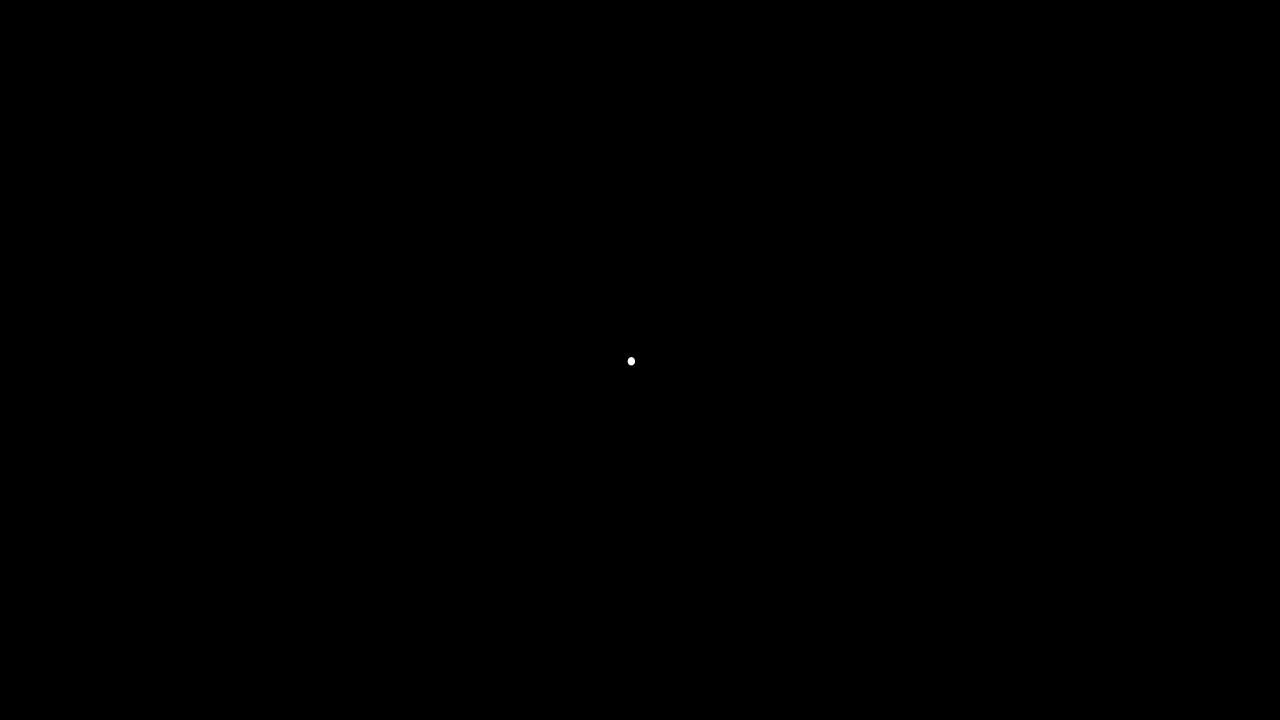
VAN course Lesson 4

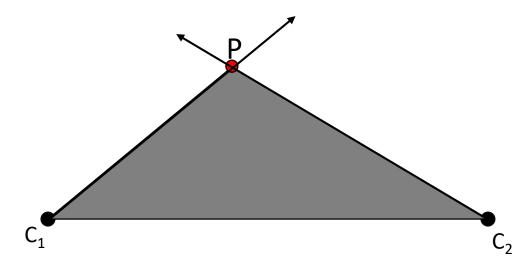
Dr. Refael Vivanti refael.vivanti@mail.huji.ac.il

Today's topics

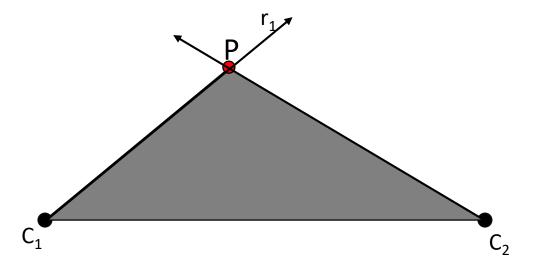
- Epipolar geometry
- Epipole
- Fundamental Matrix Calculation
- Rectification
- RANSAC



