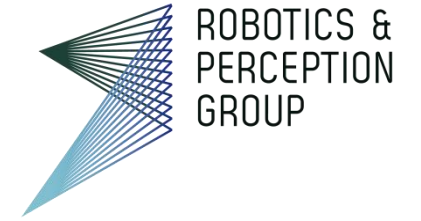




University of  
Zurich<sup>UZH</sup>



# Vision Algorithms for Mobile Robotics

## Lecture 09 Multiple View Geometry 3

Davide Scaramuzza

<http://rpg.ifi.uzh.ch>

# Lab Exercise 7 – Today

Implement the P3P algorithm and RANSAC.  
Additionally, we will outline the mini projects



# Outline

- Robust Structure from Motion
- Bundle Adjustment

# Robust Estimation

- Matched points are usually contaminated by **outliers** (i.e., wrong image matches).

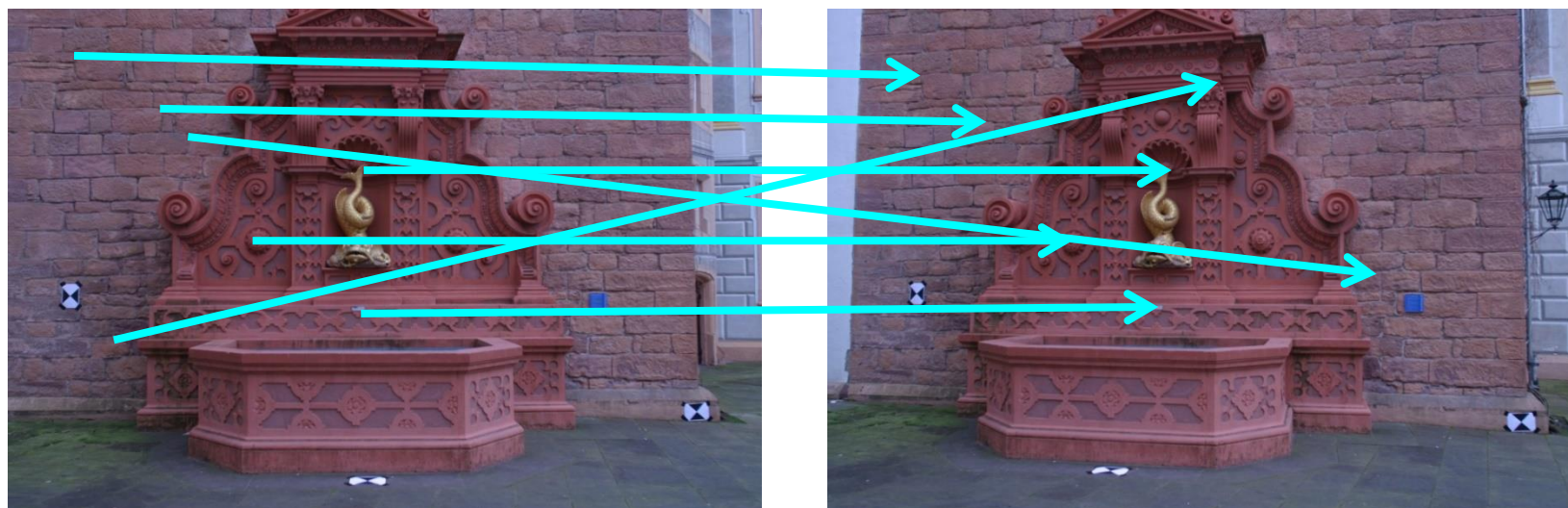


Image 1

Image 2

# Robust Estimation

- Matched points are usually contaminated by **outliers** (i.e., wrong image matches).
- Causes of outliers are:
  - Repetitive features (i.e., features with the same appearance)
  - Geometric and photometric changes to which the descriptor is not invariant
  - Large image noise
  - Occlusions
  - Moving objects
  - Image or motion blur
- For reliable and accurate visual odometry, outliers must be removed
- This is the task of **Robust Estimation**

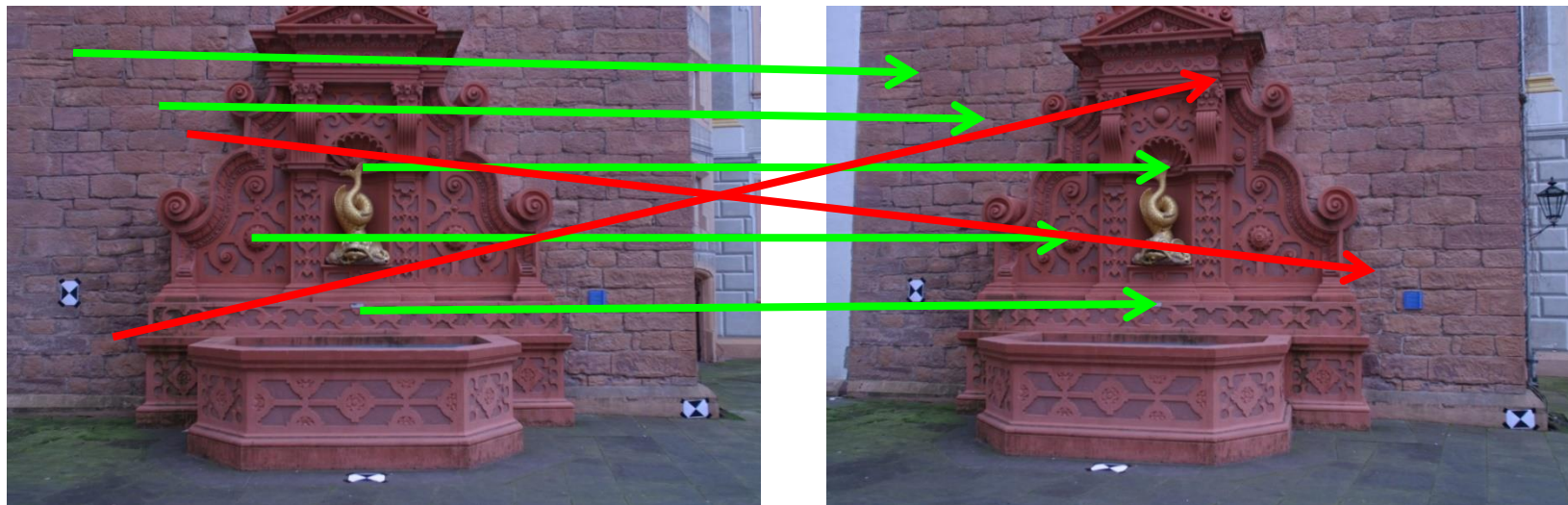
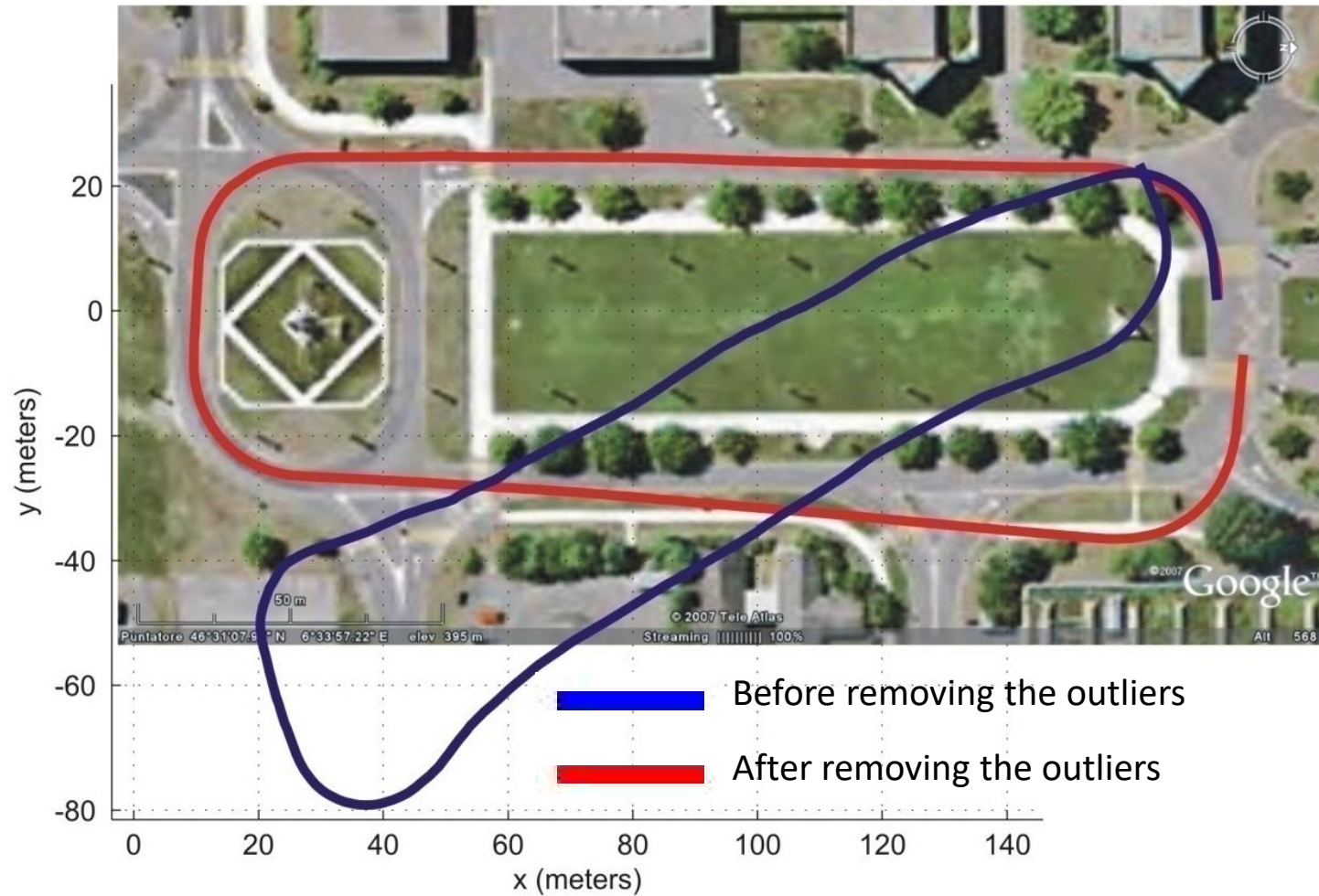


Image 1

Image 2

# Effect of Outliers on Visual Odometry



# Expectation Maximization (EM) algorithm

- EM is a simple **method for model fitting in the presence of outliers** (very noisy points or wrong data)
- It can be applied to all sorts of problems where the goal is to **estimate the parameters of a model from the data** (e.g., camera calibration, Structure from Motion, DLT, PnP, P3P, Homography, etc.)
- Let's review EM applied to the line fitting problem

[1] Dellaert, *The expectation maximization algorithm*, Georgia Institute of Technology, 2002. [PDF](#) (explains the original papers below)

[2] Hartley, *Maximum likelihood estimation from incomplete data*, Biometrics, 1958.

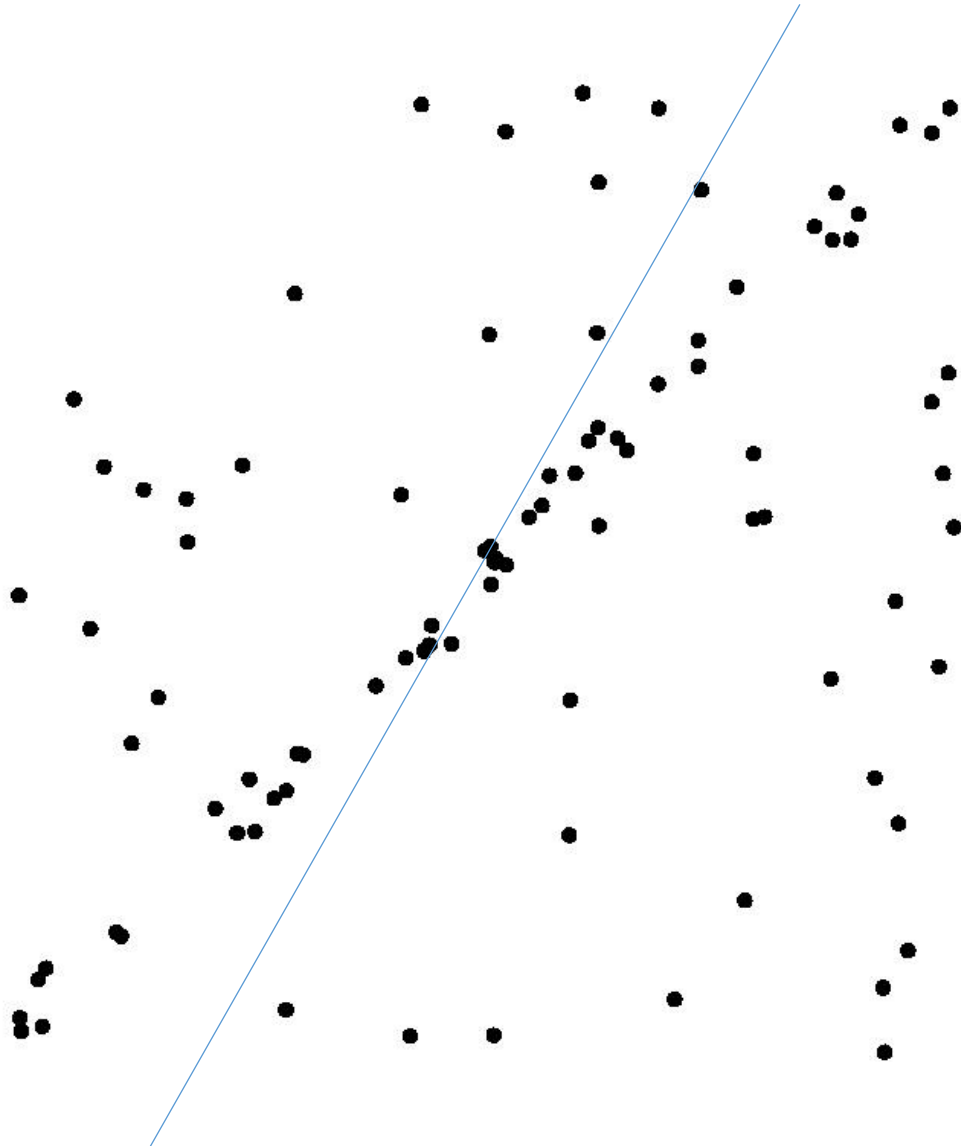
[3] Dempster, Laird, Rubin, *Maximum likelihood from incomplete data via the EM algorithm*, Journal of the Royal Statistical Society, 1977.

