

# Feature matching and model fitting

(Week-07 Lectures 29-33)



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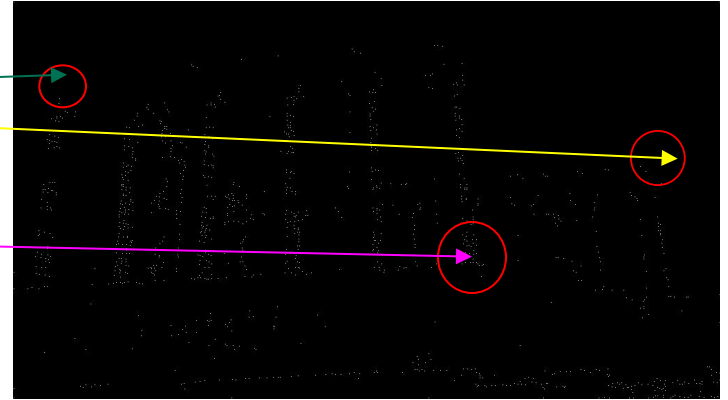
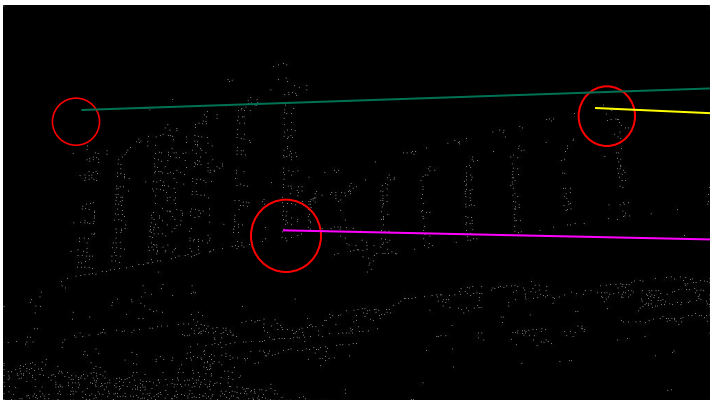
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# Can you get the 3-D structure of the scene?

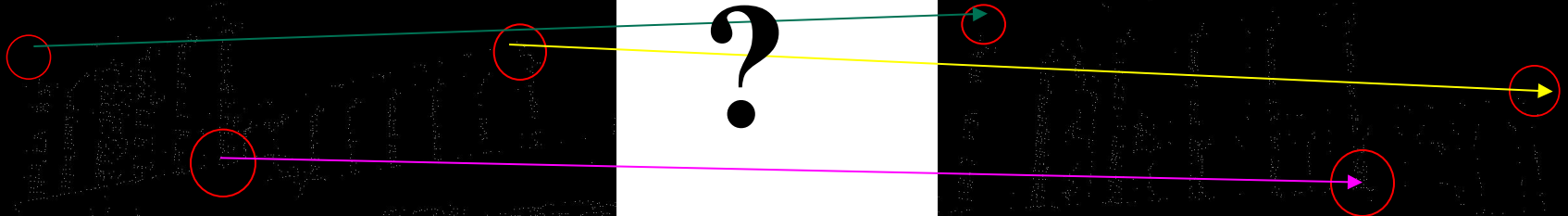


- Get a set of pairs of corresponding points.
- Compute fundamental matrix.
- Derive camera matrices.
- Solve for 3-D coordinates of scene points for each pair of corresponding points.

# Feature matching



?





# Matching with Features

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- Detect feature points in both images.
- Describe them by local statistics.
- Find corresponding pairs (Matching).



# Matching key points

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- Representation of a key-point by a feature vector.
  - e.g.  $[f_0 f_1 \dots f_n]^T$
- Use distance functions / similarity measures.

- $L_1$  norm

$$L_1(\vec{f}, \vec{g}) = \sum_{i=0}^n |f_i - g_i|$$

- $L_2$  norm

$$L_2(\vec{f}, \vec{g}) = \left( \sum_{i=0}^n |f_i - g_i|^2 \right)^{\frac{1}{2}}$$

- $L_p$  norm

$$L_p(\vec{f}, \vec{g}) = \left( \sum_{i=0}^n |f_i - g_i|^p \right)^{\frac{1}{p}}$$



# Weighted Distance function

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- More weights to reliable components.

$$d_w(\mathbf{f}, \mathbf{g}) = \sqrt{(\mathbf{f} - \mathbf{g})^T A (\mathbf{f} - \mathbf{g})}$$

$\mathbf{f}, \mathbf{g}$ : Column vectors of dimension  $n$ .

$A$ : A +ve semidefinite matrix

- Symmetric.
- $\mathbf{v}^T A \mathbf{v} \geq 0$  for all  $\mathbf{v}$ .
- Typical example:  $\text{Diag}(w_0, w_2, \dots, w_{n-1})$ ,  $w_i \geq 0$

$$d_w(\vec{f}, \vec{g}) = \sqrt{\sum_{i=0}^{n-1} w_i (f_i - g_i)^2}$$



# A few other similarity measures

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- Normalized Cross Correlation.

$$\begin{aligned}\rho(\vec{f}, \vec{g}) &= \frac{\text{cov}(f, g)}{s.d.(f) \times s.d.(g)} \\ &= \frac{\frac{1}{n} \sum_{i=0}^{n-1} (f_i - \bar{f})(g_i - \bar{g})}{\sqrt{\frac{1}{n} \sum_{i=0}^{n-1} (f_i - \bar{f})^2} \sqrt{\frac{1}{n} \sum_{i=0}^{n-1} (g_i - \bar{g})^2}}\end{aligned}$$

- Cosine similarity.  $\frac{\vec{f} \cdot \vec{g}}{\|\vec{f}\| \|\vec{g}\|}$



# Matching criteria

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- Distance based

- Fixed threshold (FT):

- Report all matches within the threshold value.

- Nearest neighbor (NN):

- Report the nearest neighbor.