• The relations between views as appeared in the image

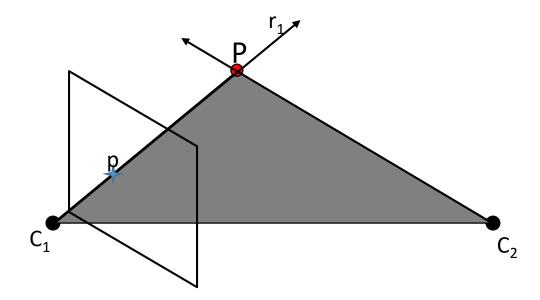


- The relations between views as appeared in the image
- In 3D: Ray r₁ intersects P



• The relations between views as appeared in the image

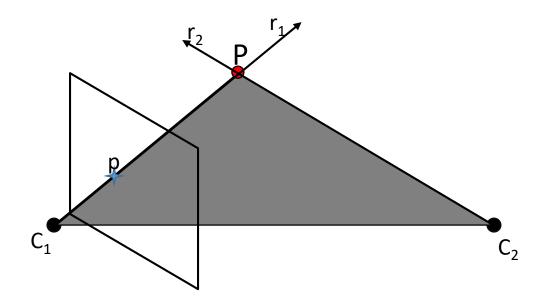
• In 3D: Ray r₁ intersects P In image: P is projected to p



• The relations between views as appeared in the image

• In 3D: Ray r₁ intersects P In image: P is projected to p

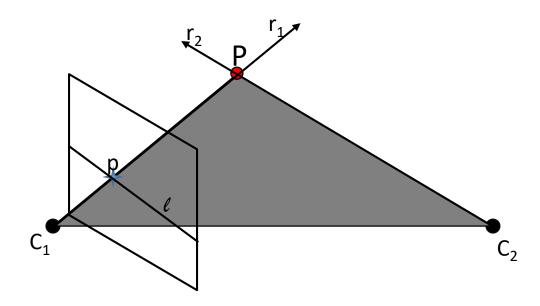
• In 3D: Ray r₂ intersect P



• The relations between views as appeared in the image

• In 3D: Ray r₁ intersects P In image: P is projected to p

• In 3D: Ray r_2 intersect P In Image: r_2 is projected to line ℓ



• The relations between views as appeared in the image

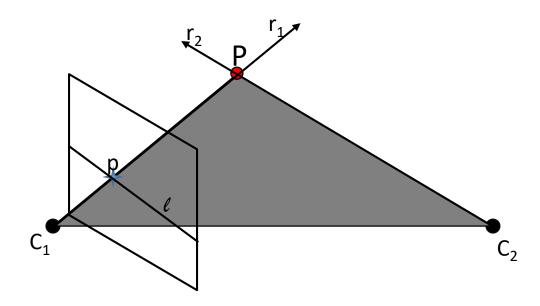
• In 3D: Ray r₁ intersects P In i

• In 3D: Ray r₂ intersect P

 $ullet \ell$ is the **epipolar line**

In image: P is projected to p

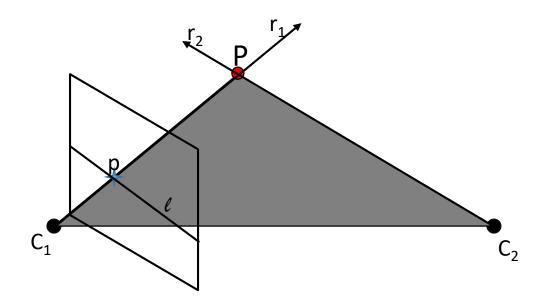
In Image: r_2 is projected to line ℓ



- The relations between views as appeared in the image
- In 3D: Ray r₁ intersects P
- In 3D: Ray r₂ intersect P
- $ullet \ell$ is the **epipolar line**
- $ullet \ell$ intersects p in image 1

In image: P is projected to p

In Image: r_2 is projected to line ℓ



• The relations between views as appeared in the image

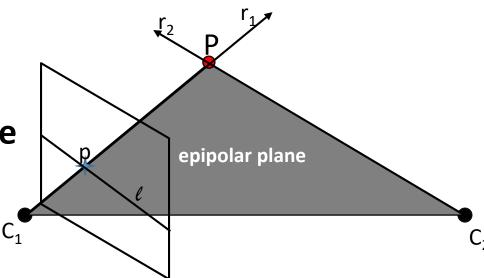
• In 3D: Ray r₁ intersects P In image: P is projected to p