# EM applied to line fitting



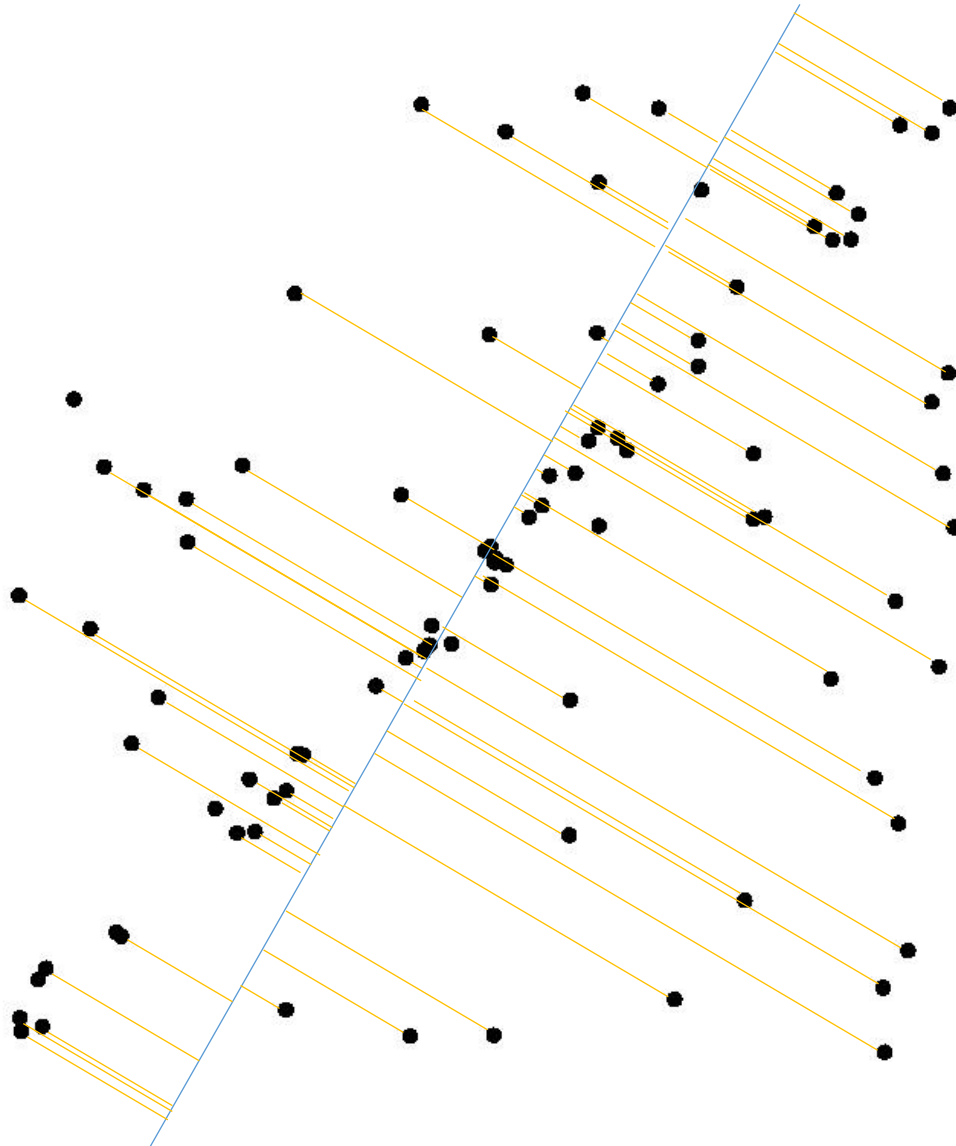


# EM applied to line fitting



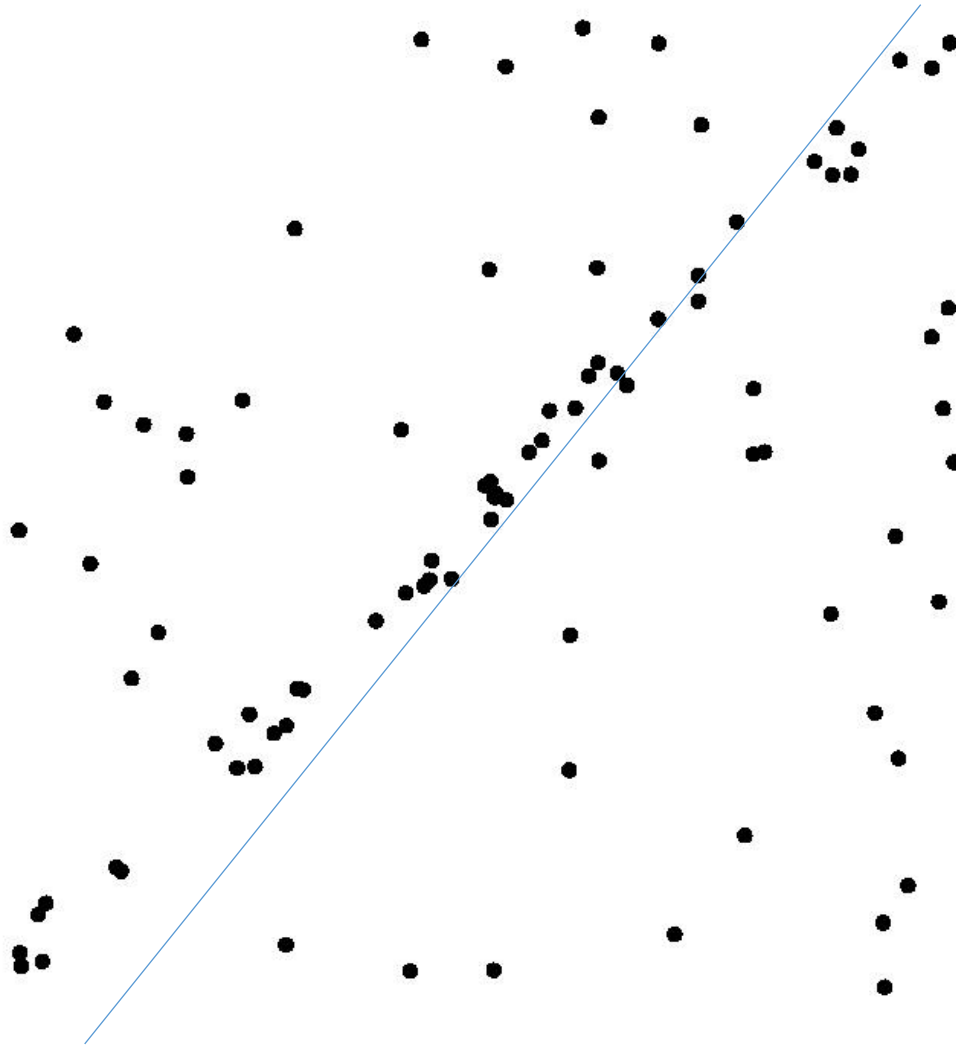
1. Estimate line parameters that fit all data points (e.g., using least-square:  $\min \sum r_i^2$ , where  $r_i$  is the point-to-line distance)

# EM applied to line fitting



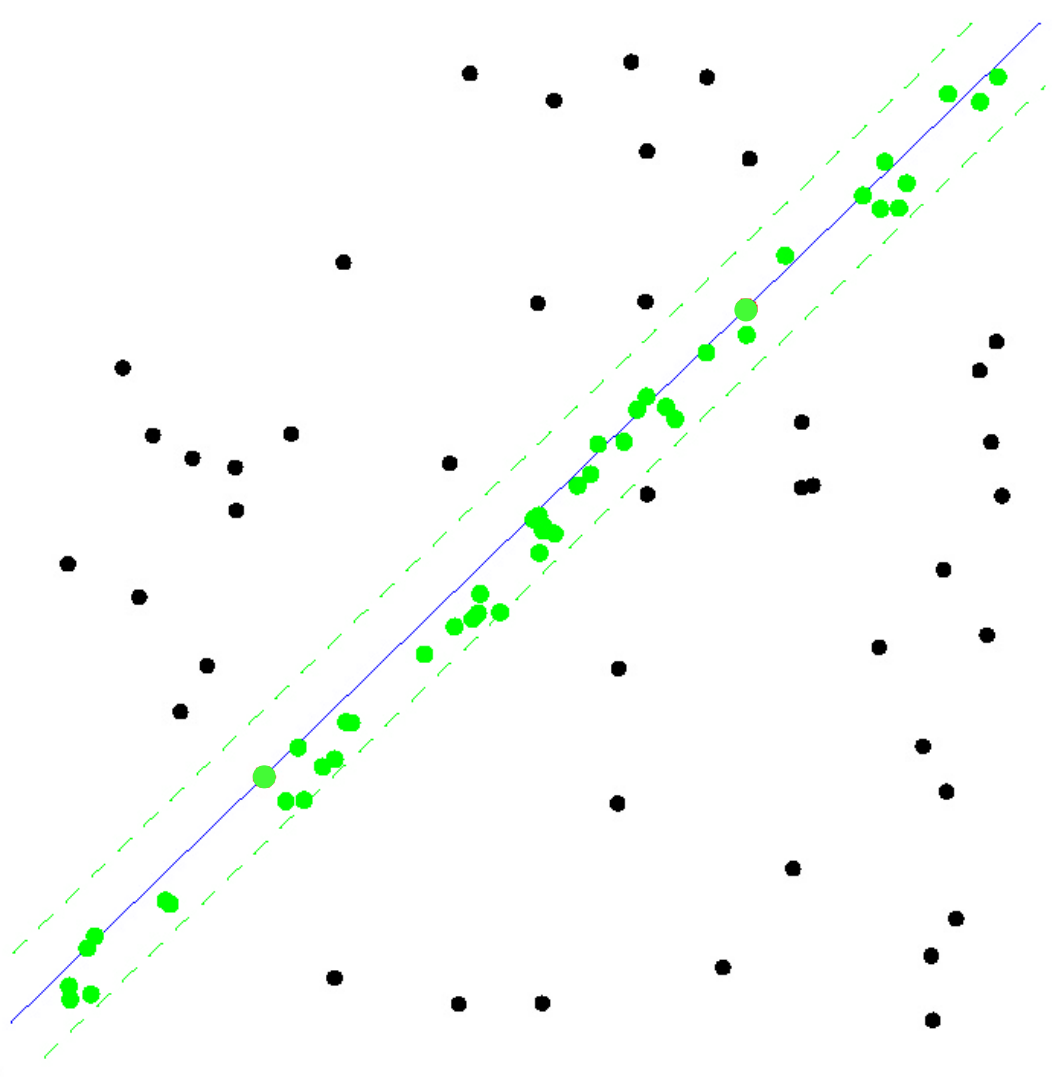
1. Estimate line parameters that fit all data points (e.g., using least-square:  $\min \sum r_i^2$ , where  $r_i$  is the point-to-line distance)
2. Calculate residual error  $r_i$  for each data point and assign it a weight (e.g.,  $w_i = e^{-r_i^2}$  representing the likelihood that such assignment is correct (estimates the **Expectation**))

# EM applied to line fitting



1. Estimate line parameters that fit all data points (e.g., using least-square:  $\min \sum r_i^2$ , where  $r_i$  is the point-to-line distance)
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3. Re-estimate line parameters (e.g., using weighted least-squares:  $\min \sum w_i r_i^2$ ) (**Maximization Step**)

# EM applied to line fitting



1. Estimate line parameters that fit all data points (e.g., using least-square:  $\min \sum r_i^2$ , where  $r_i$  is the point-to-line distance)
2. Calculate residual error  $r_i$  for each data point and assign it a weight (e.g.,  $w_i = e^{-r_i^2}$  representing the likelihood that such assignment is correct (estimates the **Expectation**))
3. Re-estimate line parameters (e.g., using weighted least-squares:  $\min \sum w_i r_i^2$ ) (**Maximization Step**)
4. Iterate 2 and 3 till convergence
5. Select as **inliers** the data points with weight higher than a threshold

# Problem of EM algorithm

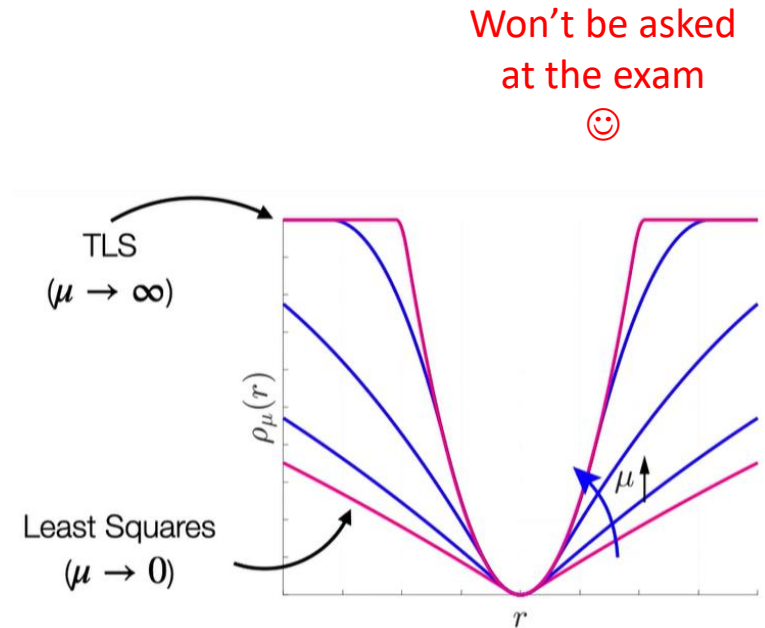
## **Very sensitive to initial condition:**

- This is because EM selects the initial condition by minimizing the sum of squared residuals  $\sum r_i^2$ .
- While this is a convex function, the result is strongly influenced by a few large error values (e.g., outliers).
- Thus, EM converges to the wrong solution if initial condition is far from the true one
  
- Alternative options:
  - GNC algorithm
  - RANSAC algorithm

# Graduated Non-Convexity algorithm (GNC)

**Idea: optimize a surrogate function  $\sum \rho_{\mu}(r_i)$ , where  $\mu$  controls the amount of non-convexity.**

- Start by solving the non-robust convex optimization function ( $\mu \rightarrow 0$ , i.e., least squares)
- At each iteration, gradually increase non-convexity ( $\mu \rightarrow \infty$ ) and recompute weights  $w_i$  till we achieve the desired level of robustness.
- It is shown in [1] to be robust up to 90% of outliers with five times fewer iterations than RANSAC.
- However, RANSAC can cope with even more than 90% outliers.



[1] Yang, Antonante, Tzoumas, Carlone, *Graduated Non-Convexity for Robust Spatial Perception: From Non-Minimal Solvers to Global Outlier Rejection*, International Conference on Robotics and Automation (ICRA), 2020. **Best paper award in Robot Vision.** [PDF](#). [Code](#).

[2] Blake, Zisserman, *Visual Reconstruction*. MIT Press, Cambridge, Massachusetts, 1987.

# RANSAC (RANdom SAmple Consensus)

- RANSAC is the **standard method for model fitting in the presence of outliers** (very noisy points or wrong data)
- It is **non-deterministic**: you get a different result everytime you run it
- It is **not sensitive to the initial condition**, and **does not get stuck in local maxima**
- It can be applied to all sorts of problems where the goal is to **estimate the parameters of a model from the data** (e.g., camera calibration, Structure from Motion, DLT, PnP, P3P, Homography, etc.)
- Let's review RANSAC for line fitting and see how we can use it to do Structure from Motion

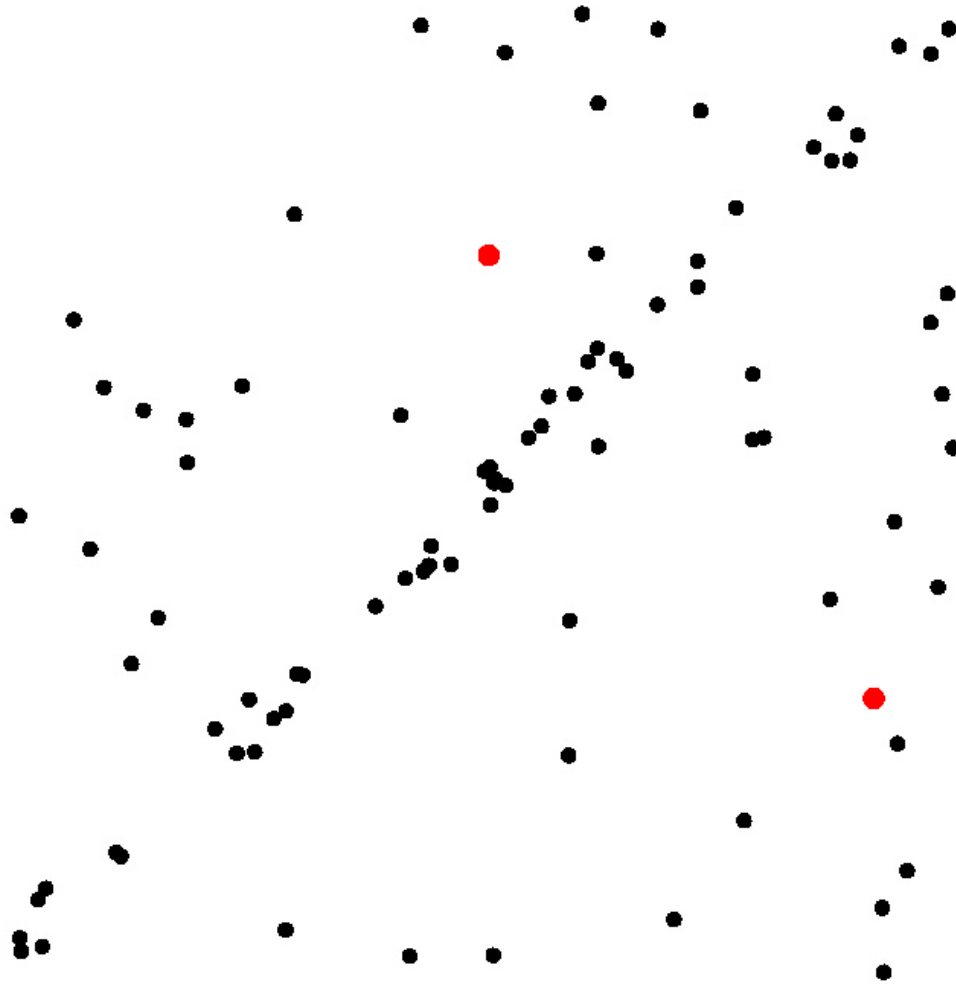
M. A. Fischler and R. C. Bolles. Random sample consensus: A paradigm for model fitting with applications to image analysis and automated cartography. Graphics and Image Processing, 1981. [PDF](#).