- Nearest Neighbor Distance Ratio (NNDR):

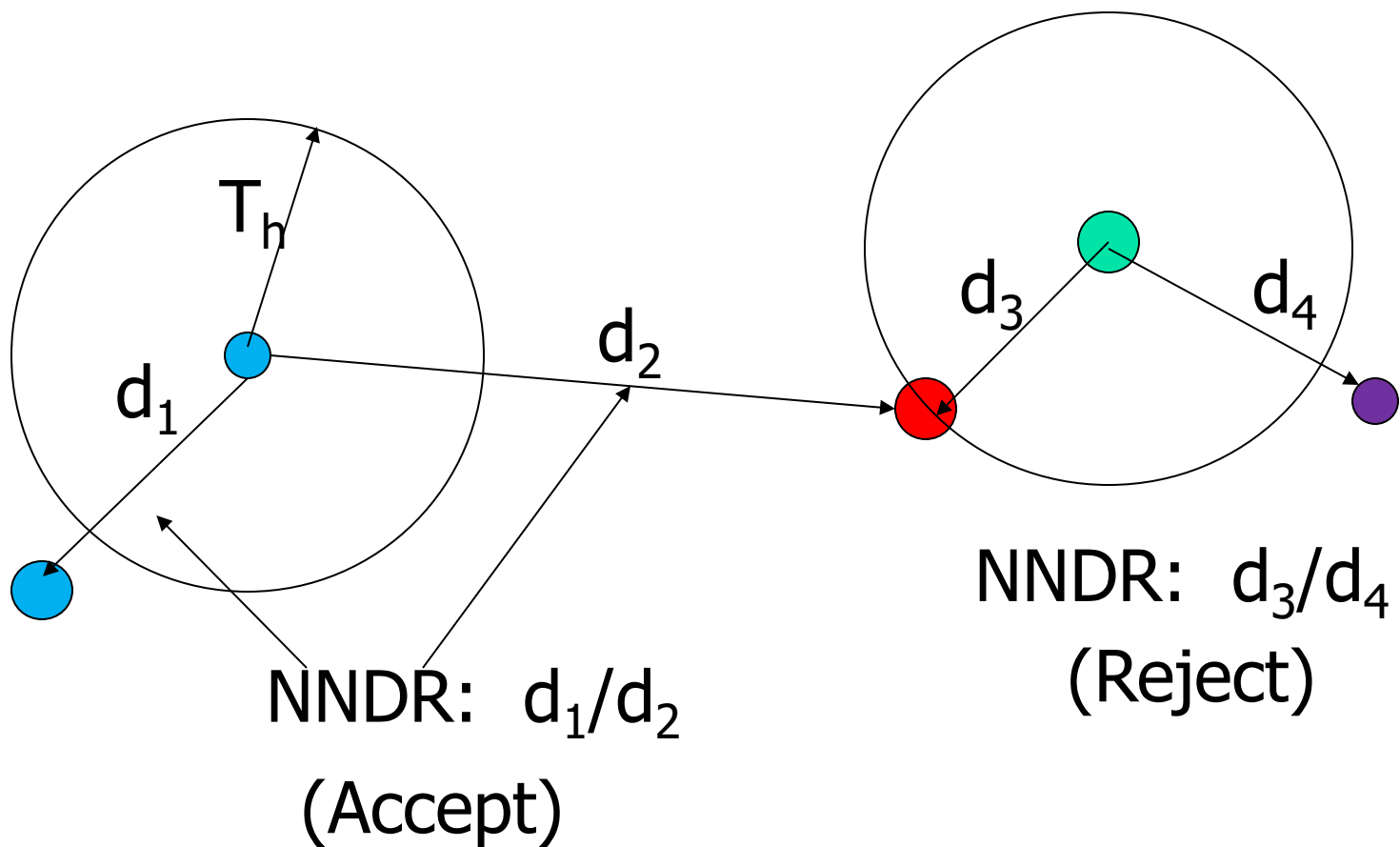
- Report the NN if ratio of distances between the NN and 2<sup>nd</sup> NN is small.





# Matching criteria

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# Matching histograms

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- $L_p$  norm of bins (representing same intervals of covariates).
  - Usually  $L_1$  used.

- Kullback-Leiber divergence (of pdf).

$$D_{KL}(P||Q) = - \sum_x P(x) \ln \frac{P(x)}{Q(x)}$$

- Earth Mover's Distance (EMD)



# Earth Mover's Distance (EMD)

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- EMD of two histograms  $P$  and  $Q$  measures the minimum accumulated cost of transferring masses from any bin of  $P$  to any bin of  $Q$  so that the histogram  $P$  gets transformed into  $Q$ .
  - Cost defined by the product of transferred mass and distance between bin.
    - $m \times |i-j|$  for transferring mass  $m$  from  $i$ th bin of  $P$  to  $j$ th bin of  $Q$ .
    - This would make  $P[i]=P[i]-m$  and  $Q[j]=Q[j]+m$ .
  - Total mass of  $P$  and  $Q$  should be same.
  - Accumulated cost is normalized w.r.t. to the total transfer of mass.

# Computing EMD

- Two normalized histograms:
  - $P=\{p_i\}$  and  $Q=\{q_i\}$ , for  $i=0,1,\dots,N-1$ .
  - $m_{ij}$  transferred from  $i$  th bin of  $P$  to  $j$  th bin of  $Q$ .
  - $d_{ij}$  distance between  $i$  th and  $j$  th bin.
- EMD is the minimum *normalized* work (transfer of masses) required for transforming  $P$  into  $Q$ .

$$EMD(P, Q) = \min_{M=\{m_{ij}\}} \left( \frac{\sum_{i,j} m_{ij} d_{ij}}{\sum_{i,j} m_{ij}} \right) \quad m_{ij} \geq 0$$

$$\sum_j m_{ij} \leq p_i \quad \sum_i m_{ij} \leq q_j \quad \sum_i \sum_j m_{ij} \leq \min \left( \sum_i p_i, \sum_j q_j \right)$$

# Computing EMD

- EMD is the minimum *normalized* work (transfer of masses) required for transforming  $P$  into  $Q$ .
- Before initiating transfer operation, bins of  $Q$  is initialized to zero. The input specification  $q_j$ 's are used for specifying constraints of the optimization equations.

$$EMD(P, Q) = \min_{M=\{m_{ij}\}} \left( \frac{\sum_{i,j} m_{ij} d_{ij}}{\sum_{i,j} m_{ij}} \right) \quad m_{ij} \geq 0$$

$$\sum_j m_{ij} \leq p_i \quad \sum_i m_{ij} \leq q_j \quad \sum_i \sum_j m_{ij} \leq \min \left( \sum_i p_i, \sum_j q_j \right)$$



# Efficient computation of feature matching

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- Range query

- Nearest Neighbor of the query.
- Report all the features within a distance from the query.
- Ordinarily of linear time complexity  $O(N)$ ,
  - $N$  is the number of features in the target image / database.