• In 3D: Ray r_2 intersect P In Image: r_2 is projected to line ℓ

ullet ℓ is the **epipolar line**

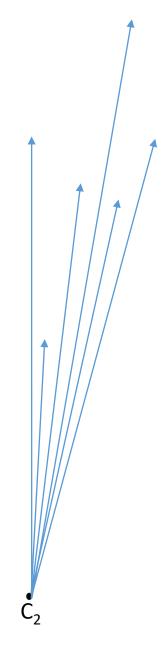
 $ullet \ell$ intersects p in image 1

• C₁ PC₂ defines the **epipolar plane**

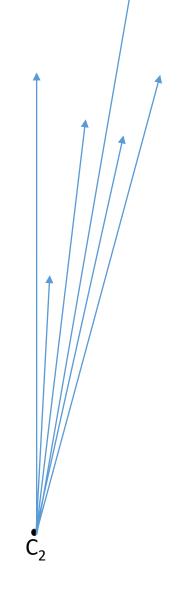


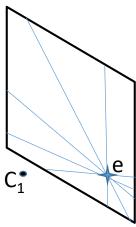
- What epipolar line is good for?
- If we search for a match for p₁
 - It will be on the epipolar line ℓ
- If we suspect the match is wrong
 - We can decide it is an outlier if it's not on ℓ

• In 3D: All 3D rays coming from C₂ create a pencil

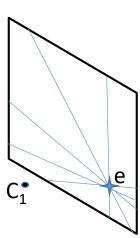


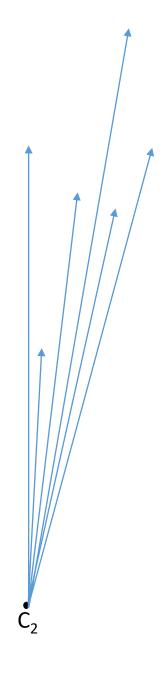
- In 3D: All 3D rays coming from C₂ create a pencil
- In image: all epipolar lines intersect at point e



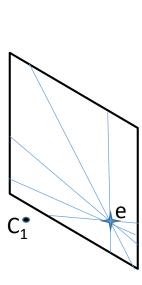


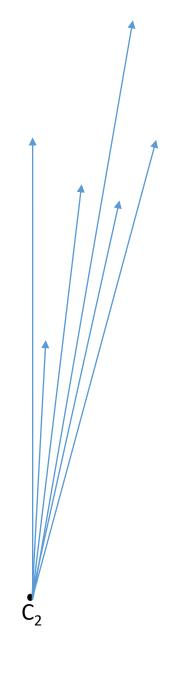
- In 3D: All 3D rays coming from C₂ create a pencil
- In image: all epipolar lines intersect at point e
- e is the **epipole**



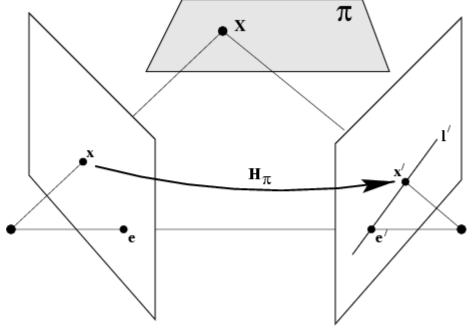


- In 3D: All 3D rays coming from C₂ create a pencil
- In image: all epipolar lines intersect at point e
- e is the **epipole**
- e is the projection of c₂ 3D location





geometric derivation of F:



$$x' = H_{\pi}x$$

$$1' = e' \times x' = [e']_{\times} H_{\pi}x = Fx$$

- If we don't know K_1 , K_2 , R, or t, can we still estimate F?
- Yes, given enough correspondences.

Many algorithms:

- Linear (the normalized 8-point algorithm)
- Minimal (7-point)
- Robust (RANSAC)
- Non-linear refinement (MLE, Algebraic minimization)
- We use 8-point algorithm
 - Although it's inaccurate
 - Because it's fast