# RANSAC



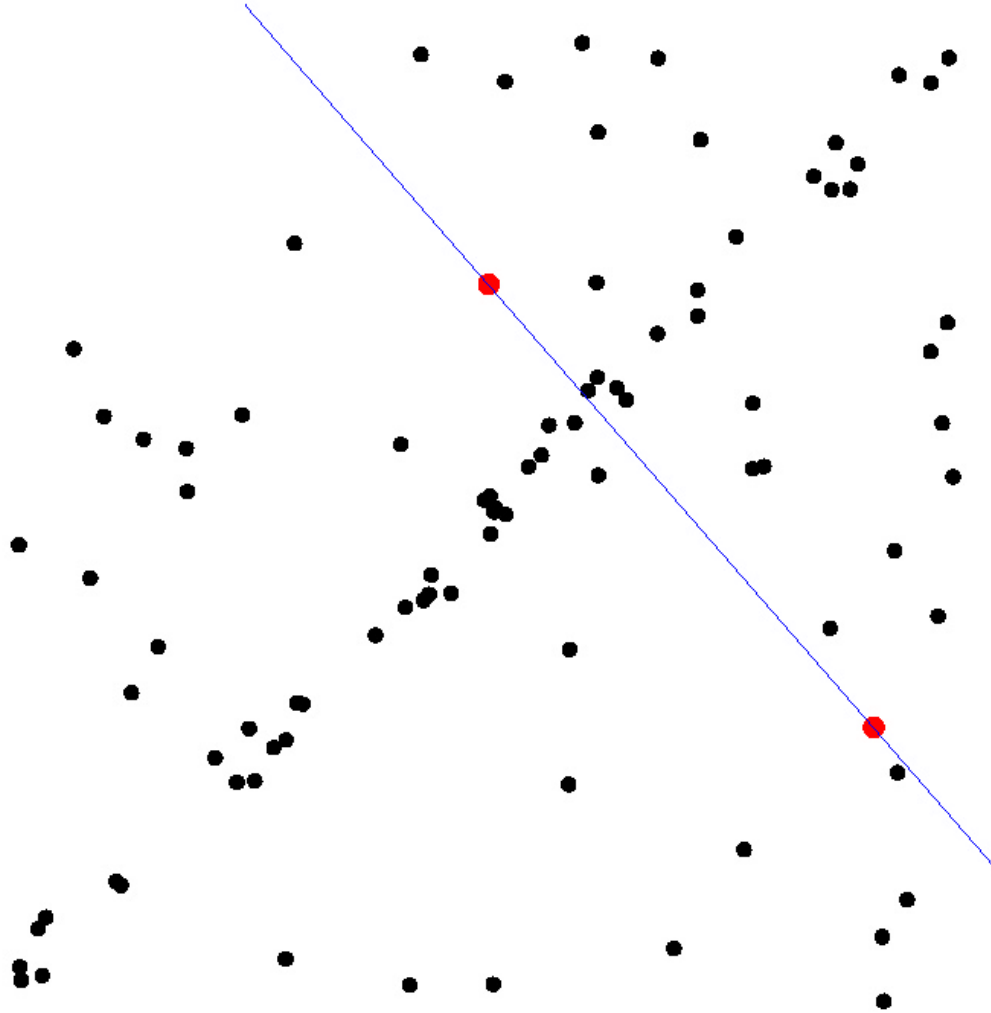


# RANSAC



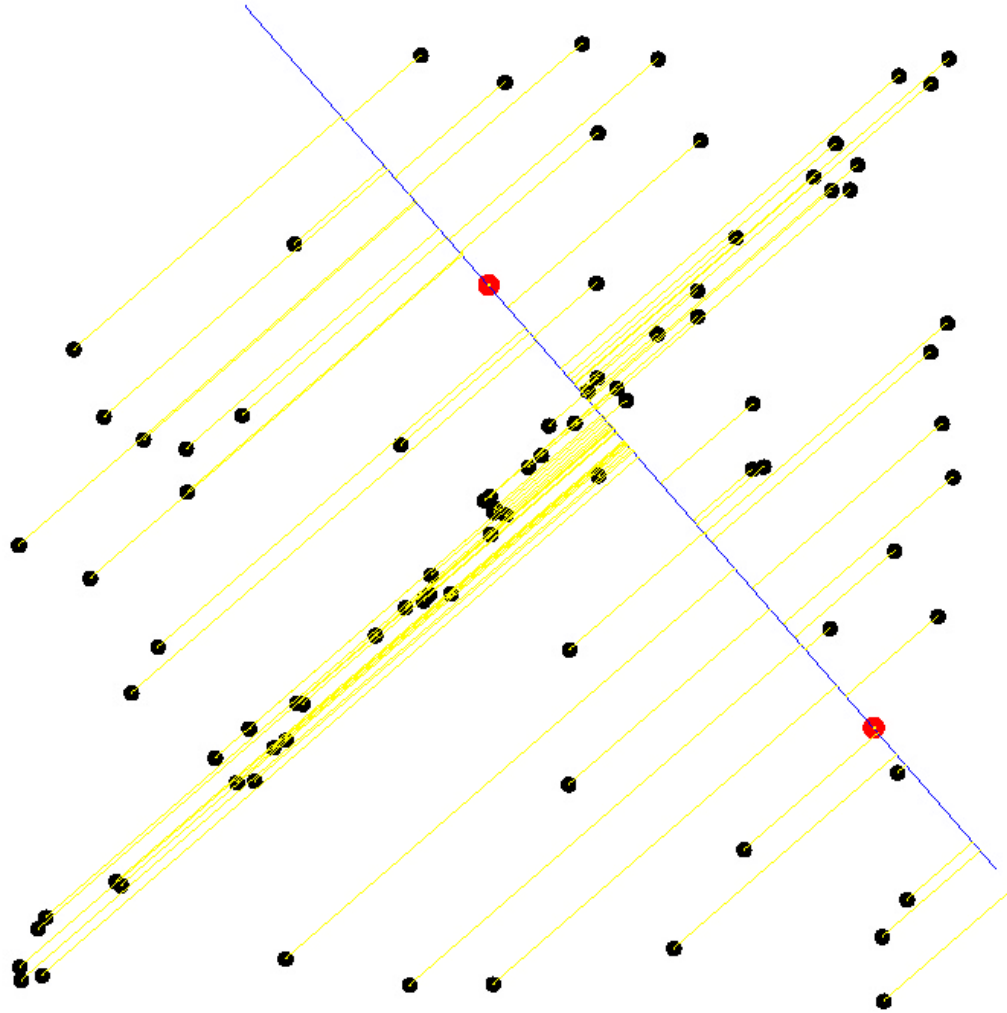
1. Select a sample of 2 points at *random*

# RANSAC



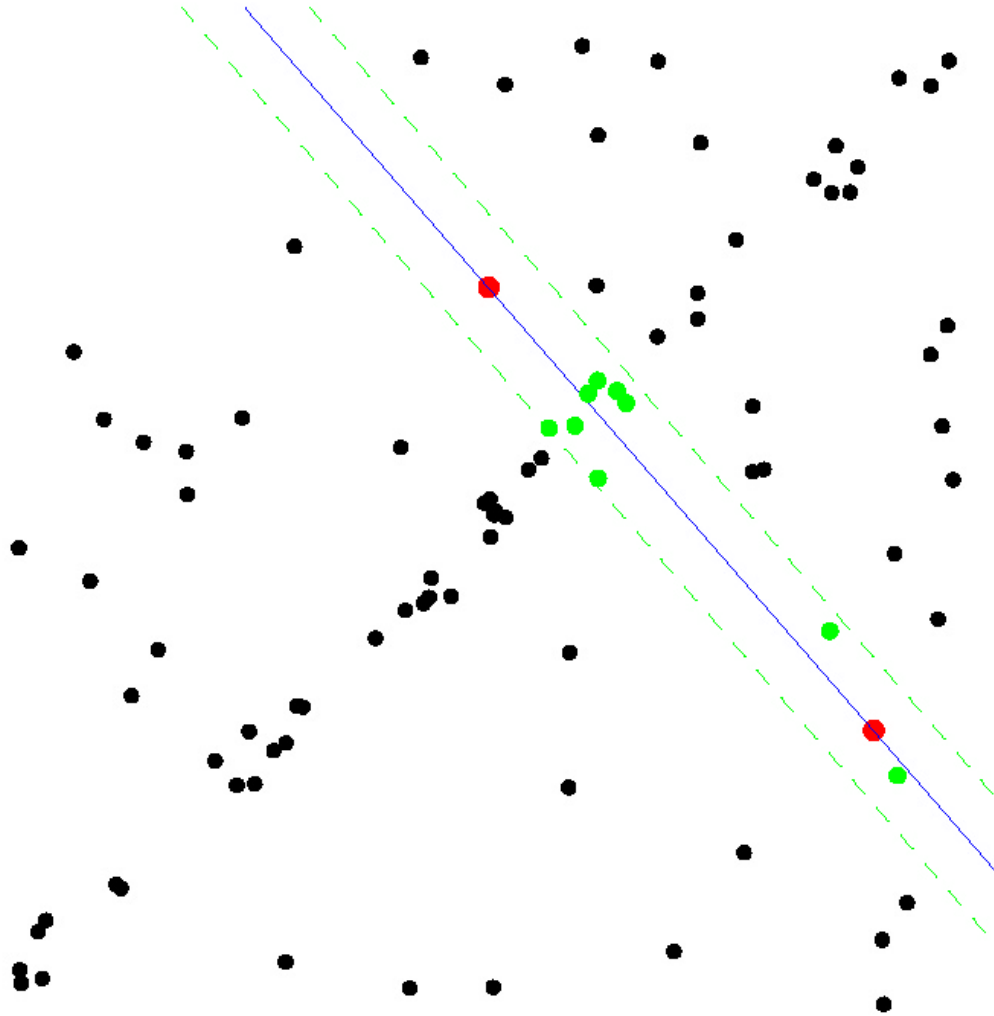
1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample

# RANSAC



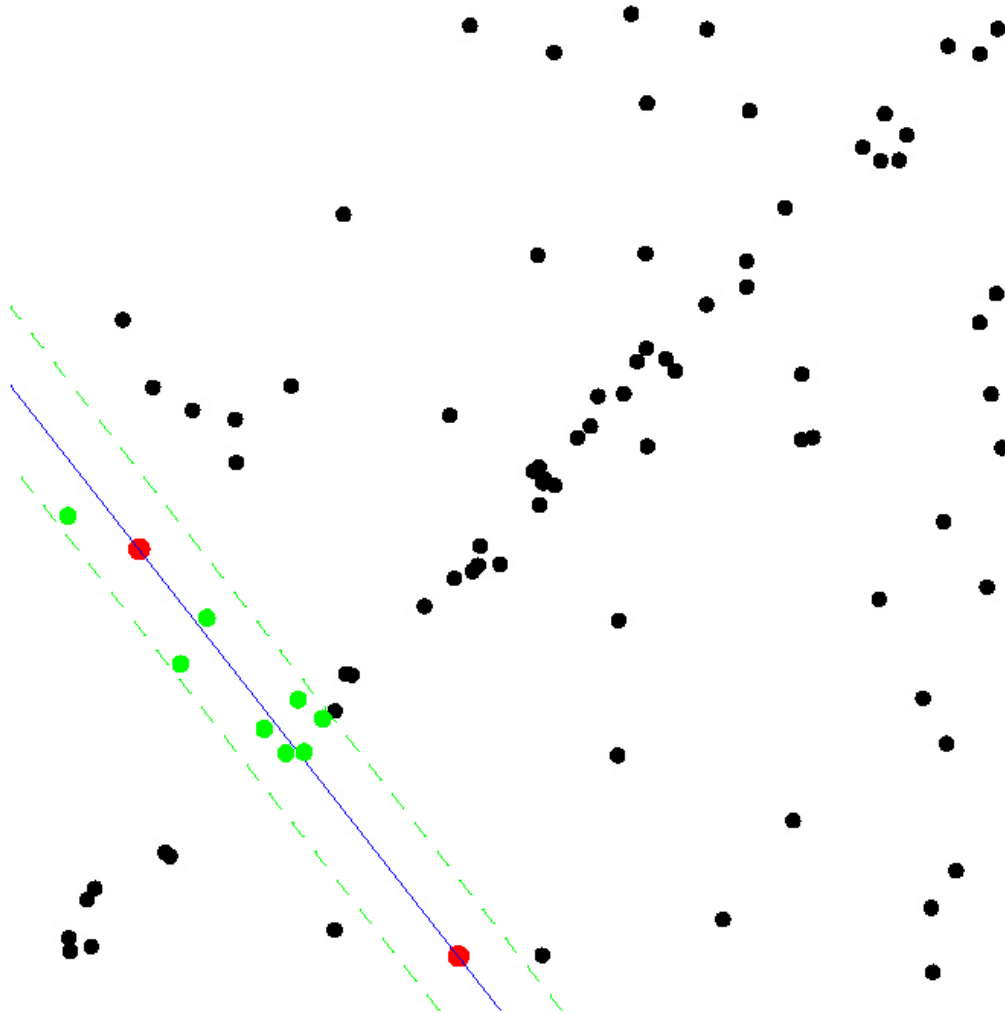
1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample
3. Calculate the residual error for each data point

# RANSAC



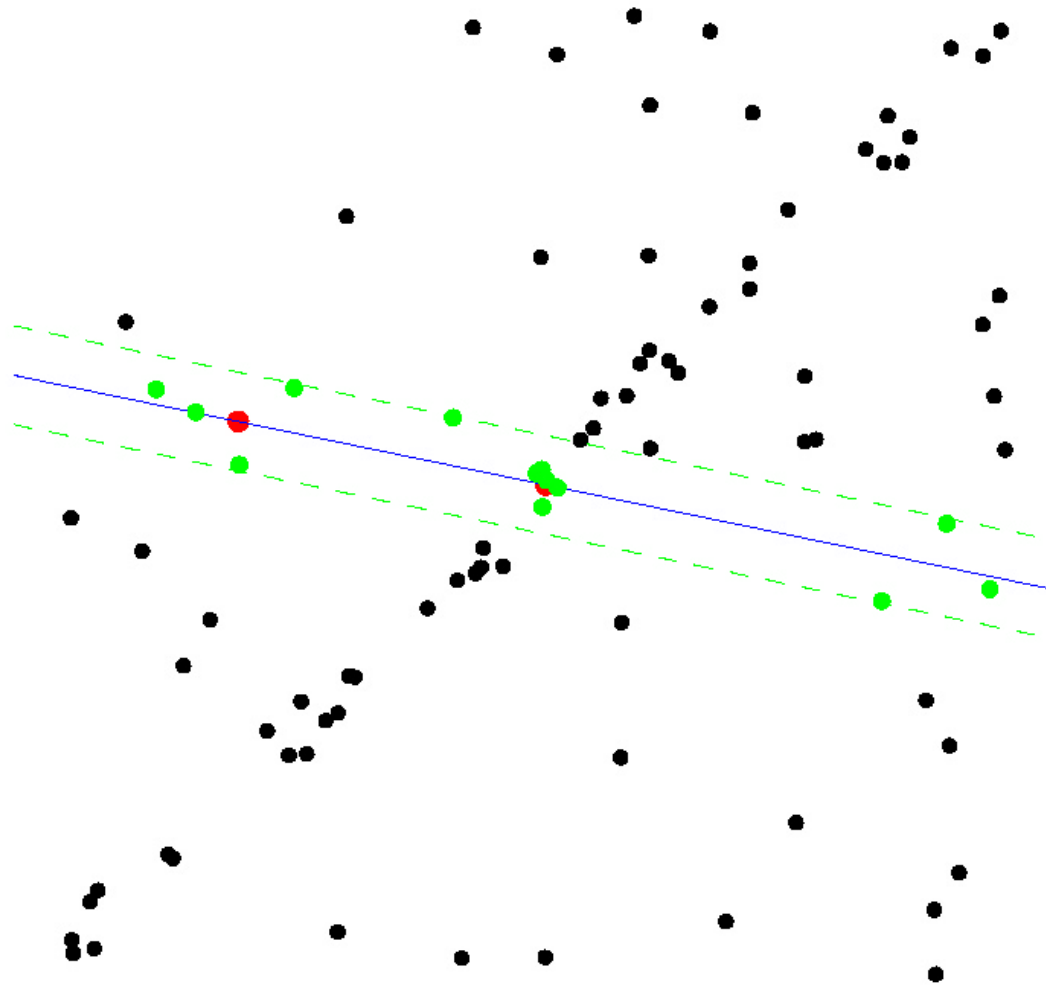
1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample
3. Calculate the residual error for each data point
4. **Select data that support current hypothesis**