- Use of indexing and hashing.

- K-D Tree
- Locality Sensitive Hashing.



# K-D tree

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- A binary tree.
- Each node contains a (key) feature vector.
- Each node behaves like a node of binary search tree (BST) for values of a particular dimension
  - Called cut-dimension.
  - Alternate periodically on nodes along any path.
  - Places a key feature vector following the rules of BST comparing on the value of the cut-dimension.



# NN Search using K-D tree

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- As you walk through the tree nodes
  - Store the current smallest distance and the respective key-feature.
  - Prune sub-trees by comparing the minimum distance with the corner nodes of the hypervolume (bounding boxes) represented by the sub-tree.
  - Search the sub-tree to maximize the chance of pruning.
    - Go to the sub-tree closer to the query.





# NN Search using K-D tree

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- May require to traverse all the nodes of the tree.
  - Worst case:  $O(N)$
  - In practice close to:  $O(\log(N) + 2^d)$ , where  $d$  is the dimension of the feature space.
    - $\log(N)$  to find the cell near the query point.
    - $2^d$  search around cells in the neighborhood.



# Locality sensitive hashing (LSH)

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- Locality preserving hashing function ( $h(.)$ ) w.r.t a distance function  $d(.)$ .
  - Prob. ( $h(\mathbf{x})=h(\mathbf{y})$ ) should be high, if  $d(\mathbf{x},\mathbf{y})$  is small,
    - else the probability should be small.



# LSH: A typical example

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- Choose a random unit vector  $\mathbf{r}$  of dimension  $n$  (e.g. following Normal distribution  $(N(0,1))$  independently in each dimension).
- For any vector  $\mathbf{x}$  of dimension  $n$  define a hash function as follows:

$$h_{\mathbf{r}}(\mathbf{x}) = \begin{cases} 1 & \text{if } \mathbf{x} \cdot \mathbf{r} > 0 \\ 0 & \text{otherwise} \end{cases}$$

It can be shown, for any  $\mathbf{x}$  and  $\mathbf{y}$

$$\Pr[h_{\mathbf{r}}(\mathbf{x}) = h_{\mathbf{r}}(\mathbf{y})] = 1 - \frac{\theta(\mathbf{x}, \mathbf{y})}{\pi}$$

← Angle between  $\mathbf{x}$  and  $\mathbf{y}$ .



# LSH multidimensional bucketing

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- You may also have a set of  $k$  random vectors independently generating  $k$  hash values, and define the multidimensional bucket for hashing to place the input vector.
  - $H(\mathbf{x}) = [h_{r1}(\mathbf{x}) \ h_{r2}(\mathbf{x}) \ \dots \ h_{rk}(\mathbf{x})]$
- You may have  $L$  multiple buckets (corresponding to  $L$  sets of  $k$  random vectors) for the same input.
  - $H_1(\mathbf{x}), H_2(\mathbf{x}), \dots, H_L(\mathbf{x})$
- Given a query  $\mathbf{y}$  compute  $L$  buckets and search for NN in all of them.



# Model fitting

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- Given a set of data points fit a model to establish relationship among the data points.
  - Homography matrix between pixels in two images of the same planar scene.
  - Fundamental matrix between corresponding points in two stereo images.
  - Given a set of 2-D points fit a straight line / parabolic curve / circle / a high degree polynomial curve passing through them.
- Obtain data points and fit appropriate model.
  - Preprocessing, feature detection-description-matching.



# Knowledge of models

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- Mathematical form of the class / family of Models.
  - 3x3 non-singular homography matrix for projective transformation.
  - 3x3 singular fundamental matrix for stereo geometry.
  - 3x4 Projection matrix.



# Choice of a model

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- Error of fitting
  - Mean square error
- Likelihood
  - $\Pr(\text{Data} \mid \text{Model})$
- Size of a model
  - Number of independent parameters
- Training and Test Error
  - Large training error: Under-fitting.
  - Low training error, but large test error: Over-fitting
  - Low training and test error: A good fit.

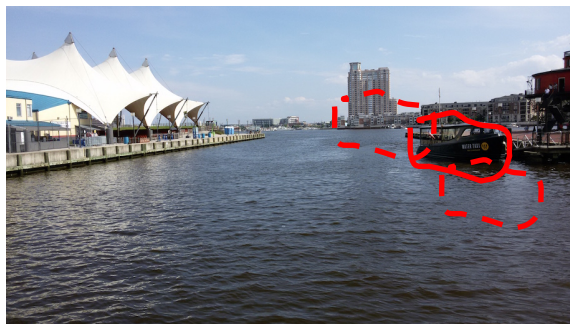
# Fitting curves and lines



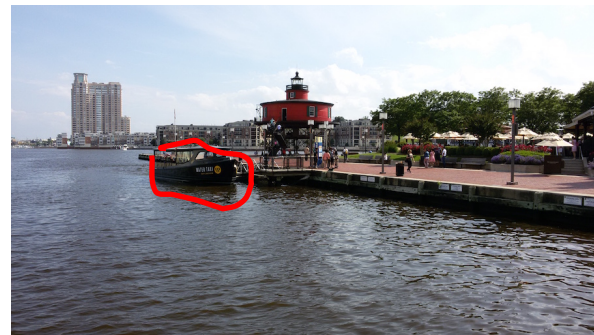
simple model: lines



simple model: circles



To decide  
about  
appropriate  
parametric  
models.



complicated model: Steam-boat





# Various Issues

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- **Noise / Error**
  - in estimation of feature locations.
- **Extraneous data**
  - clutter (outliers), multiple lines.
- **Missing data**
  - occlusions



# Fitting a straight line over 2D points

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- Fitting techniques
  - Least Squares
  - Total Least Squares
  - Random Sample Consensus (RANSAC)
  - Hough Voting



# Line fitting in varying contexts

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- Given points belonging to a line, to find the “optimal” line parameters.
  - Least squares
- To handle outliers.
  - RANSAC
- Presence of many lines.
  - Voting methods: Hough transform