The fundamental matrix F is defined by

$$\mathbf{x'}^{\mathsf{T}}\mathbf{F}\mathbf{x} = \mathbf{0}$$

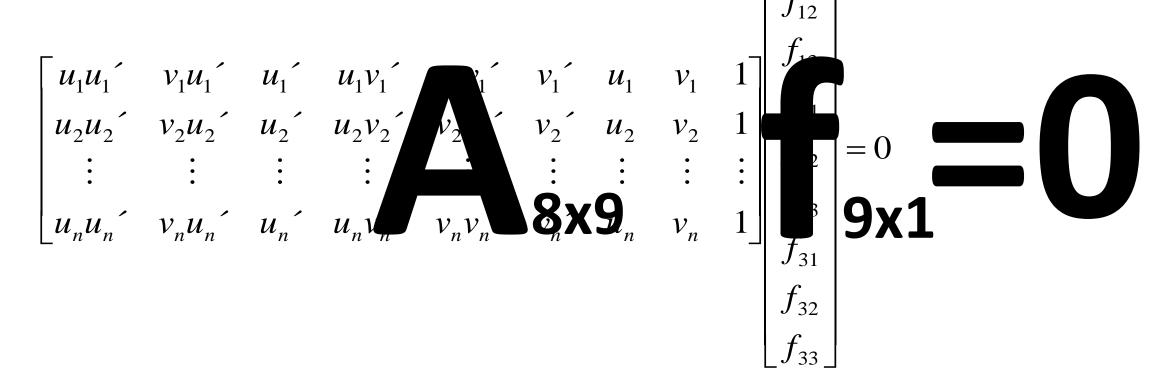
for any pair of matches x and x' in two images.

• Let $x=(u,v,1)^T$ and $x'=(u',v',1)^T$, each match gives a linear equation

$$\mathbf{F} = \begin{bmatrix} f_{11} & f_{12} & f_{13} \\ f_{21} & f_{22} & f_{23} \\ f_{31} & f_{32} & f_{33} \end{bmatrix}$$

$$uu'f_{11} + vu'f_{12} + u'f_{13} + uv'f_{21} + vv'f_{22} + v'f_{23} + uf_{31} + vf_{32} + f_{33} = 0$$

8-point algorithm
$$\begin{bmatrix} u_{1}u_{1}' & v_{1}u_{1}' & u_{1}' & u_{1}v_{1}' & v_{1}v_{1}' & v_{1}' & u_{1} & v_{1} & 1 \\ u_{2}u_{2}' & v_{2}u_{2}' & u_{2}' & u_{2}v_{2}' & v_{2}v_{2}' & v_{2}' & u_{2} & v_{2} & 1 \\ \vdots & \vdots \\ u_{n}u_{n}' & v_{n}u_{n}' & u_{n}' & u_{n}v_{n}' & v_{n}v_{n}' & v_{n}' & u_{n} & v_{n} & 1 \end{bmatrix} \begin{bmatrix} f_{11} \\ f_{12} \\ f_{13} \\ f_{21} \\ f_{22} \\ f_{23} \\ f_{31} \\ f_{32} \\ f_{33} \end{bmatrix} = 0$$



We solve it as before, using SVD decomposition:

$$Af = 0$$

$$A = U\Sigma V^{T}$$

$$f = V^{T}_{N}$$

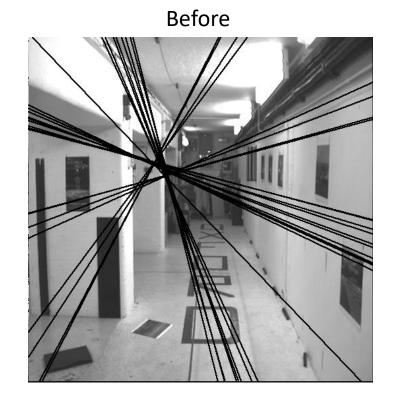
- We can use more than 8 points. M>N=8
 - But now, instead of solving $\mathbf{Af} = 0$, we seek \mathbf{f} to minimize $\|\mathbf{Af}\|$, least eigenvector of $\mathbf{A}^T\mathbf{A}$.
 - Still, we take $f = V_N^T$

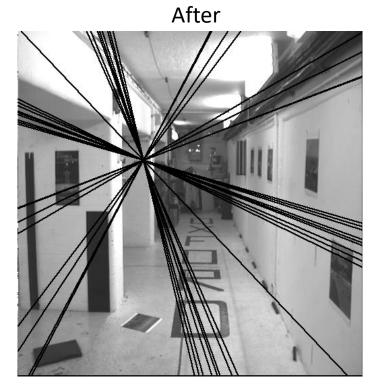
- Problem: F should have rank 2. It doesn't.
- To enforce that **F** is of rank 2, F is replaced by F' that minimizes $\|\mathbf{F} \mathbf{F}'\|$ subject to the rank constraint.