# RANSAC



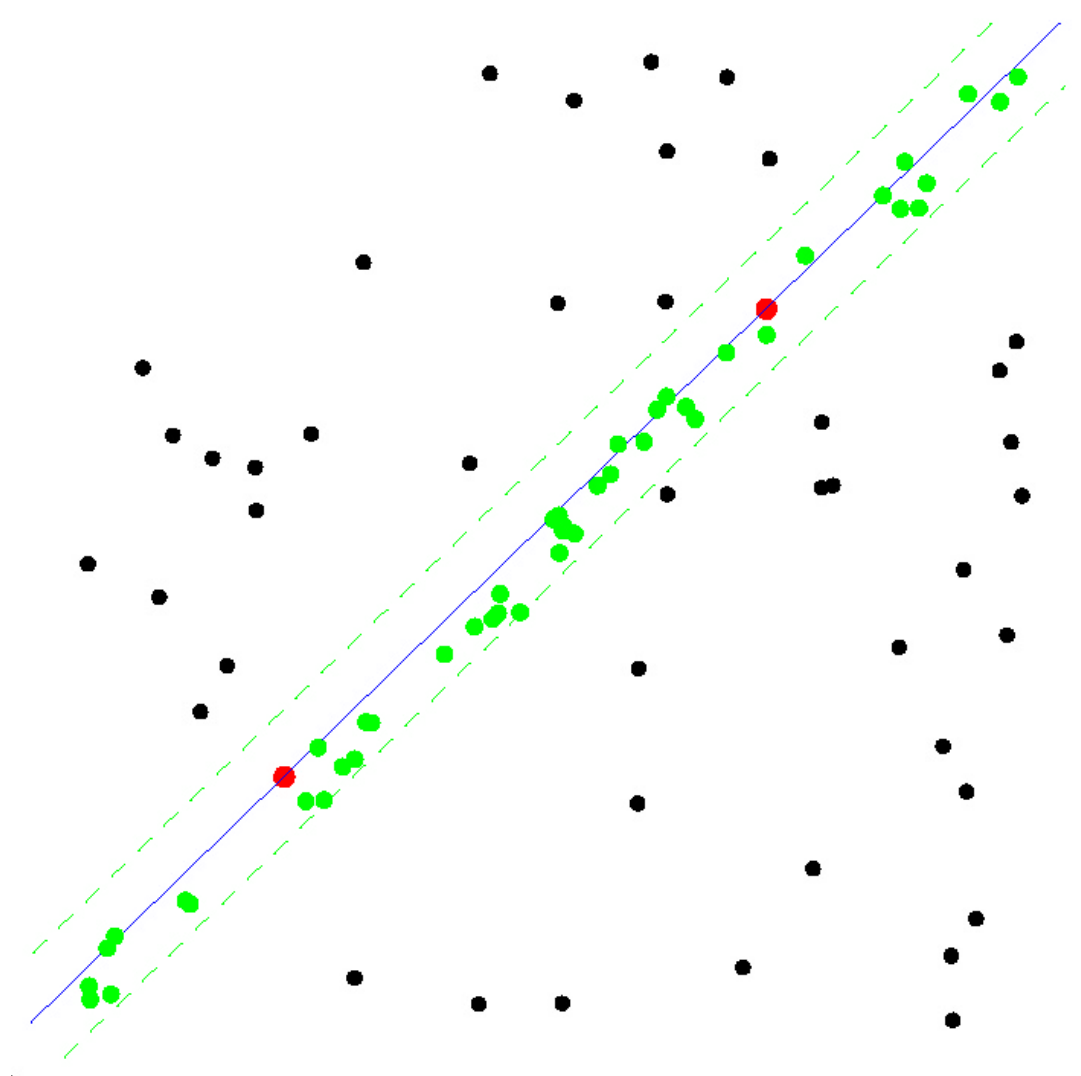
1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample
3. Calculate the residual error for each data point
4. Select data that support current hypothesis
5. **Repeat from step 1 for  $k$  times**

# RANSAC



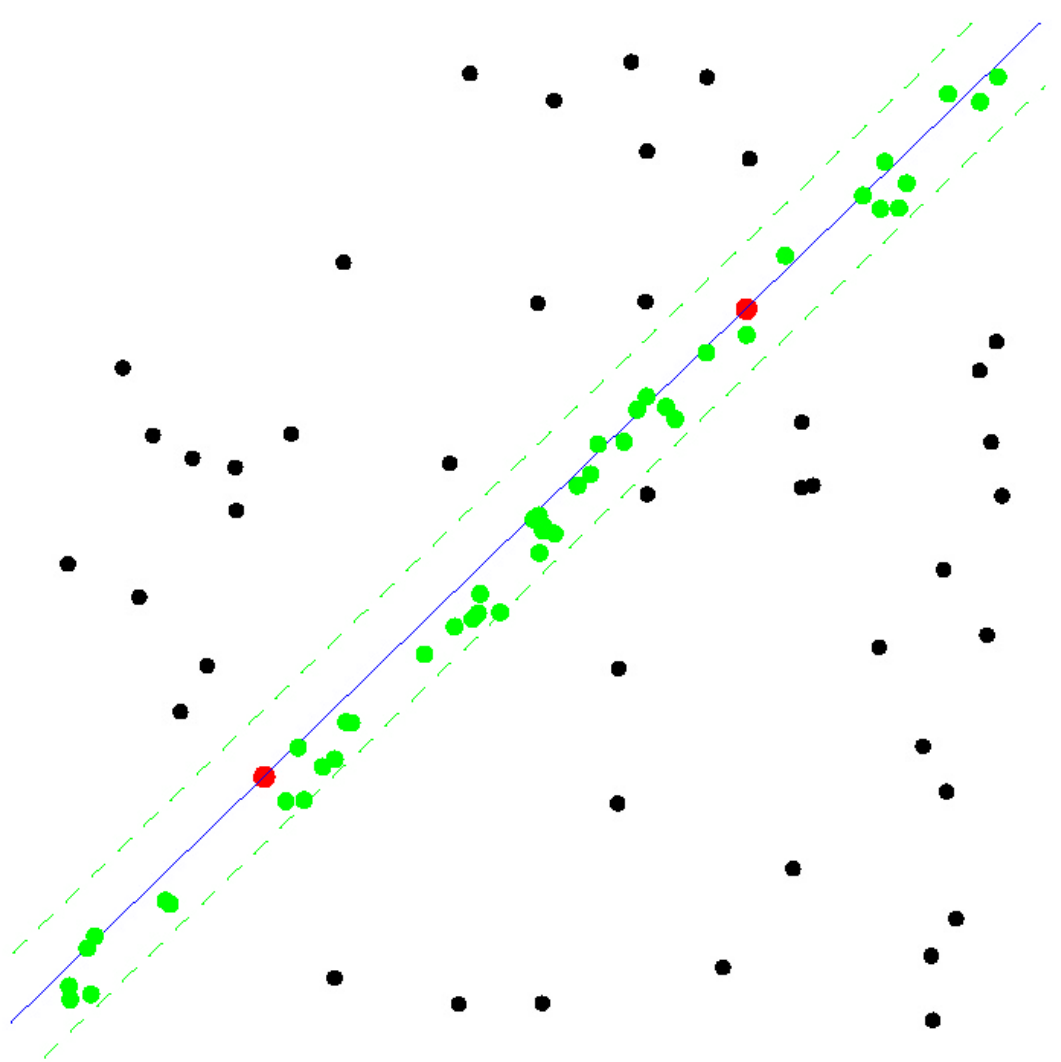
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# RANSAC



1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample
3. Calculate the residual error for each data point
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5. Repeat from step 1 for  $k$  times
6. **Select the set with the maximum number of inliers obtained within  $k$  iterations**

# RANSAC



1. Select a sample of 2 points at *random*
2. Calculate model parameters that fit the data in the sample
3. Calculate the residual error for each data point
4. Select data that support current hypothesis
5. Repeat from step 1 for  $k$  times
6. **Select the set with the maximum number of inliers obtained within  $k$  iterations**
7. Finally, calculate the model parameters using **all the inliers**

**NB: RANSAC is non deterministic:** every time you run it you may get a different result (due to the random hypotheses' generation process). Conversely, **EM and GNC are deterministic**



# RANSAC

- How many iterations does RANSAC need?
- Ideally: check all possible combinations of 2 points in a dataset of  $N$  points.
- Number of all pairwise combinations:  $\frac{N(N-1)}{2}$ 
  - computationally unfeasible if  $N$  is too large.  
**Example, for 1000 points you need to check all  $1000 \times 999 / 2 \cong 500'000$  possibilities!**
- Do we really need to check all possibilities or can we stop RANSAC after some iterations?
  - We will see that it is **enough to check a subset of all combinations if we have a rough estimate of the percentage of inliers** in our dataset
  - This can be done in a **probabilistic way**

# RANSAC

- How many iterations does RANSAC need?
- $N$  := total number of data points
- $w$  := number of inliers /  $N \rightarrow w$ : fraction of inliers in the dataset  $\rightarrow w = P(\text{selecting an inlier-point out of the dataset})$
- Assumption: the 2 points necessary to estimate a line are selected independently
  - $\rightarrow w^2 = P(\text{both selected points are inliers})$
  - $\rightarrow 1 - w^2 = P(\text{at least one of these two points is an outlier})$
- Let  $k$  be the number of RANSAC iterations executed so far
- $\rightarrow (1 - w^2)^k = P(\text{RANSAC never selected two points that are both inliers after } k \text{ iterations})$
- Let  $p$  := Probability to have selected at least two points that are both inliers after  $k$  iterations. We call  $p$  *Probability of Success*
- $\rightarrow 1 - p = (1 - w^2)^k$  and therefore:

$$k = \frac{\log(1 - p)}{\log(1 - w^2)}$$

# RANSAC

- How many iterations does RANSAC need?

$$k = \frac{\log(1 - p)}{\log(1 - w^2)}$$

→ knowing the fraction of inliers  $w$ , after  $k$  iterations we will have a probability  $p$  of finding a set of points free of outliers

- Example: if we want a probability of success  $p = 99\%$  and we know that  $w = 50\% \rightarrow k = 16$  iterations
  - these are **significantly fewer** than the number of **all possible combinations (500,000)!**
  - **Notice: the number of data points does not influence the minimum number of iterations  $k$ , only  $w$  does!**
- In practice we only need a rough estimate of  $w$ . More advanced variants of RANSAC estimate the fraction of inliers and adaptively update it at every iteration (**how?**)

# RANSAC applied to Line Fitting

1. Initial: let  $A$  be a set of  $N$  points
2. **repeat**
3. Randomly select a sample of **2** points from  $A$
4. **Fit a line** through the **2** points
5. Compute the **distances** of all other points **from this line**
6. Construct the inlier set (i.e. count the number of points whose distance  $< d$ )
7. Store these inliers
8. **until** maximum number of iterations  $k$  reached
9. The set with the maximum number of inliers is chosen as a solution to the problem

$$k = \frac{\log(1 - p)}{\log(1 - w^2)}$$

# RANSAC applied to General Model Fitting

1. Initial: let  $A$  be a set of  $N$  points
2. **repeat**
3. Randomly select a sample of  $s$  points from  $A$
4. **Fit a model** from the  $s$  points
5. Compute the **distances** of all other points **from this model**
6. Construct the inlier set (i.e. count the number of points whose distance  $< d$ )
7. Store these inliers
8. **until** maximum number of iterations  $k$  reached
9. The set with the maximum number of inliers is chosen as a solution to the problem

$$k = \frac{\log(1 - p)}{\log(1 - w^s)}$$

# RANSAC applied to General Model Fitting

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7. Store these inliers
8. **until** maximum number of iterations  $k$  reached
9. The set with the maximum number of inliers is chosen as a solution to the problem

$$k = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^s)}$$

NB: The formula is more commonly written as a function of the **fraction of outliers  $\varepsilon$**

# The Three Key Ingredients of RANSAC

In order to implement RANSAC for Structure From Motion (SFM), we need three key ingredients:

1. What's the **model** in SFM?
2. What's the **minimum number of points** to estimate the model?
3. How do we compute the distance of a point from the model? In other words, can we define a **distance metric** that measures how well a point fits the model?

# Answers

## 1. What's the model in SFM?

- The **Essential Matrix** (for calibrated cameras) or the **Fundamental Matrix** (for uncalibrated cameras)
- Alternatively, **R** and **T**

## 2. What's the **minimum number of points** to estimate the model?

1. We know that 5 points is the theoretical minimum number of points for calibrated cameras
2. However, if we use the *8-point algorithm*, then **8** is the minimum (for both calibrated or uncalibrated cameras)

## 3. How do we compute the **distance** of a point from the model?

1. Algebraic error
2. Directional error
3. Epipolar line distance
4. Reprojection error



# Example: 8-point RANSAC applied to SFM

- Let's consider the following image pair and its image correspondences (e.g., Harris, SIFT, etc.), denoted by arrows



Image 1

Image 2

# Example: 8-point RANSAC applied to SFM

- Let's consider the following image pair and its image correspondences (e.g., Harris, SIFT, etc.), denoted by arrows
- For convenience, we overlay the features of the second image on the first image and use arrows to denote the *motion vectors* of the features

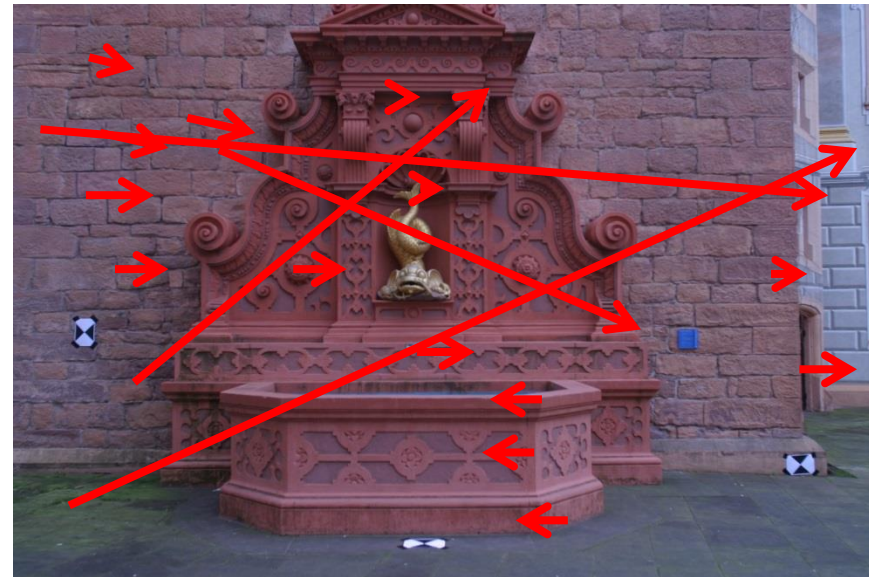


Image 1

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1. Randomly select 8 point correspondences and compute the model