# Least squares line fitting

$$E = \sum_{i=1}^n (y_i - mx_i - c)^2$$

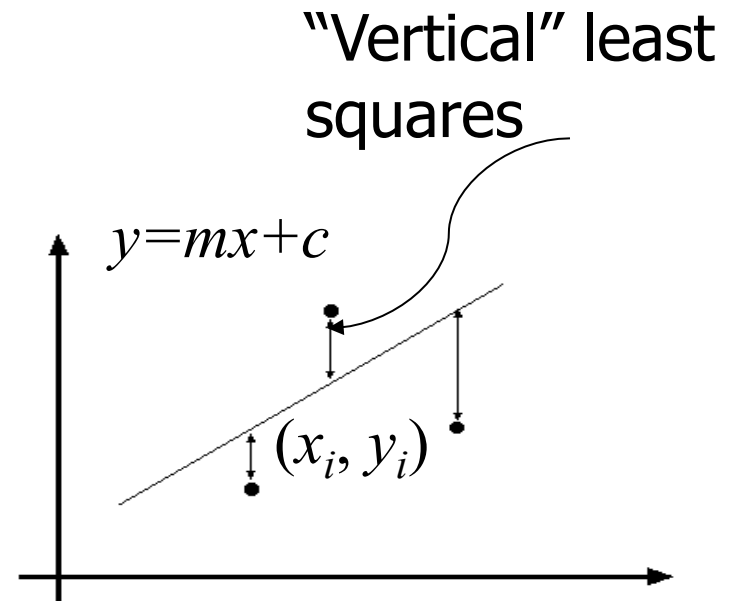
- Data:

- $(x_1, y_1), \dots, (x_n, y_n)$

- Line equation:

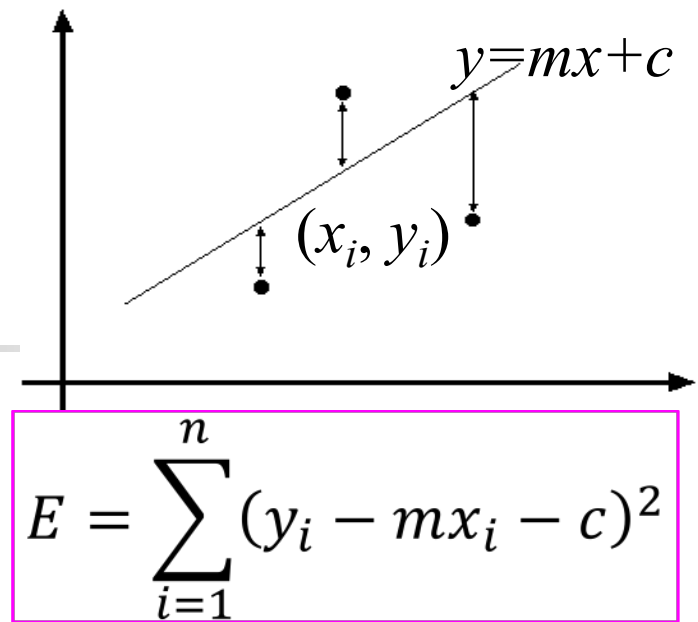
- $y_i = m x_i + c$

- Find  $(m, c)$  to minimize



# Least squares line fitting

- Data:  $(x_1, y_1), \dots, (x_n, y_n)$
- Line equation:  $y_i = mx_i + c$
- Find  $(m, c)$  to minimize

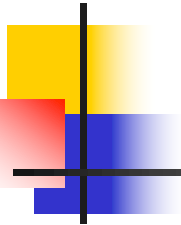


$$E = \sum_{i=1}^n \left( y_i - \begin{bmatrix} x_i & 1 \end{bmatrix} \begin{bmatrix} m \\ c \end{bmatrix} \right)^2 = \left\| \begin{bmatrix} y_1 \\ \vdots \\ y_n \end{bmatrix} - \begin{bmatrix} x_1 & 1 \\ \vdots & \vdots \\ x_n & 1 \end{bmatrix} \begin{bmatrix} m \\ c \end{bmatrix} \right\|^2 = \|Y - XC\|^2$$

$$E = (Y - XC)^T (Y - XC) = Y^T Y - 2(XC)^T Y + (XC)^T (XC)$$

$$\frac{dE}{dC} = 2X^T XC - 2X^T Y = 0 \quad \Rightarrow \quad C = (X^T X)^{-1} X^T Y$$

# Least squares line fitting



$$C = (X^T X)^{-1} X^T Y$$

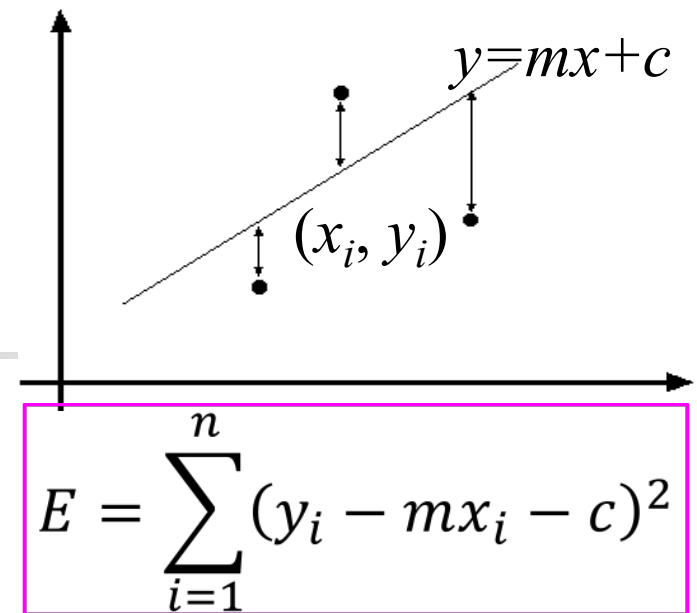
$$m = \frac{\text{cov}(X, Y)}{\text{var}(X)} = \frac{\frac{1}{n} \sum_i x_i y_i - \bar{x} \bar{y}}{\frac{1}{n} \sum_i x_i^2 - \bar{x}^2}$$

$$c = \bar{y} - m \bar{x}$$

The  $R^2$  goodness-of-fit criterion compares the variability in the measurements not explained by the model to the total variability in the measurements.