• This too is achieved by SVD. Let $\mathbf{F} = \mathbf{U} \Sigma \mathbf{V}^{\mathrm{T}}$, where

$$\Sigma = \begin{bmatrix} \sigma_1 & 0 & 0 \\ 0 & \sigma_2 & 0 \\ 0 & 0 & \sigma_3 \end{bmatrix} \qquad \Sigma' = \begin{bmatrix} \sigma_1 & 0 & 0 \\ 0 & \sigma_2 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

then $\mathbf{F'} = \mathbf{U} \mathbf{\Sigma'} \mathbf{V}^{\mathrm{T}}$ is the solution.

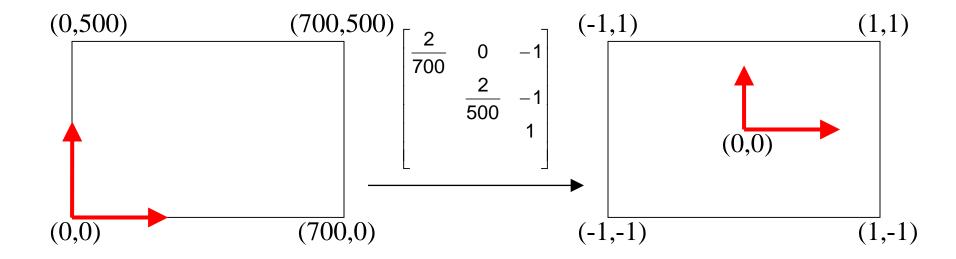


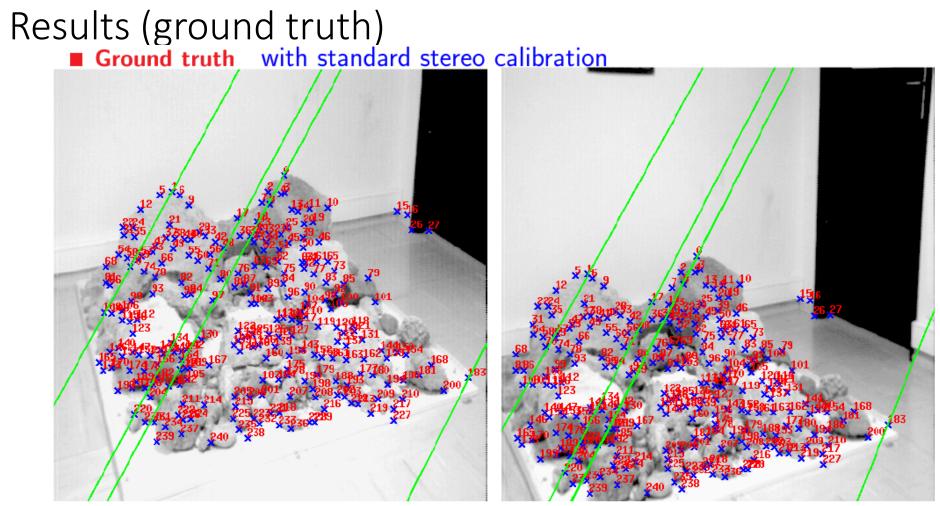




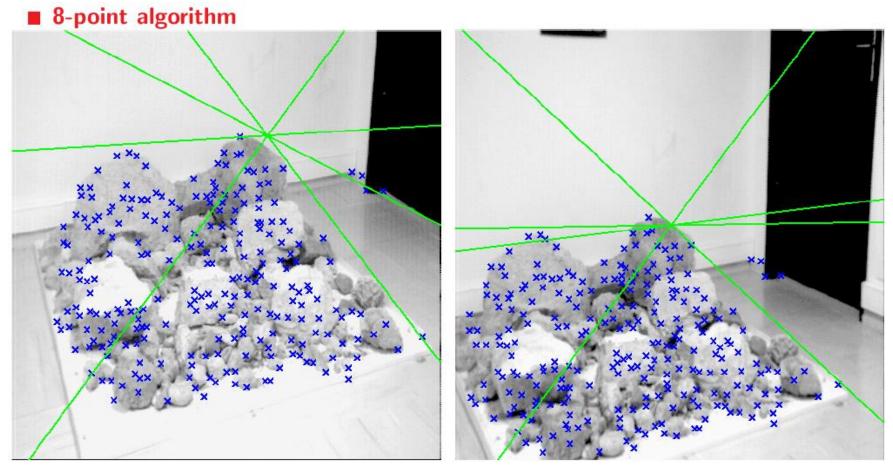
Orders of magnitude difference between column of data matrix → least-squares yields poor results