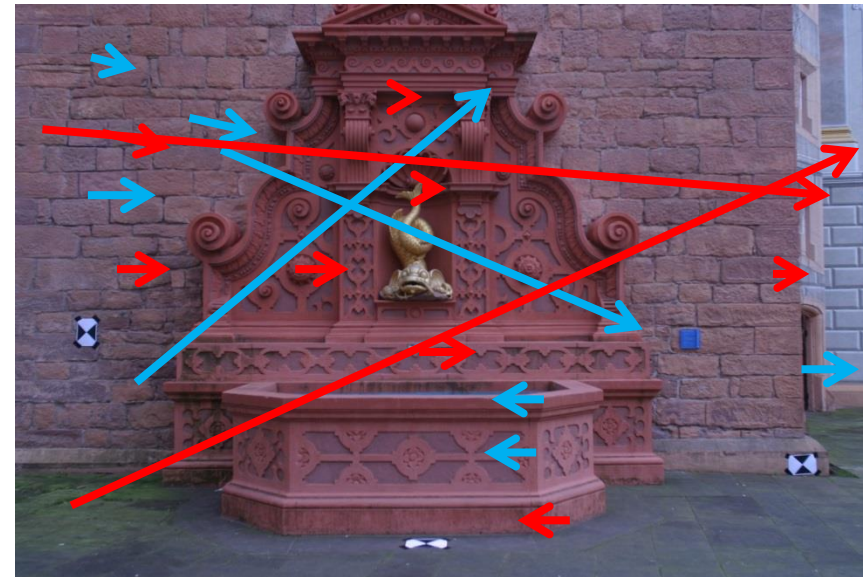


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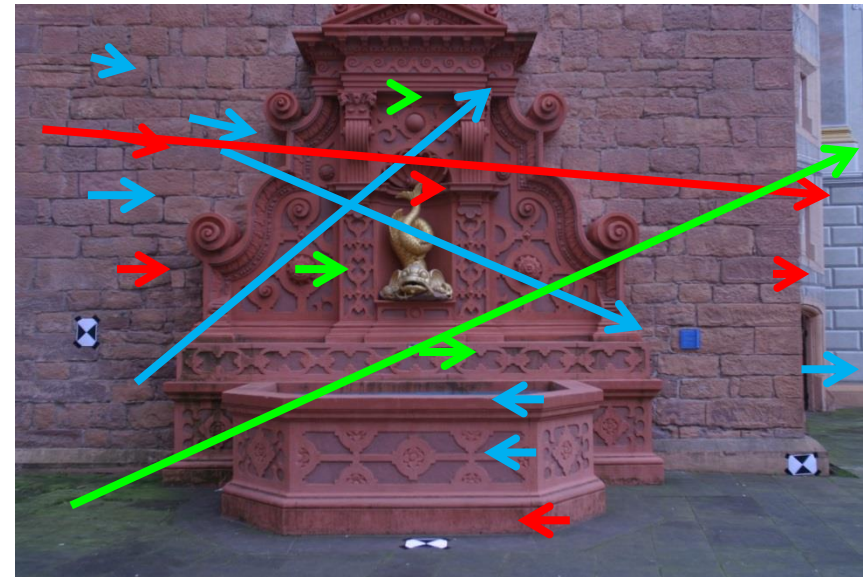


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2. Compute distance of all other points from this model and count the inliers
3. Repeat from 1

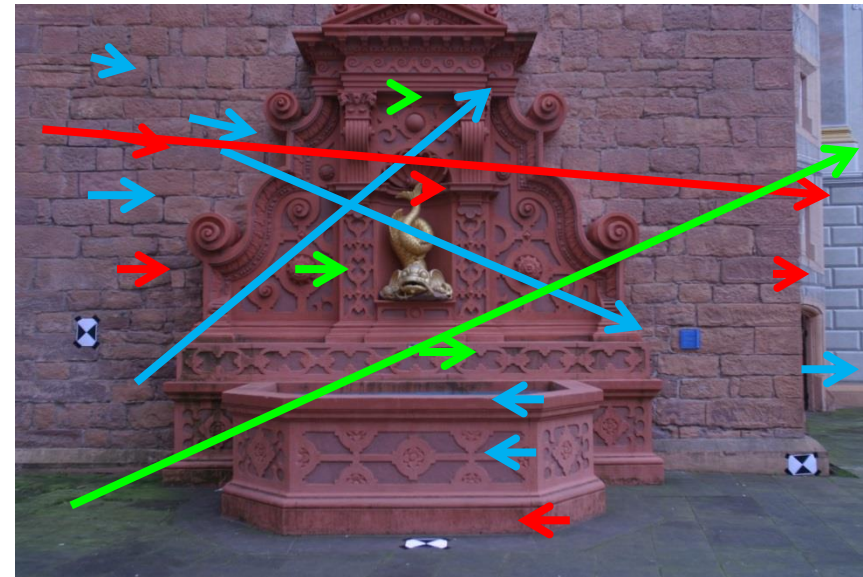


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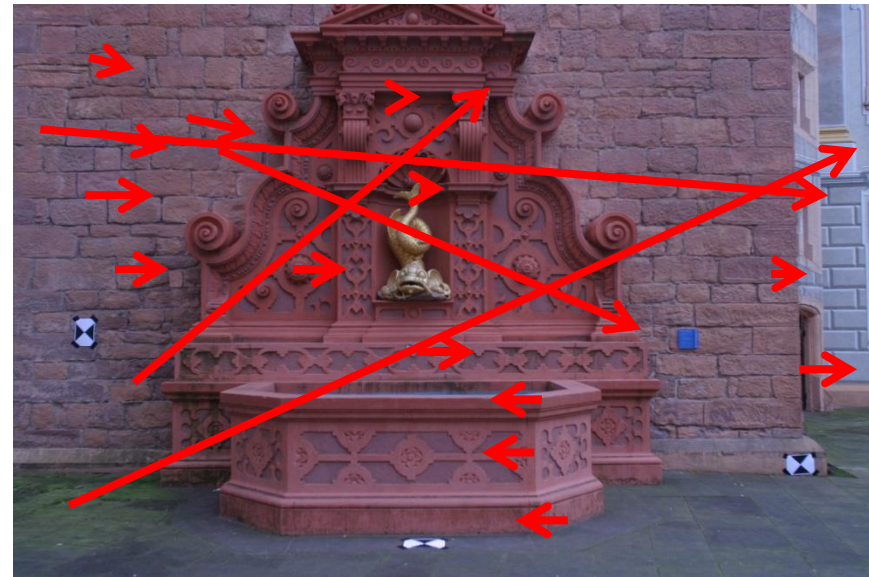


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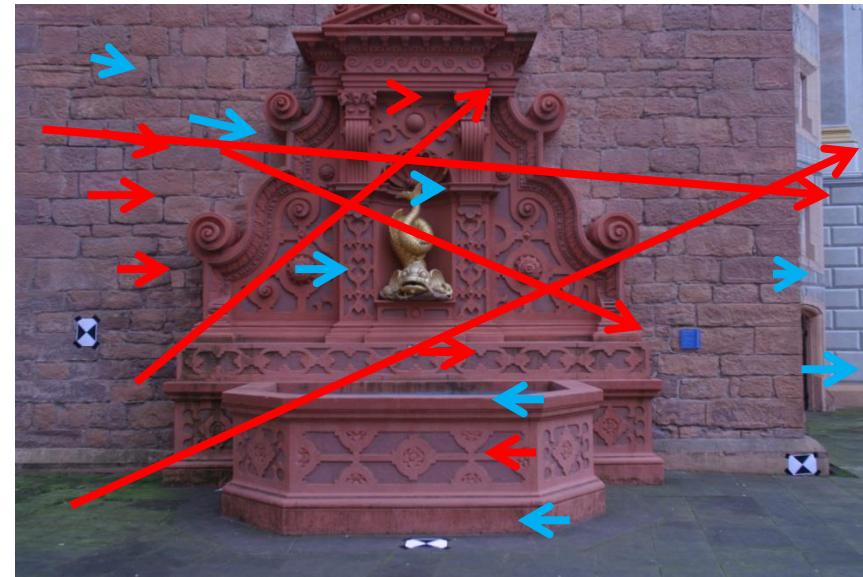


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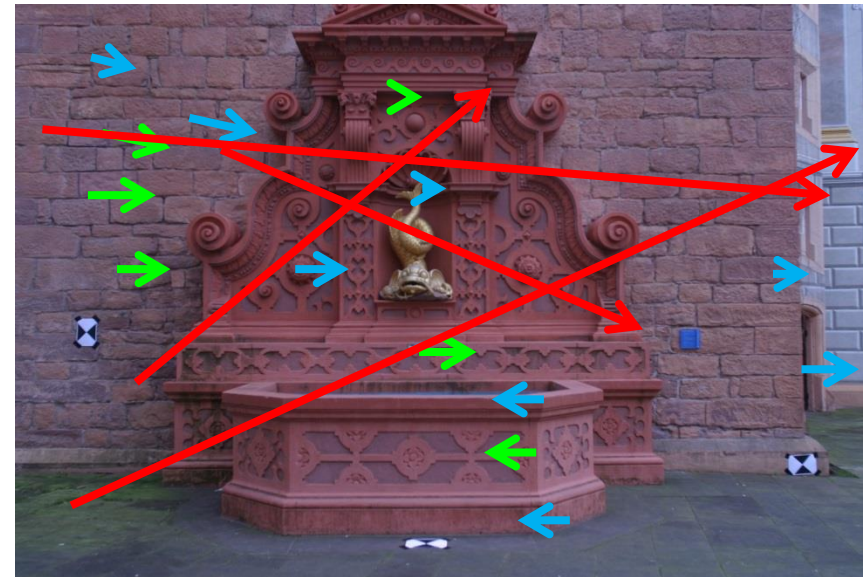


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1. Randomly select 8 point correspondences and compute the model
2. Compute distance of all other points from this model and count the inliers
3. Repeat from 1 for  $k$  times

$$k = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^8)}$$

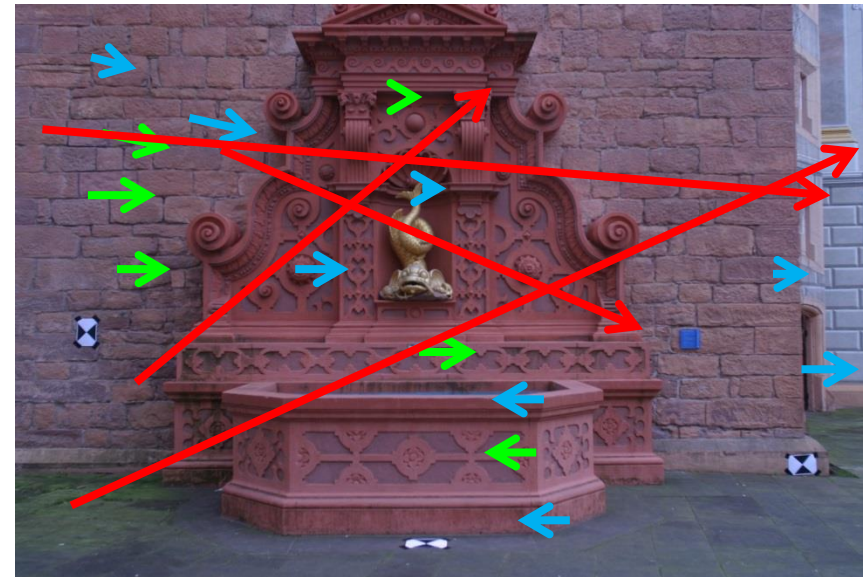


Image 1

# RANSAC iterations $k$ vs. $s$

$k$  increases exponentially with the number of points  $s$  estimate the model

Let's assume  $p = 99\%$  and  $\varepsilon = 50\%$  (fraction of outliers):

- **8-point RANSAC**

- $s = 8$  points (8-point algorithm)



$$k = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^8)} = 1177 \text{ iterations}$$

- **5-point RANSAC**

- $s = 5$  points (5-point algorithm)



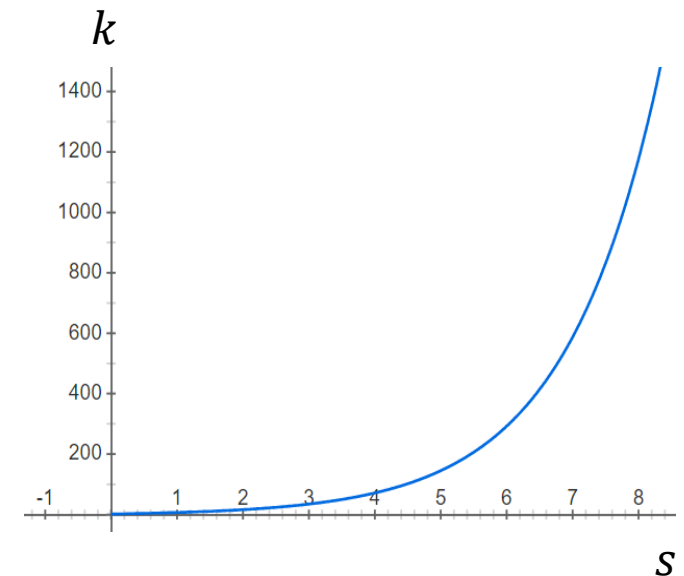
$$k = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^5)} = 145 \text{ iterations}$$

- **2-point RANSAC (e.g., line fitting)**

- $s = 2$  points

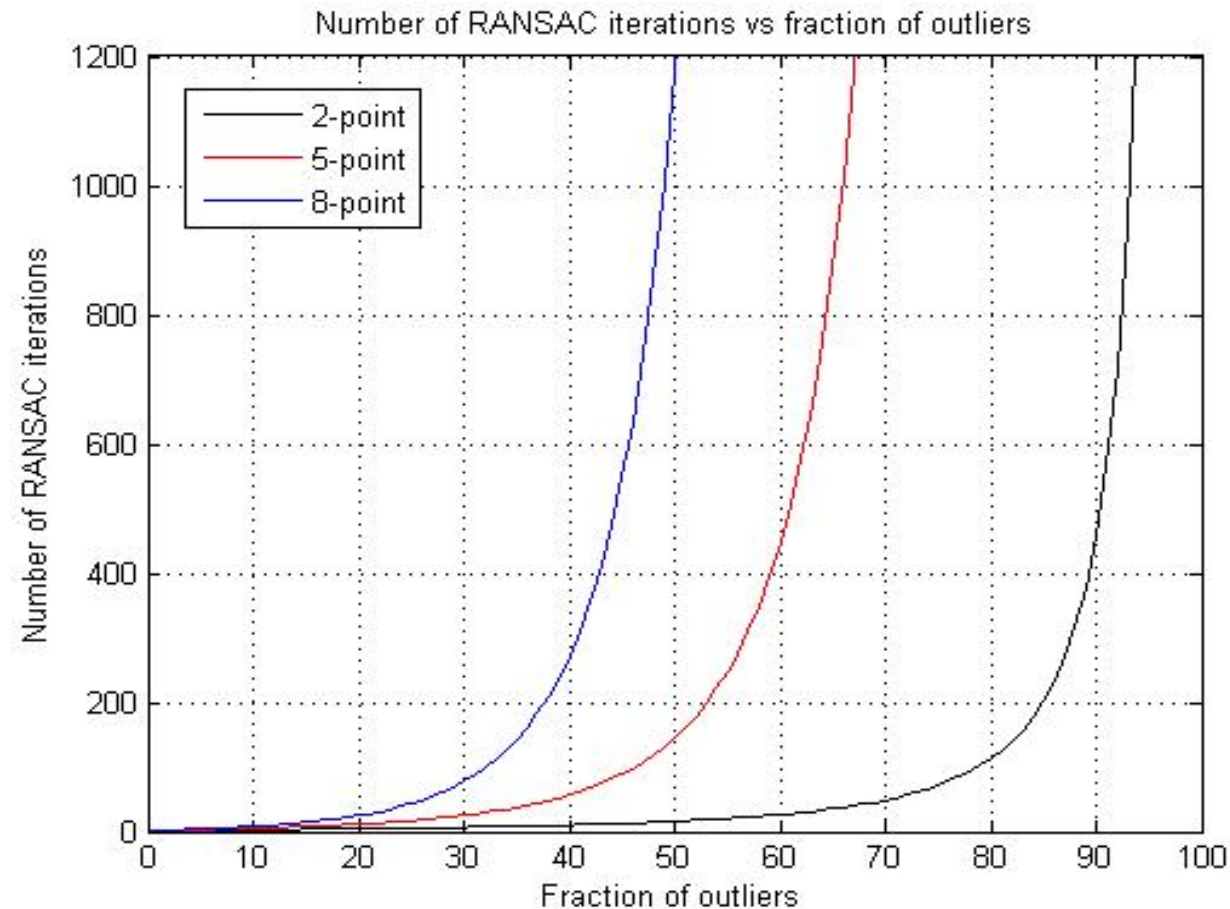


$$k = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^2)} = 16 \text{ iterations}$$



# RANSAC iterations $k$ vs. $\epsilon$

$k$  increases exponentially with the fraction of outliers  $\epsilon$ :



These plots were computed assuming  $p = 99\%$

# RANSAC iterations

- As observed,  $k$  is exponential with the number of points  $s$  necessary to estimate the model
- The **8-point algorithm** is extremely simple and was very successful; however, it requires more than **1177 iterations**
- Because of this, there has been a large interest by the research community in **using smaller motion parameterizations** (i.e., smaller  $s$ )
- The first efficient solution to the minimal-case solution (5-point algorithm) took almost a century (Kruppa 1913 → Nister 2004)
- The **5-point RANSAC** (Nister 2004) only requires **145 iterations**; however:
  - The **5-point algorithm** can return **up to 10 solutions of E (worst case scenario)**
  - The **8-point algorithm** only returns a **unique solution of E**

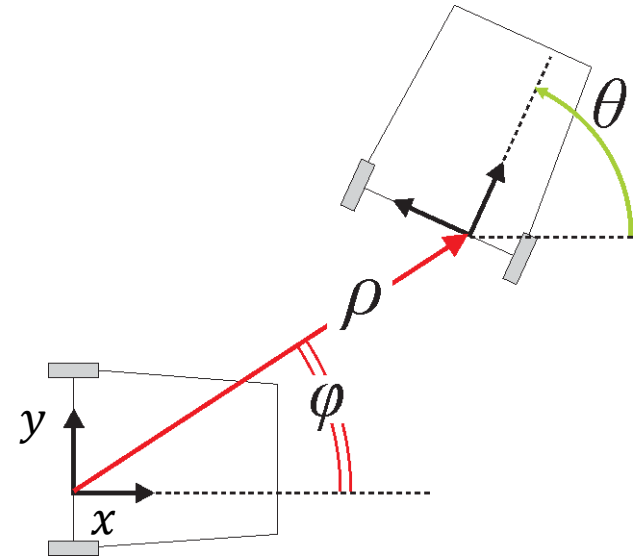
Can we use less than 5 points?