Goodness of fit: 
$$R^2 = 1 - \frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\sum_{i=1}^n (y_i - \bar{y})^2} \Rightarrow R^2 = 1 - \frac{E}{n \cdot \text{var}(y)}$$

- Not rotation-invariant.
- Fails completely for vertical lines.



$$E = n(\text{var}(y) - m^2 \text{var}(x))$$

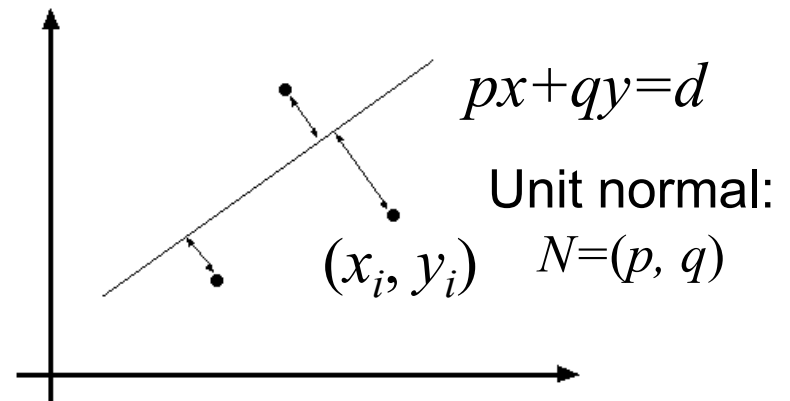
# Total least squares

- Distance between a point  $(x_i, y_i)$  and the line  $px + qy = d$  :  $|px_i + qy_i - d|$  given  $p^2 + q^2 = 1$ .

$$E = \sum_{i=1}^n (px_i + qy_i - d)^2$$

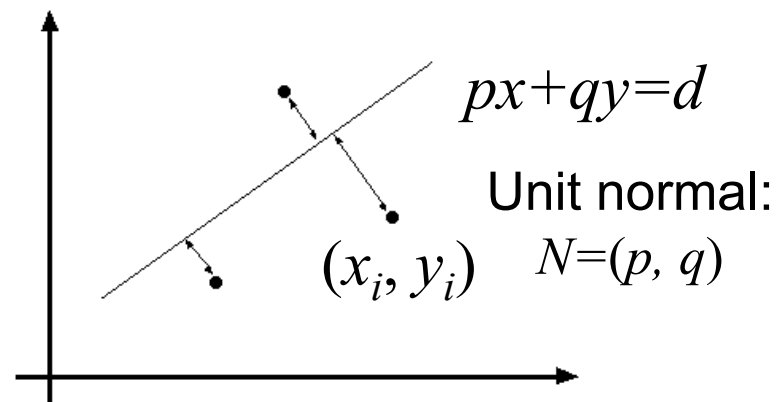
$$\frac{\partial E}{\partial d} = \sum_{i=1}^n -2(px_i + qy_i - d) = 0$$

$$\Rightarrow d = p\bar{x} + q\bar{y} \Rightarrow E = \sum_{i=1}^n (p(x_i - \bar{x}) + q(y_i - \bar{y}))^2$$



# Total least squares

$$E = \sum_{i=1}^n (p(x_i - \bar{x}) + q(y_i - \bar{y}))^2$$



$$E = \left\| \begin{matrix} \overbrace{\begin{bmatrix} x_1 - \bar{x} & y_1 - \bar{y} \\ \vdots & \vdots \\ x_n - \bar{x} & y_n - \bar{y} \end{bmatrix}}^U \underbrace{\begin{bmatrix} p \\ q \end{bmatrix}}_N \right\|^2 = (UN)^T (UN)$$

$$\frac{dE}{dN} = 2(U^T U)N = 0 \quad \text{Subject to } \|N\| = 1.$$

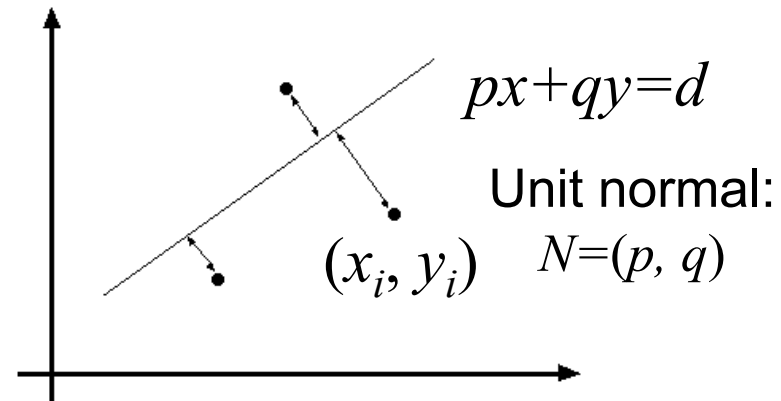
Eigen vector of  $U^T U$  corresponding to the smallest eigen value.



# Total least squares

$$E = \left\| \begin{bmatrix} x_1 - \bar{x} & y_1 - \bar{y} \\ \vdots & \vdots \\ x_n - \bar{x} & y_n - \bar{y} \end{bmatrix} \begin{bmatrix} p \\ q \end{bmatrix} \right\|^2 = (UN)^T(UN)$$

$$\frac{dE}{dN} = 2(U^T U)N = 0 \quad \text{Subject to } \|N\|=1.$$



$$U^T U = \begin{bmatrix} \sum_{i=1}^n (x_i - \bar{x})^2 & \sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y}) \\ \sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y}) & \sum_{i=1}^n (y_i - \bar{y})^2 \end{bmatrix}$$

A 2D coordinate system showing an ellipse centered at the origin, representing the confidence region for the total least squares fit. The ellipse is elongated along the direction of the line. A point  $(\bar{x}, \bar{y})$  is marked on the ellipse. A red vector  $N = (p, q)$  points from the origin to the ellipse. A blue vector  $(x_i - \bar{x}, y_i - \bar{y})$  points from the center of the ellipse to a point on the ellipse. The distance  $d$  is indicated as the perpendicular distance from the origin to the line.

Eigen vector of  $U^T U$  corresponding to the smallest eigen value.

# Random sample consensus (RANSAC)

- A general framework for model fitting in the presence of outliers.
- Outline
  - Choose a small subset of points uniformly at random.
  - Fit a model to that subset.
  - Find all remaining points that are “close” to the model and reject the rest as outliers.
  - Do this many times and choose the best model.