- normalized least squares yields good results
- Transform image to \sim [-1,1]x[-1,1]

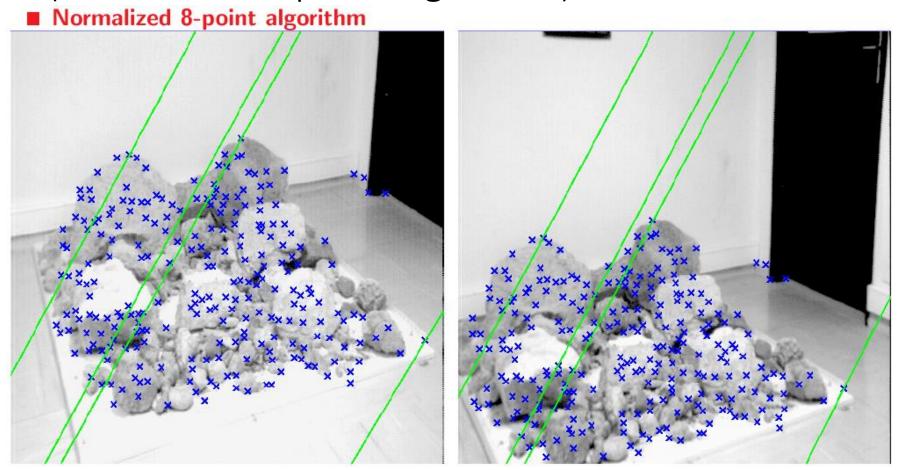




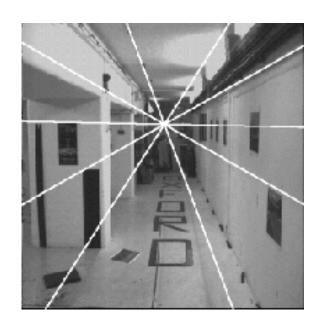
8-point algorithm Results (8-point algorithm)



Results (normalized 8-point algorithm)

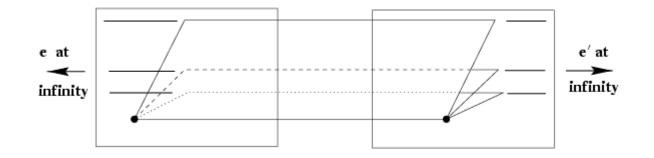


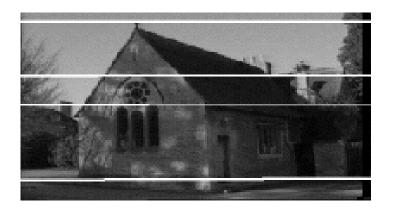
Example: forward motion

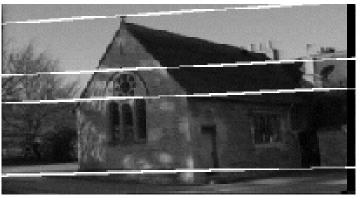




Example: motion parallel with image plane







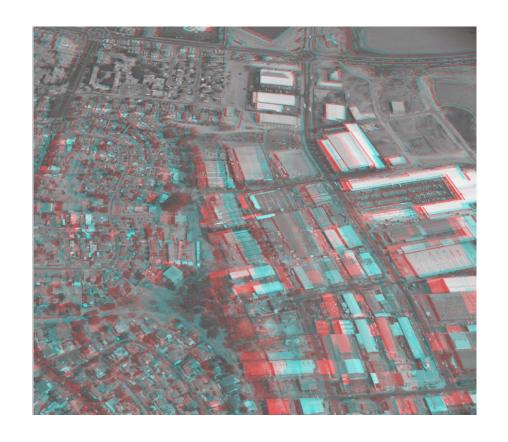
• In stereo rectification

We wish all epipolar lines to be:

$$F(x,y,1)^{T} \rightarrow (0, 1, -y)$$

So F is from the shape:

$$[t]_{x} = \begin{bmatrix} 1 \\ 0 \\ 0 \end{bmatrix}_{x} \begin{bmatrix} 1 \\ -1 \end{bmatrix}$$