Yes, if you use motion constraints!

# Planar Motion

Planar motion is described by three parameters:  $\vartheta$ ,  $\varphi$ ,  $\rho$

$$R = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix} \quad T = \begin{bmatrix} \rho \cos \varphi \\ \rho \sin \varphi \\ 0 \end{bmatrix}$$



Let's compute the Epipolar Geometry

$$E = [T_x]R \quad \text{Essential matrix}$$

$$\bar{p}_2^T E \bar{p}_1 = 0 \quad \text{Epipolar constraint}$$

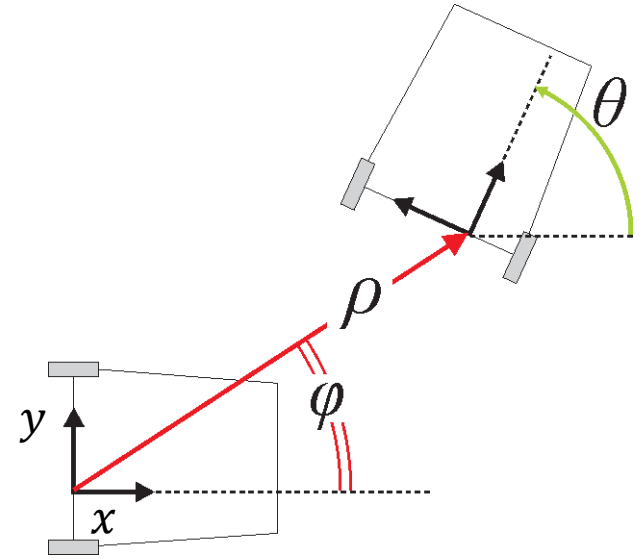
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Let's compute the Epipolar Geometry

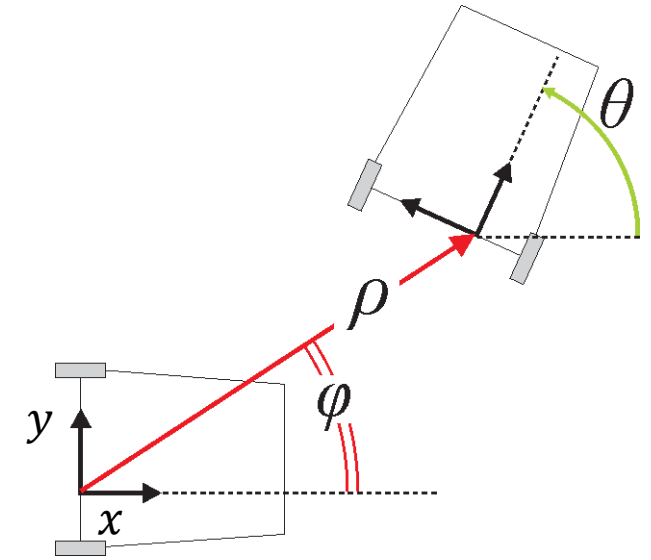
$$[T_x] = \begin{bmatrix} 0 & 0 & \rho \sin \varphi \\ 0 & 0 & -\rho \cos \varphi \\ -\rho \sin \varphi & \rho \cos \varphi & 0 \end{bmatrix}$$



# Planar Motion

Planar motion is described by three parameters:  $\vartheta$ ,  $\varphi$ ,  $\rho$

$$R = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix} \quad T = \begin{bmatrix} \rho \cos \varphi \\ \rho \sin \varphi \\ 0 \end{bmatrix}$$



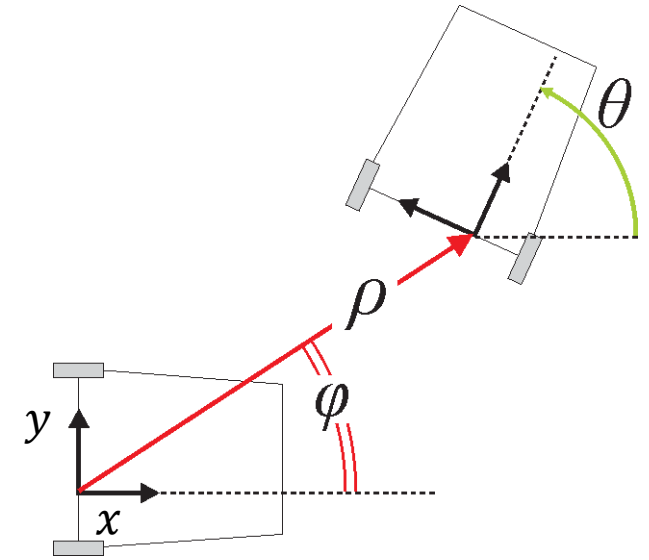
Let's compute the Epipolar Geometry

$$E = [T_x]R = \begin{bmatrix} 0 & 0 & \rho \sin \varphi \\ 0 & 0 & -\rho \cos \varphi \\ -\rho \sin \varphi & \rho \cos \varphi & 0 \end{bmatrix} \cdot \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

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Let's compute the Epipolar Geometry

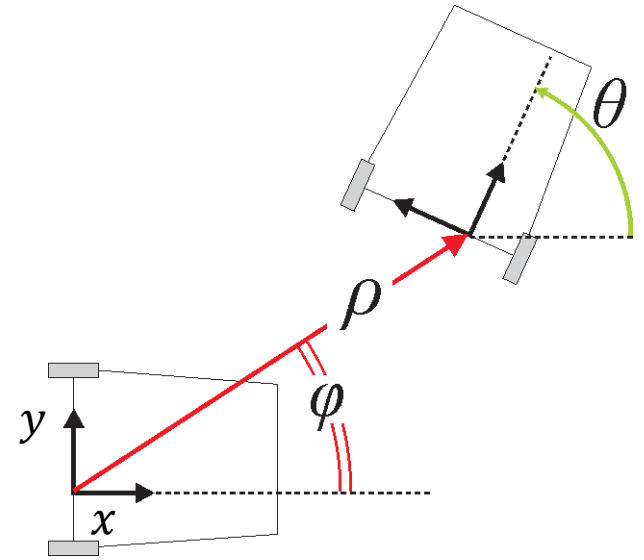
$$E = [T_x]R = \begin{bmatrix} 0 & 0 & \rho \sin(\varphi) \\ 0 & 0 & -\rho \cos(\varphi) \\ -\rho \sin(\varphi - \theta) & \rho \cos(\varphi - \theta) & 0 \end{bmatrix}$$



# Planar Motion

Planar motion is described by three parameters:  $\vartheta$ ,  $\varphi$ ,  $\rho$

$$R = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix} \quad T = \begin{bmatrix} \rho \cos \varphi \\ \rho \sin \varphi \\ 0 \end{bmatrix}$$



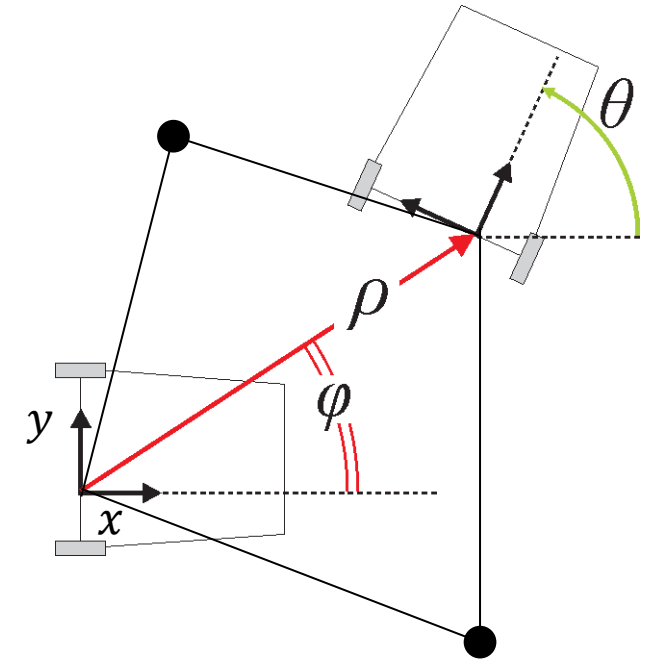
Let's compute the Epipolar Constraint:  $\bar{p}_2^T E \bar{p}_1 = 0$

$$-u_1 \sin(\phi - \theta) + v_1 \cos(\phi - \theta) + u_2 \sin(\phi) - v_2 \cos(\phi) = 0$$

# Planar Motion

Planar motion is described by three parameters:  $\vartheta$ ,  $\varphi$ ,  $\rho$

$$R = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix} \quad T = \begin{bmatrix} \rho \cos \varphi \\ \rho \sin \varphi \\ 0 \end{bmatrix}$$



Observe that  $\rho$  was cancelled out. Since only  $\theta$ ,  $\varphi$  can be determined and every point correspondence provides one scalar equation, then **2 point correspondences are sufficient** to estimate  $\theta$  and  $\varphi$

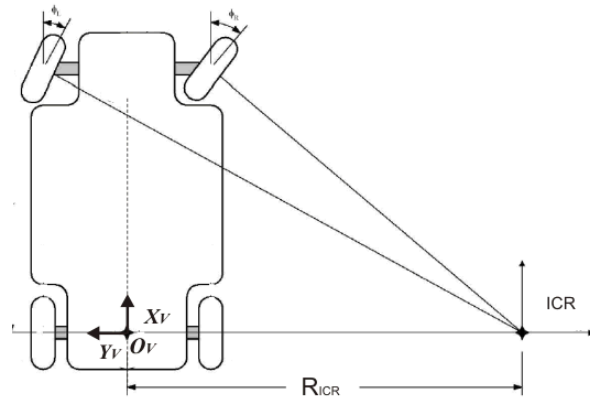
$$-u_1 \sin(\phi - \theta) + v_1 \cos(\phi - \theta) + u_2 \sin(\phi) - v_2 \cos(\phi) = 0$$

# Less than 2 points?

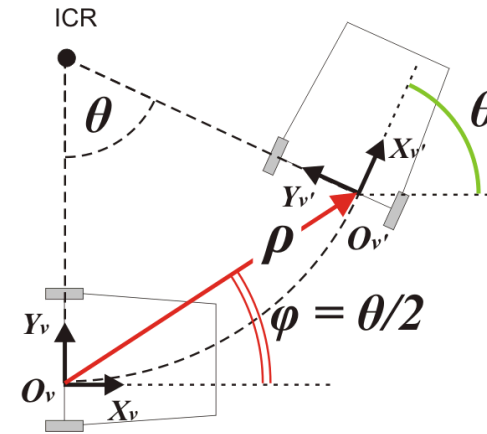
- Can we use less than 2 point correspondences?
  - Yes, if we exploit wheeled vehicles with **non-holonomic** constraints

# Planar & Circular Motion (e.g., cars)

Wheeled vehicles, like cars, follow locally-planar circular motion about the Instantaneous Center of Rotation (ICR)



Example of Ackerman steering principle

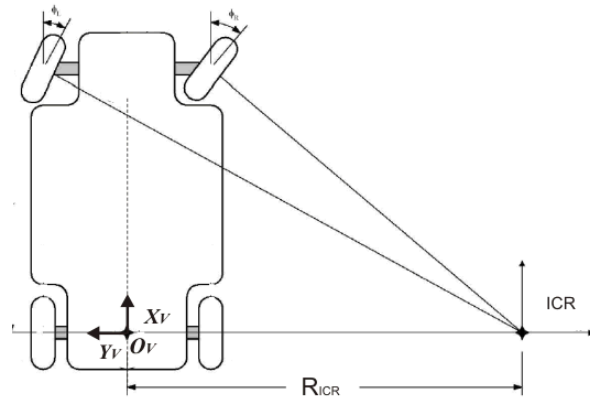


Locally-planar circular motion

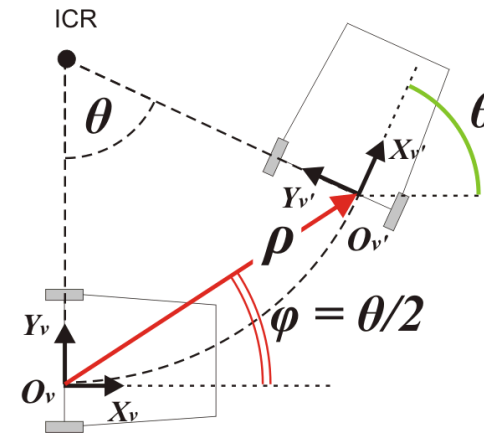


# Planar & Circular Motion (e.g., cars)

Wheeled vehicles, like cars, follow locally-planar circular motion about the Instantaneous Center of Rotation (ICR)



Example of Ackerman steering principle



Locally-planar circular motion

$\varphi = \theta/2 \Rightarrow$  only 1 DoF ( $\theta$ ); thus, **only 1 point correspondence** is sufficient [Scaramuzza, 2011]

**This is the smallest parameterization possible and results in the most efficient algorithm for removing outliers**

# Planar & Circular Motion (e.g., cars)

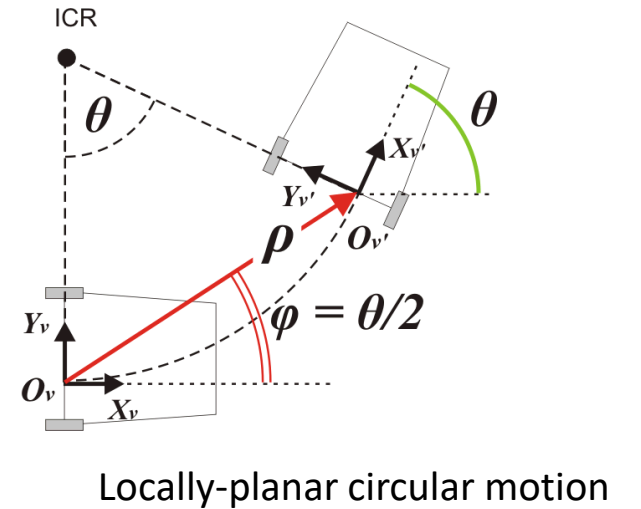
Let's compute the Epipolar Geometry

$$E = [T_{\times}]R \quad \text{Essential matrix}$$

$$\bar{p}_2^T E \bar{p}_1 = 0 \quad \text{Epipolar constraint}$$

$$R = \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

$$T = \begin{bmatrix} \rho \cos \frac{\theta}{2} \\ \rho \sin \frac{\theta}{2} \\ 0 \end{bmatrix}$$



# Planar & Circular Motion (e.g., cars)

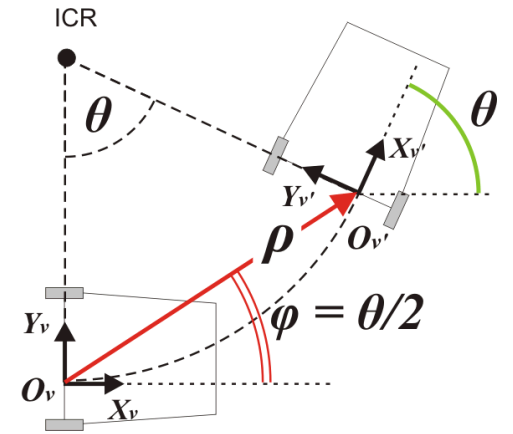
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Locally-planar circular motion