# RANSAC for line fitting

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- Perform the following  **$N$**  times:
  - Select  **$s$**  points uniformly at random.
  - Fit a line using the LSE method.
  - Find inliers to this line
    - points whose distance from the line is less than  **$t$** .
  - For sufficient number of inliers (say  $> d$ ), accept the line and refit using all inliers.



# Choice of parameters

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- Initial number of points  **$s$** 
  - Typically minimum number needed to fit the model.
- Distance threshold  **$t$** 
  - Choose  **$t$**  so probability for inlier is  $p$  (e.g. 0.95).
  - Zero-mean Gaussian noise with std. dev.  $\sigma$ :  $t=1.96\sigma$ .
- Number of trials  **$N$** 
  - Choose  **$N$**  so that, with probability  $p$ , at least in one trial random selections are free from any outlier. (e.g.  $p=0.99$ ) (outlier ratio:  $e$ ).
- Consensus set size  **$d$** 
  - Should match expected inlier ratio:  $(1-e) \times n$ .

$n$ =No. of data points



# Estimating number of trials $N$

Outlier ratio:  $e$  ← Prob. that a sample is outlier.

Prob. that all  $s$  samples are inliers:  $(1 - e)^s$

Prob. that at least one sample is an outlier in a trial:  $\Rightarrow (1 - (1 - e)^s)$

Prob. that all  $N$  trials have an outlier:  $(1 - (1 - e)^s)^N$

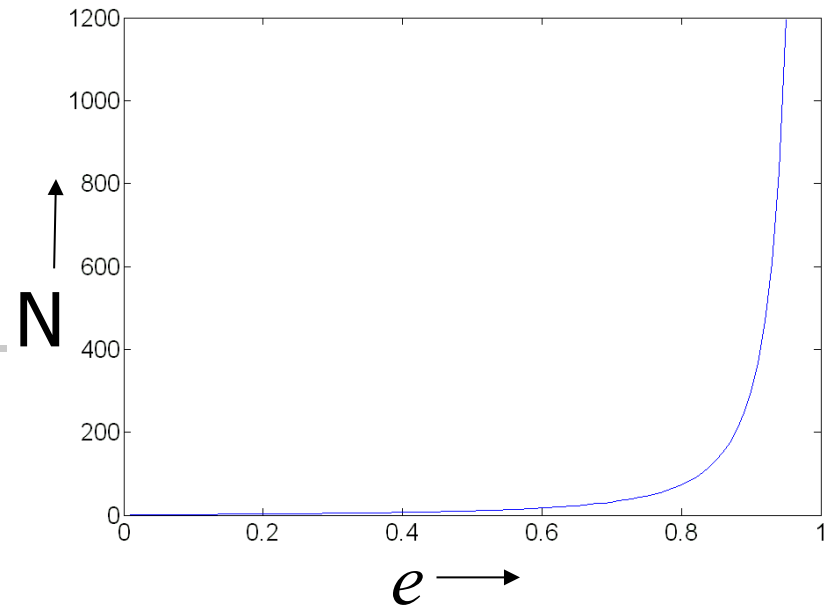
Given probability  $p$ , so that at least one random sample is free from outliers.  $\Rightarrow (1 - (1 - e)^s)^N = 1 - p$

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

# Choosing N

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

$$p=0.99$$



s	proportion of outliers $e$						
	5%	10%	20%	25%	30%	40%	50%
2	2	3	5	6	7	11	17
3	3	4	7	9	11	19	35
4	3	5	9	13	17	34	72
5	4	6	12	17	26	57	146
6	4	7	16	24	37	97	293
7	4	8	20	33	54	163	588
8	5	9	26	44	78	272	1177



# RANSAC pros and cons

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- Pros

- Simple and general
- Applicable to many different problems
- Often works well in practice

- Cons

- Many parameters to tune
- Can't always get a good initialization of the model based on the minimum number of samples
- Sometimes too many iterations are required
- Not appropriate for low inlier ratio.



# Voting schemes

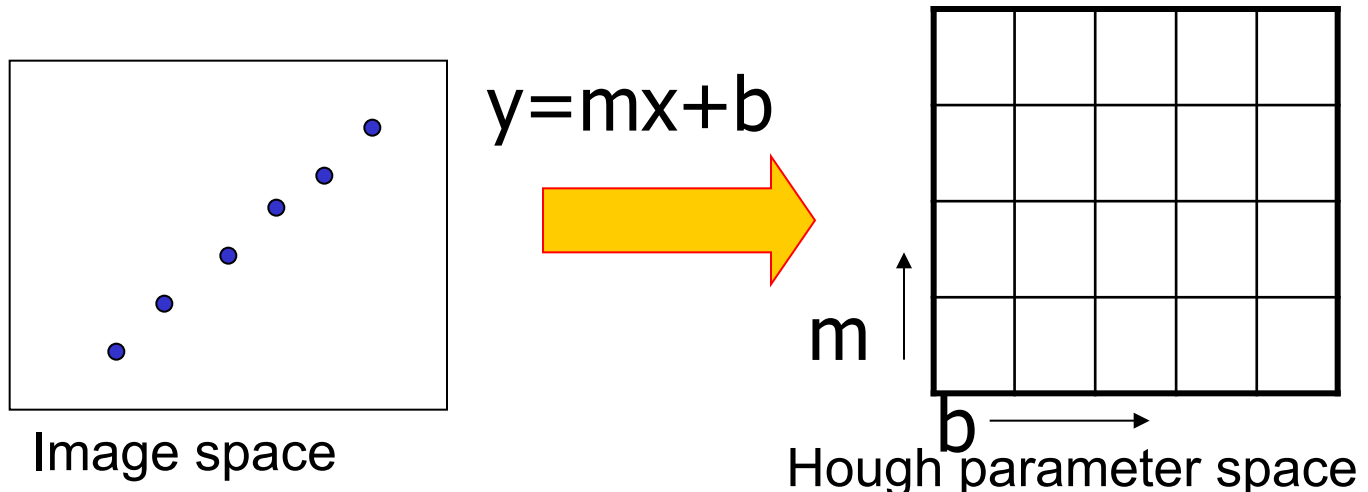
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- Let each feature vote for all the models that are compatible with it.
- Hopefully the noise features will not vote consistently for any single model
- Missing data doesn't matter as long as there are enough features remaining to agree on a good model.