- RANdom SAmple Consensus
- Problem:
 - All inliers obey some model
 - But there are some unknown outliers.
- Example: inliers are on a curve
- Chicken and egg situation:
 - If we had the curve, we could spot the outliers
 - If we knew the inliers, we could estimate the curve
- Key to solution:
 - The model can be estimated using a small set





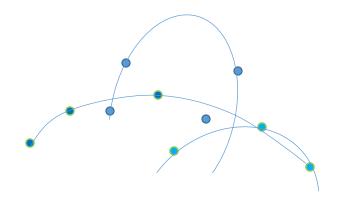
- Algorithm:
 - 1. Repeat:
 - 1. Sample a minimal set
 - 2. Estimate a model
 - 3. Check how many points obey the model
 - 2. Choose model with maximal #points
 - 3. Repeat:
 - 1. Estimate model from all inliers
 - 2. Calc inliers of new model
- Output: inliers, outliers, and model

- Example:
- #inliers = 3. max #inliers = 3
- #inliers = 5. max #inliers = 5
- #inliers = 3. max #inliers = 5

•

•

• Output: #inliers = 5



- When to stop the first loop of RANSAC?
- Goal: one sample that will have only inliers, with high prob p.
- Prob of being an outlier: ϵ
- P(being an inlier) = 1ϵ
- P(all inliers-sample) = $(1 \epsilon)^s$
- P(bad sample)= $1-(1-\epsilon)^{s}$
- P(All samples are bad) = $(1 (1 \epsilon)^s)^I$
- We wish it to be small: $(1-(1-\varepsilon)^s)^I < 1-p$

$$\log(1-(1-\varepsilon)^s)^I < \log(1-p)$$

$$I\log(1-(1-\varepsilon)^s) < \log(1-p)$$

$$I > \log(1-p)/\log(1-(1-\varepsilon)^s)$$

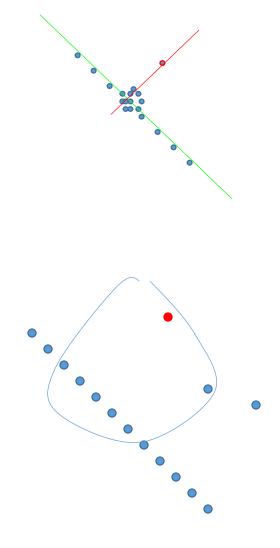
This can be really high:

$\mathbf{s} \setminus \mathbf{\epsilon}$	25%	50%	60%	70%	80%	85%
2	6	16	26	49	113	202
3	8	34	70	168	573	1362
7	33	588	2808	21055	2.5E05	2.6E06

- What if we don't know ε ?
- We can estimate it online:
 - We calc #inliers at each sample
 - This gives an ever-decreasing upper-bound on ϵ
 - Hence the needed iteration number I is also decreasing

- Which models are used with RANSAC?
- 2D points matching:
 - Fundamental matrix
 - Homography transformation
 - Essential Matrix
 - Trifocal Tensor
- 3D points:
 - Point cloud registration
 - Perspective-n-Point (PNP)
 - Plane fitting
 - Curve fitting

- Limitations of RANSAC with FM:
 - Efficiency: unknown
 - because outliers ratio ε in unknown
 - Accuracy
 - Even good sample may give a bad model
 - Sensitive to inlier threshold
 - Degeneracy
 - The plain+paralax problem
 - Many tricks and extensions:
 - PROSAC
 - USAC



Thanks

3D points cloud registration

- In the matched case
 - Each 3D point in X have a correspondence in Y)

$$y_{3x1} = R_{3x3}x_{3x1} + t_{3x1}$$

$$\widetilde{Y} = Y - \overline{Y}, \widetilde{X} = X - \overline{X}$$

$$t = \overline{X} - \overline{Y}$$

$$\widetilde{Y}_{3xN} = R_{3x3}\widetilde{X}_{3xN}$$

$$U_{3x3}\Sigma_{3x3}V_{3x3}^{T} = SVD(X^{T}_{Nx3}Y_{Nx3})$$

$$R = V^{T}U$$

3D points cloud registration

- In the not-matched case
- The ICP Iterative Closest Point algorithm
 - Repeat until convergence:
 - Find temporary matches:
 - For each point in X:
 - Set the closest point in Y to match it
 - Calculate R and t using the SVD algorithm above
 - May use RANSAC for outlier removal
 - Transform X using R and t

