
 - Each entry (i,j) is a hypothetical correspondence between point i in image 1 and point j in image 2
- Set entry (*i,j*) =
 - a matching score between point *i* and point *j*
- For each column or row: keep only the largest entry
 - lacktriangle Reduces m while keeping m_0 constant
- w increases $\Rightarrow r$ decreases for fixed p
- Run RANSAC on remaining tentative pairs

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Matching matrix

Score matrix

Each entry in the matching matrix describes how well a certain point in image 1 matches another point in image 2. For example: high score = good match

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Matching matrix

- The matching score can be based on *similarity of visual appearance* or other a priori knowledge about the scene (rather than geometric properties)
- For example
 - SIFT features [see previous lecture!]
 - MSER [see previous lecture!]
 - Color description
 - Camera motions in relation to scene depth
 - Tracking quality
- The resulting correspondences are referred to as
 - Tentative correspondences
 - *Putative* correspondences

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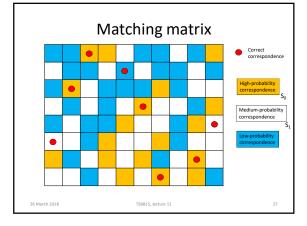
Matching matrix

- Threshold the matching scores to remove highprobability outliers and to identify high-probability inliers (two thresholds!)
 - Remove high-probability outliers
 - High probability inliers means > 50% probability
- From the original set D of possible correspondences, we have form two sets D_1 and D_0 such that
 - D₀ contains the high-probability inliers
 - A.k.a. putative correspondences
 - D₁ contains the remaining correspondences that are not high-probability outliers

 $\blacksquare \ D_0 \subset D_1 \subset D$

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Visual appearance and RANSAC

- Remove the low-probability correspondences before RANSAC
- Use the RANSAC algorithm for finding corresponding points based on the tentative correspondences
 - Use only high-probability inliers (D_0) in the initial selection of n points: w > 0.5
 - \Rightarrow fewer iterations are needed
 - Use medium and high-probability correspondences (D_1) to form the consensus step \Rightarrow increases the probability of including correct correspondences in the consensus set

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E vs. F

- If we estimate F in each RANSAC iteration, then we need N = 8 correspondences to determine F
- If instead E is determined, it is sufficient with N = 5 correspondences
 - In practice 6, since we get multiple solutions for E
- If the internal calibration K is known, we can reduce r = number of RANSAC iterations, by using E instead of F