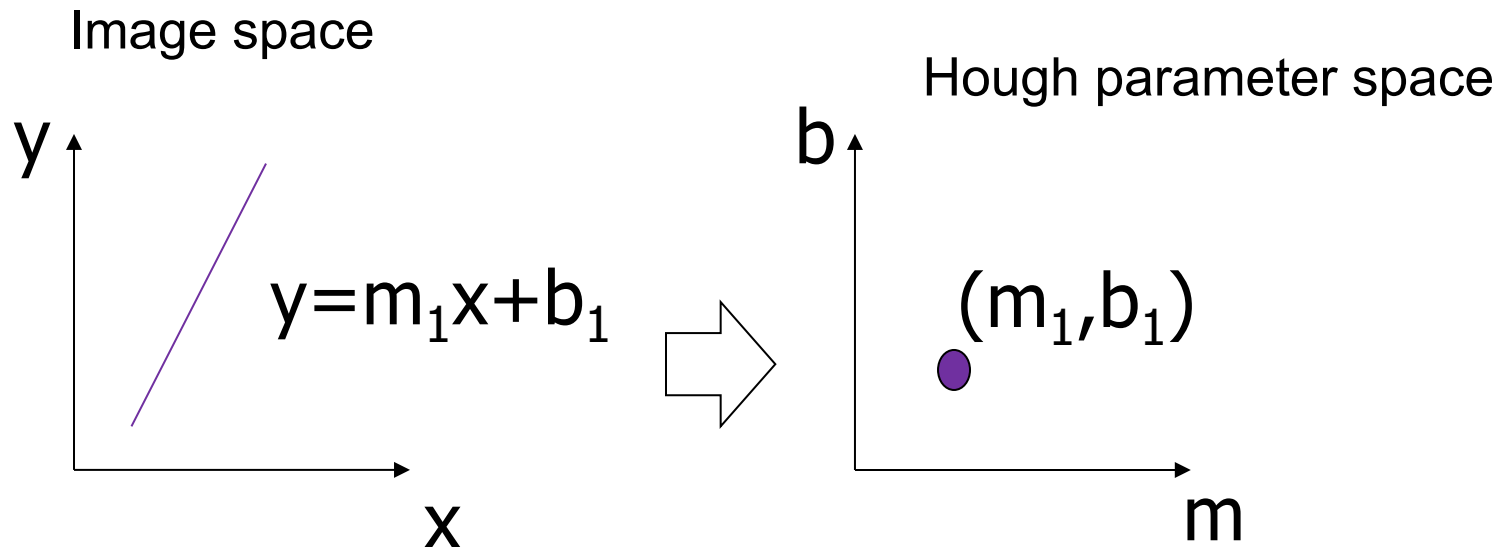
# Hough transform

- ❑ Discretize parameter space into bins
- ❑ For each feature point in the image, put a vote in every bin in the parameter space that could have generated this point.
- ❑ Find bins that have the most votes.



# Parameter space representation

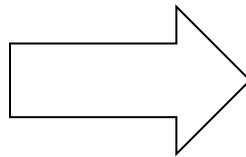
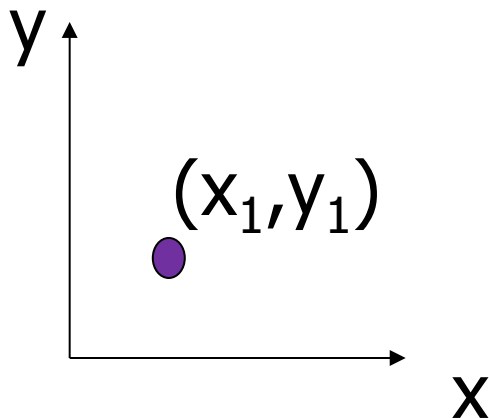
- A straight line in the image corresponds to a point in Hough space.



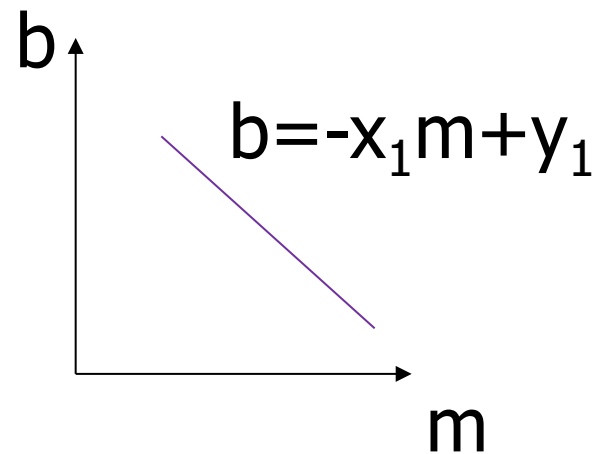
# Parameter space representation

- A point in the image corresponds to a line in the Hough space.