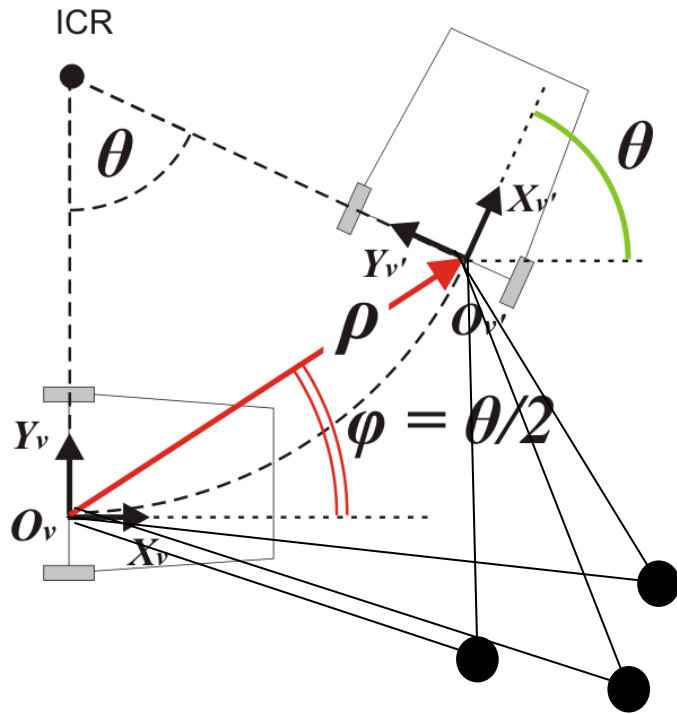
$$E = [T_{\times}]R = \begin{bmatrix} 0 & 0 & \rho \sin \frac{\theta}{2} \\ 0 & 0 & -\rho \cos \frac{\theta}{2} \\ -\rho \sin \frac{\theta}{2} & \rho \cos \frac{\theta}{2} & 0 \end{bmatrix} \cdot \begin{bmatrix} \cos \theta & -\sin \theta & 0 \\ \sin \theta & \cos \theta & 0 \\ 0 & 0 & 1 \end{bmatrix} = \begin{bmatrix} 0 & 0 & \rho \sin \frac{\theta}{2} \\ 0 & 0 & \rho \cos \frac{\theta}{2} \\ \rho \sin \frac{\theta}{2} & -\rho \cos \frac{\theta}{2} & 0 \end{bmatrix}$$

Notice that  $\rho$  can be cancelled out

$$\bar{p}_2^T E \bar{p}_1 = 0 \Rightarrow \sin\left(\frac{\theta}{2}\right) \cdot (u_2 + u_1) + \cos\left(\frac{\theta}{2}\right) \cdot (v_2 - v_1) = 0$$

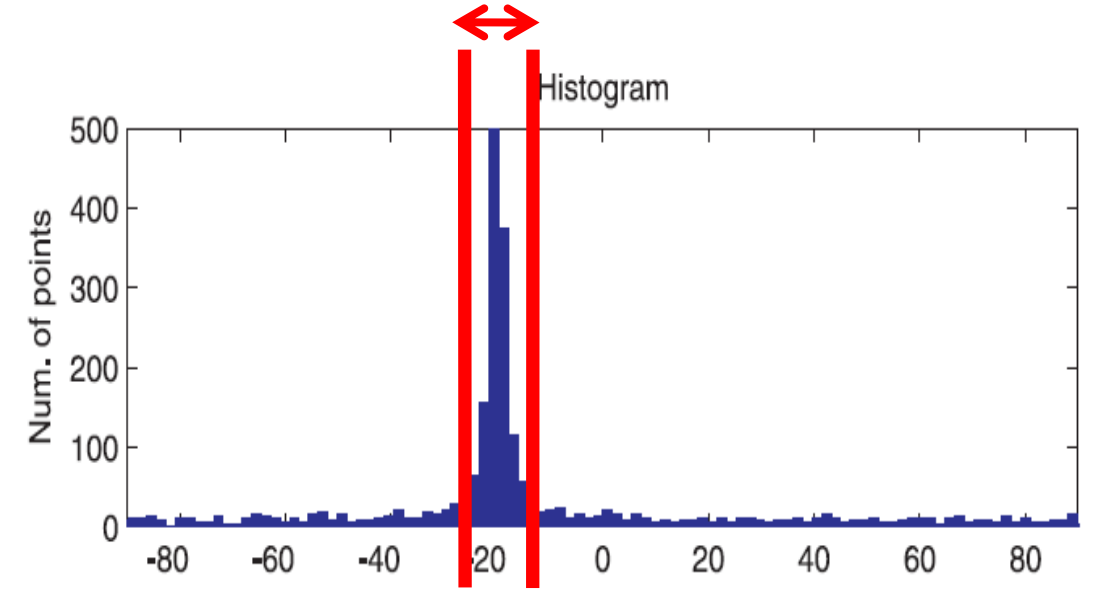
$$\theta = -2 \tan^{-1} \left( \frac{v_2 - v_1}{u_2 + u_1} \right)$$

# 1-Point RANSAC Algorithm



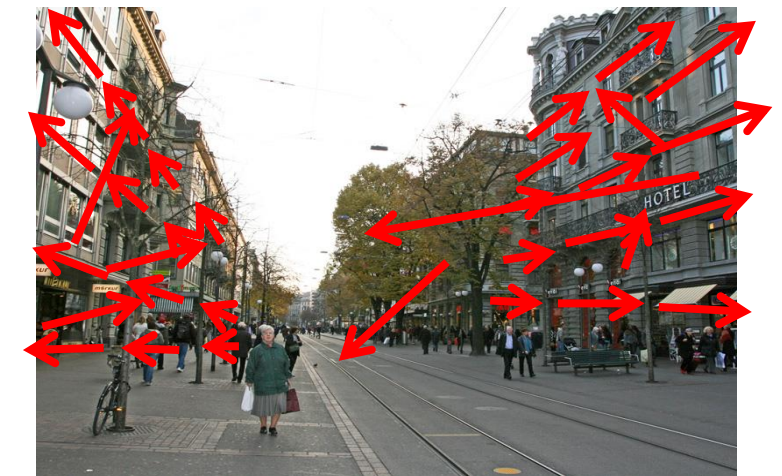
Compute  $\theta$  for every point correspondence

$$\theta = -2 \tan^{-1} \left( \frac{v_2 - v_1}{u_2 + u_1} \right)$$



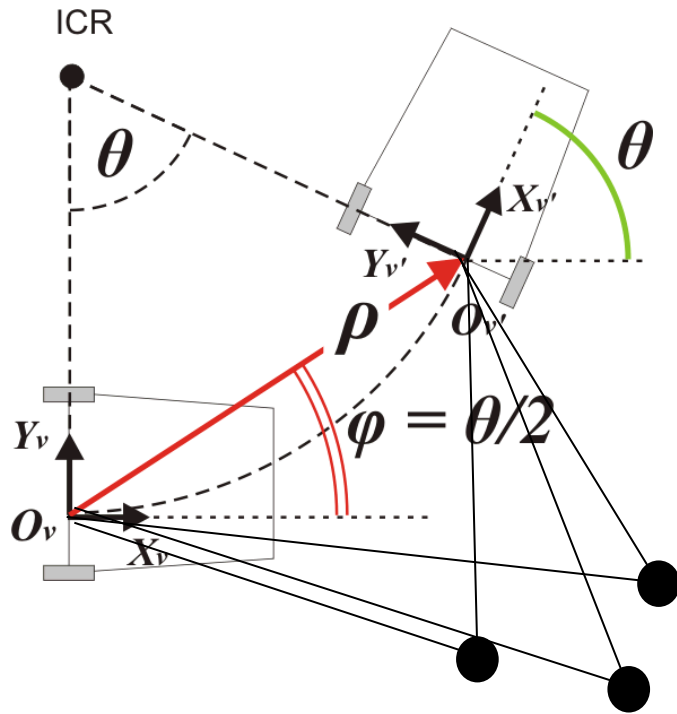
Only 1 iteration!  
The most efficient algorithm for removing outliers (<1ms)

1-Point RANSAC is ONLY used to find the inliers.  
Motion is then estimated from them in 6DOF



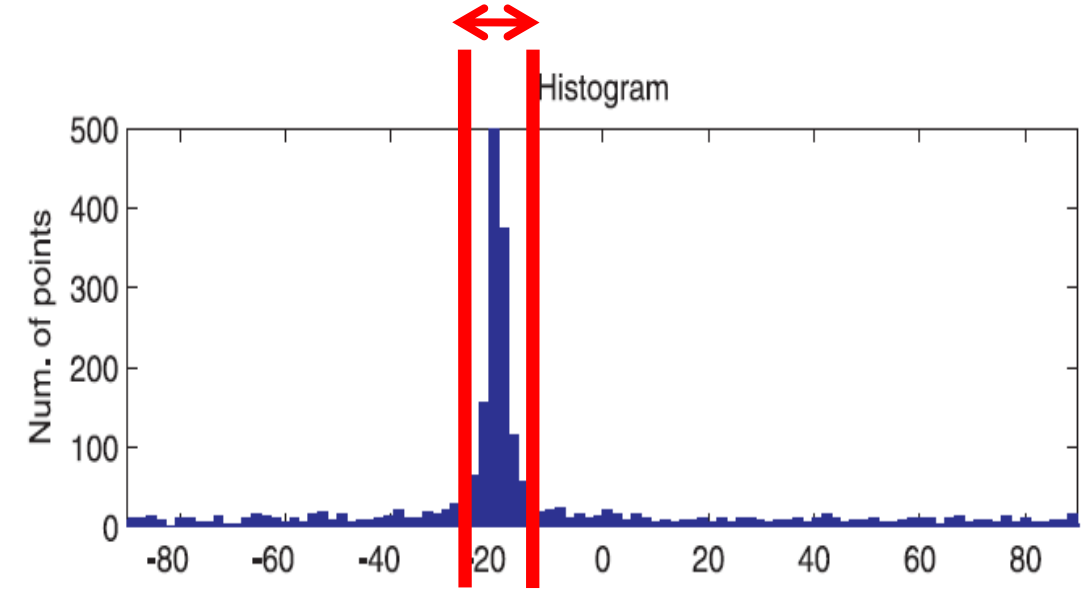


# 1-Point RANSAC Algorithm



Compute  $\vartheta$  for every point correspondence

$$\theta = -2 \tan^{-1} \left( \frac{v_2 - v_1}{u_2 + u_1} \right)$$

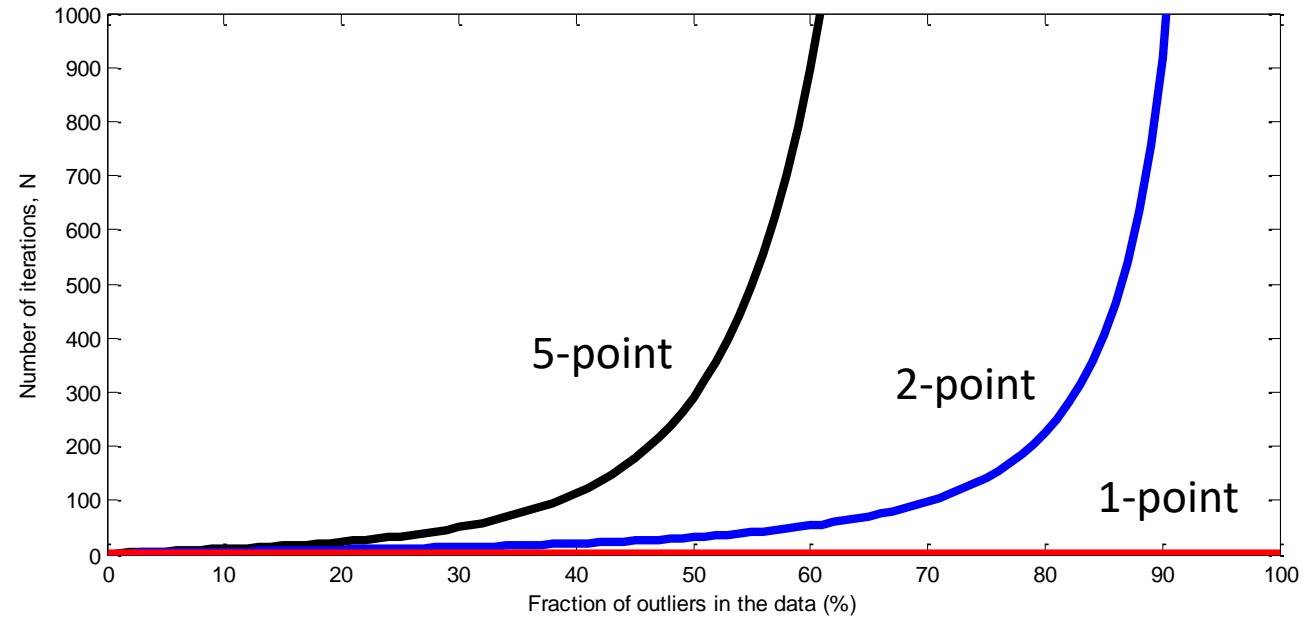


Only 1 iteration!  
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# Comparison of RANSAC algorithms

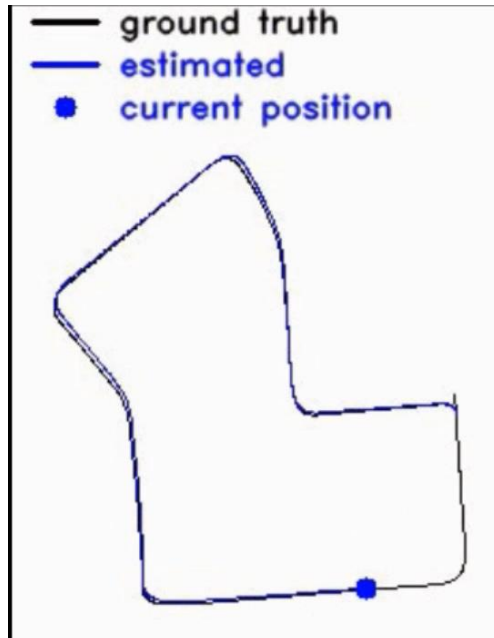


$$N = \frac{\log(1 - p)}{\log(1 - (1 - \varepsilon)^s)}$$

where we typically use  $p = 99\%$

	8-Point RANSAC [Longuet-Higgins'81]	5-Point RANSAC [Nister'04]	2-Point RANSAC [Ortin'01]	1-Point RANSAC [Scaramuzza'11]
Numb. of iterations	> 1177	>145	>16	=1

# Visual Odometry with 1-Point RANSAC



Work in different environments

Urban

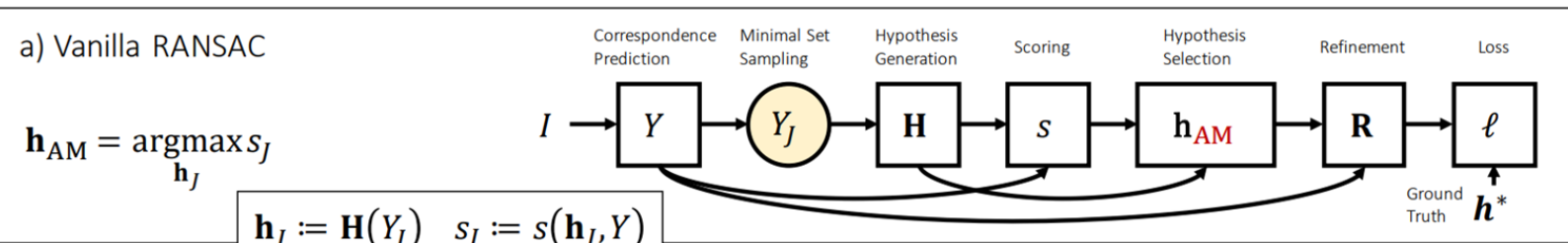


Latest and Greatest 😊

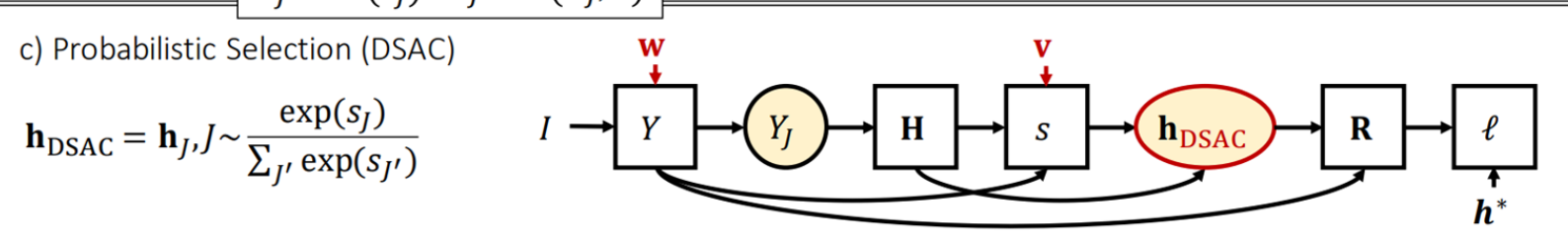
# Differentiable RANSAC

- RANSAC is not differentiable since it relies on selecting a hypothesis based on maximizing the number of inliers (i.e., argmax).
- DSAC shows how sample consensus can be used in a differentiable way
- This enables the use of sample consensus in a variety of learning tasks.

**Choose the best based on score**



**Randomly sample based on score**



# Deep Fundamental Matrix Estimation

- **Input:** two sets of noisy local features (coordinates + descriptors) contaminated by outliers
- **Output:** fundamental matrix
- **Idea:** solve a weighted homogeneous least-squares problem, where robust weights are estimated using deep networks
- **Robust:** handles extreme wide-baseline image pairs



Top-bottom as image-pair

Red: inlier correspondences

Blue: outlier correspondences

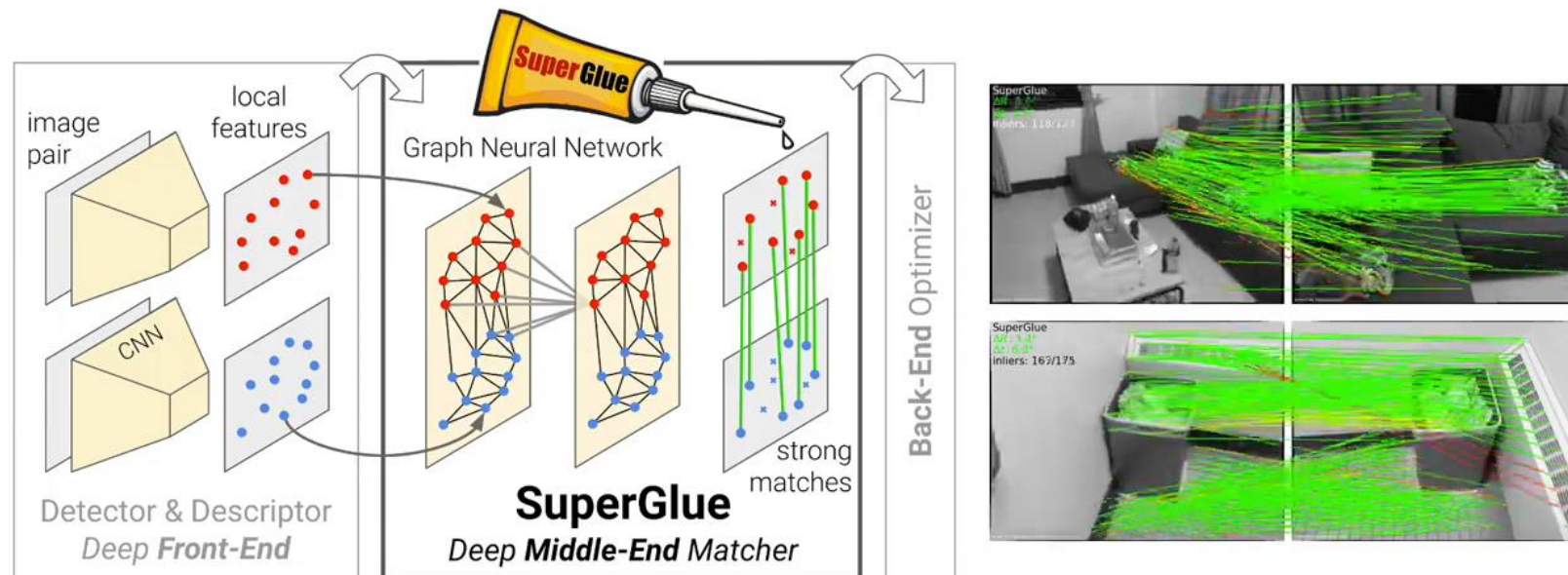
Epipolar lines

Green: estimated

Blue: ground-truth

# SuperGlue: Learning Feature Matching with Graph Neural Networks

- **Input:** two sets of noisy local features (coordinates + descriptors) contaminated by outliers
- **Output:** strong & outlier-free matches
- **Combines deep learning with classical optimization** (Graph Neural Networks, Attention, Optimal Transport)
- **Robust:** handles extreme wide-baseline image pairs



Sarlin, DeTone, Malisiewicz, Rabinovich, *SuperGlue: Learning Feature Matching with Graph Neural Networks*, International Conference on Computer Vision and Pattern Recognition (CVPR), 2020. [PDF](#). [Code](#).

# Outline

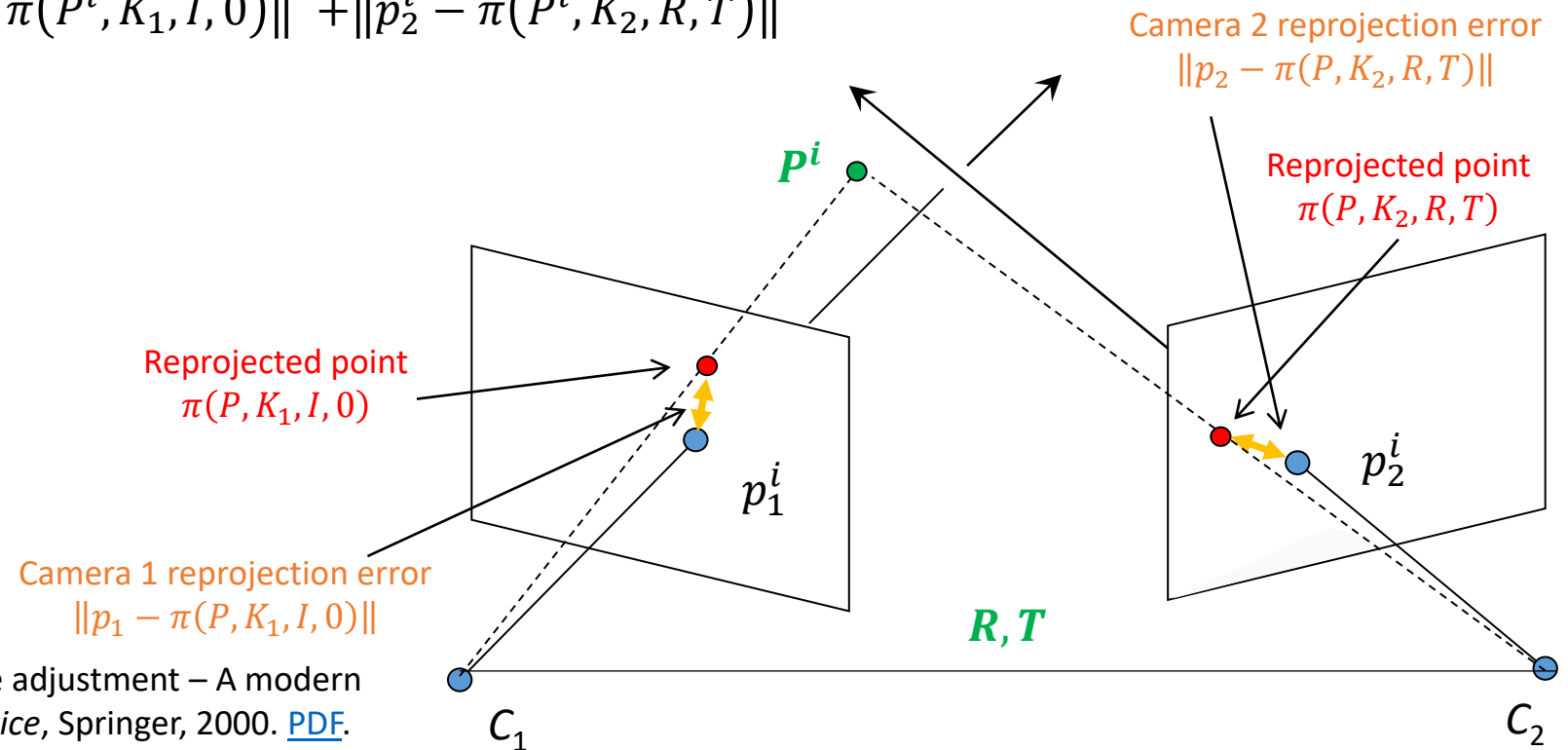
- Robust Structure from Motion
- Bundle Adjustment



# 2-View Bundle Adjustment (BA)

- **Non-linear, joint optimization of structure,  $P^i$ , and motion  $R, T$**
- Commonly used after least square estimation of  $R$  and  $T$  (e.g., after 8- or 5-point algorithm)
- Optimizes  $P^i, R, T$  by minimizing the **Sum of Squared Reprojection Errors**:

$$P^i, R, T = \operatorname{argmin}_{P^i, R, T} \sum_{i=1}^N \|p_1^i - \pi(P^i, K_1, I, 0)\|^2 + \|p_2^i - \pi(P^i, K_2, R, T)\|^2$$



# 2-View Bundle Adjustment (BA)

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## Good to know:

- Like in the formula, we typically assume the first camera as the world frame, but it's arbitrary
- Occasionally, the residual terms are weighted
- In order to not get stuck in local minima, the **initial values of  $P^i, R, T$  should be close to the optimum**
- Can be minimized using **Levenberg–Marquardt** (more robust than Gauss-Newton to local minima)
- **Can be modified to also optimize the intrinsic parameters**
- Implementation details in **Exercise 9**

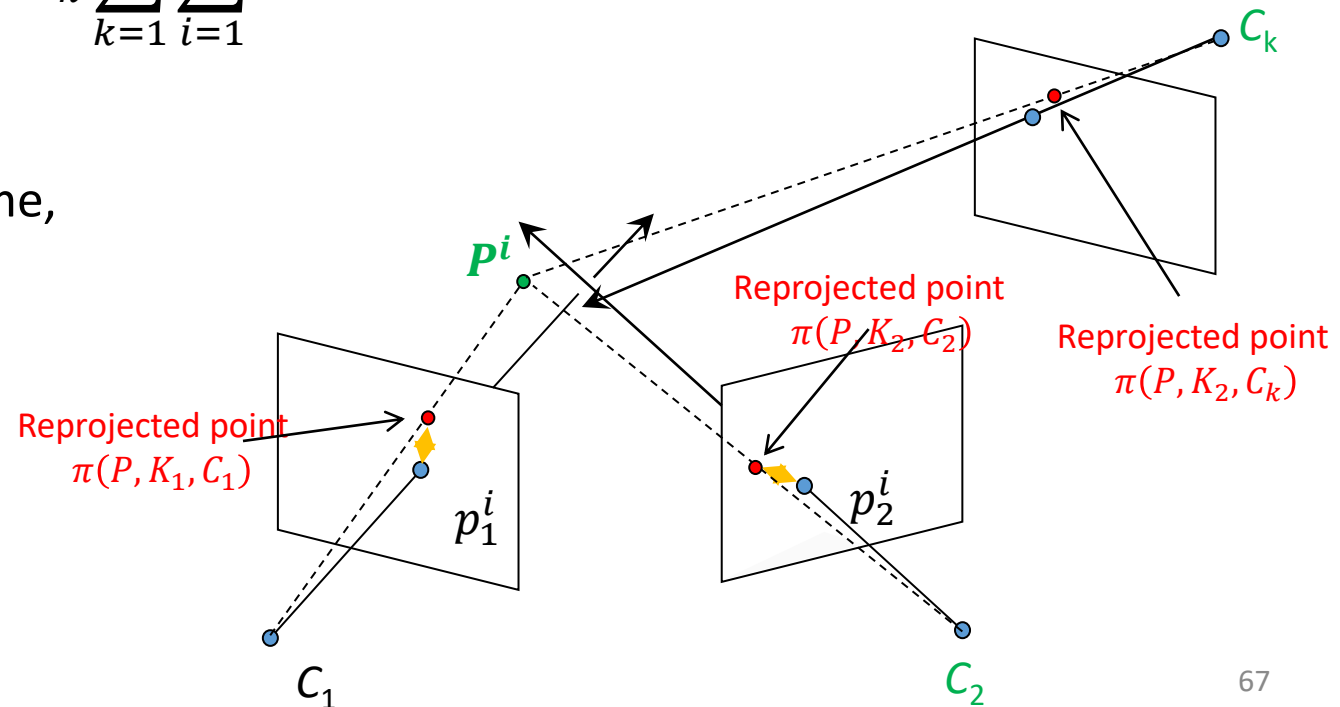
What is the key difference with the reprojection error minimization seen in previous lectures (Lecture 3, slide 21, and Lecture 7, slide 26)?

# $n$ -View Bundle Adjustment (BA)

- **Non-linear, joint optimization of structure,  $P^i$ , and camera poses  $C_1 = [I, 0], \dots, C_k = [R_k, T_k]$**
- **Minimizes the Sum of Squared Reprojection Errors across all views**

$$P^i, C_2, \dots, C_n = \operatorname{argmin}_{P^i, C_2, \dots, C_n} \sum_{k=1}^n \sum_{i=1}^N \|p_k^i - \pi(P^i, K_k, C_k)\|^2$$

- **NB:** we assume the first camera as the world frame, that's why  $C_1 = [I, 0]$



# Huber and Tukey Norms

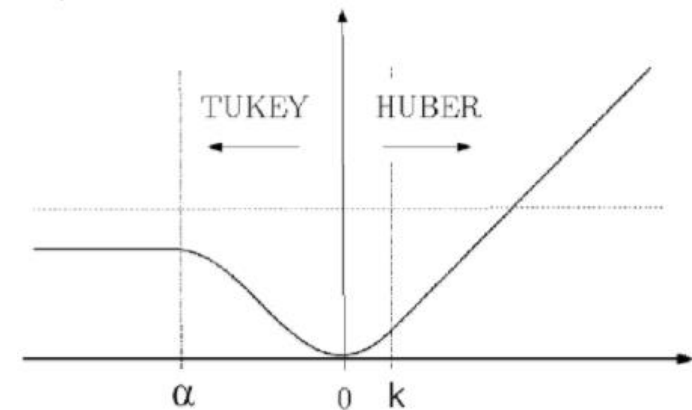
- To prevent that large reprojection errors can negatively impact the optimization, a more robust norm  $\rho(\cdot)$  is used instead of the  $L_2$ :

$$P^i, C_2, \dots, C_n = \underset{P^i, C_2, \dots, C_n}{\operatorname{argmin}} \sum_{k=1}^n \sum_{i=1}^N \rho(p_k^i - \pi(P^i, K_k, C_k))$$

- $\rho(\cdot)$  is a robust cost function (**Huber or Tukey**) to alleviate the contribution of wrong matches:

- Huber norm:** 
$$\rho(x) = \begin{cases} x^2 & \text{if } |x| \leq k \\ k(2|x| - k) & \text{if } |x| \geq k \end{cases}$$

- Tukey norm:** 
$$\rho(x) = \begin{cases} \alpha^2 & \text{if } |x| \geq \alpha \\ \alpha^2 \left( 1 - \left( 1 - \left( \frac{x}{\alpha} \right)^2 \right)^3 \right) & \text{if } |x| \leq \alpha \end{cases}$$



These formulas are not asked at the exam but their plots and meaning is asked 😊

# Things to remember

- EM algorithm
- RANSAC algorithm and its application to SFM
- 8 vs 5 vs 1 point RANSAC, pros and cons
- Bundle Adjustment

# Reading

- CH. 8.1.4, 8.3.1, 11.3 of Szeliski book, 2<sup>nd</sup> edition
- Ch. 14.2 of Corke book

# Understanding Check

Are you able to answer the following questions?

- What are the causes of outliers?
- What effects may outliers have on VO?
- How does EM work? What are the issues?
- Why do we need RANSAC?
- What is the theoretical maximum number of combinations to explore?
- After how many iterations can RANSAC be stopped to guarantee a given success probability?
- What is the trend of RANSAC vs. iterations, vs. the fraction of outliers, vs. the number of points to estimate the model?
- How do we apply RANSAC to the 8-point algorithm, DLT, P3P?
- How can we reduce the number of RANSAC iterations for the SFM problem? (1- and 2-point RANSAC)
- Bundle Adjustment. Mathematical expression and illustration. Tukey and Huber norms.