Image space

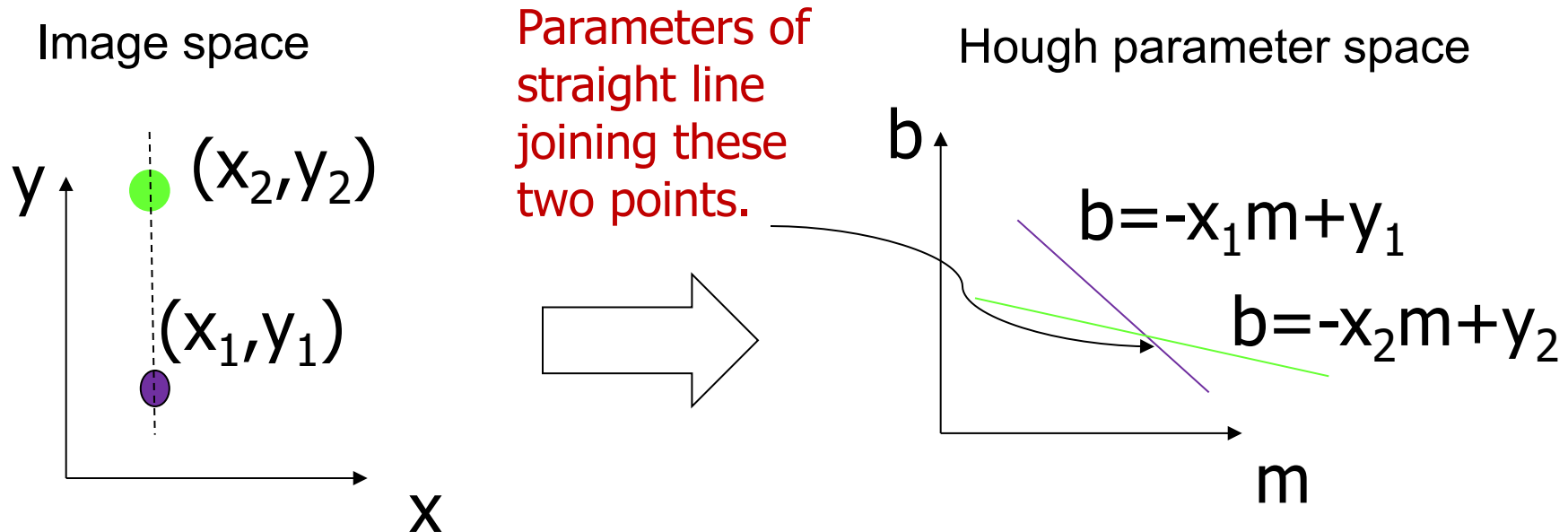


Hough parameter space



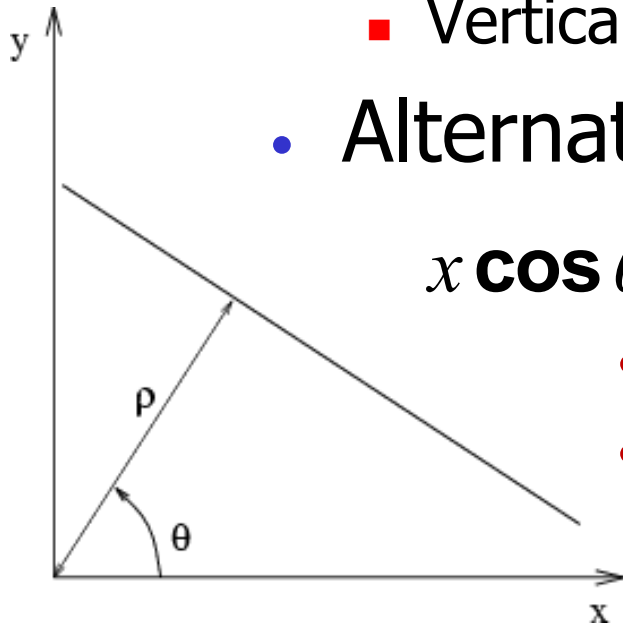
# Parameter space representation

- Two points in the image correspond to two lines in the Hough space.



# Parameter space representation

- Problems with the (m,b) space:
  - Unbounded parameter domain
  - Vertical lines require infinite m
- Alternative: polar representation



$$x \cos \theta + y \sin \theta = \rho$$

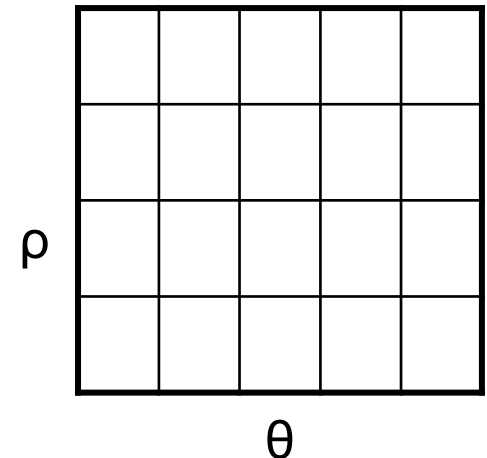
- $\theta$  varies from 0 to 180°
- $\rho$  varies from 0 to the length of diagonal of the image grid.

Each point will add a sinusoid in the  $(\theta, \rho)$  parameter space

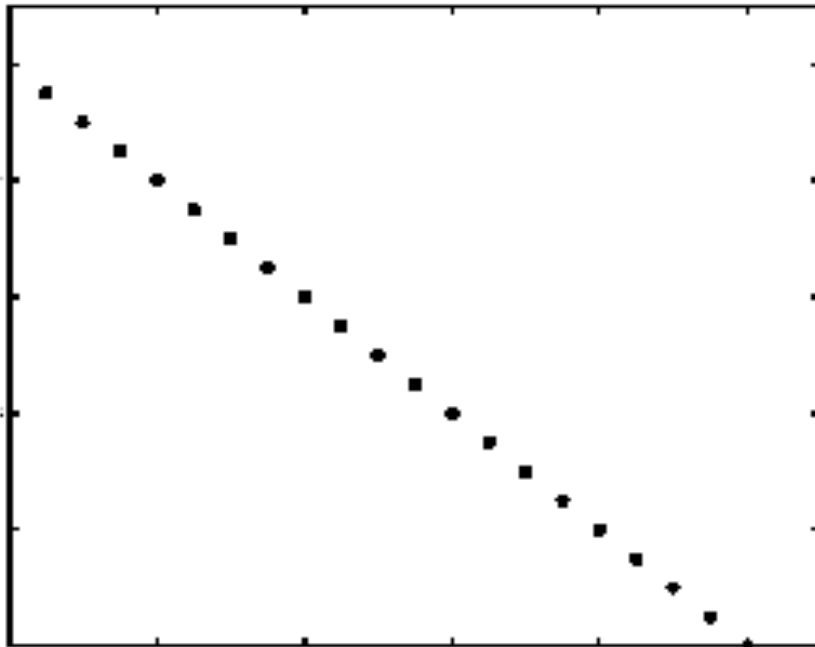
# Algorithm

- Initialize accumulator  $A$  to all zeros
- For each edge point  $(x,y)$  in the image
- {
  - For  $\theta = 0$  to  $180$
  - {
$$\rho = x \cos \theta + y \sin \theta$$
$$A(\theta, \rho) = A(\theta, \rho) + 1$$
}}
- Find the value(s) of  $(\theta, \rho)$  where  $A(\theta, \rho)$  is a local maximum
- The detected line in the image is given by
$$\rho = x \cos \theta + y \sin \theta$$

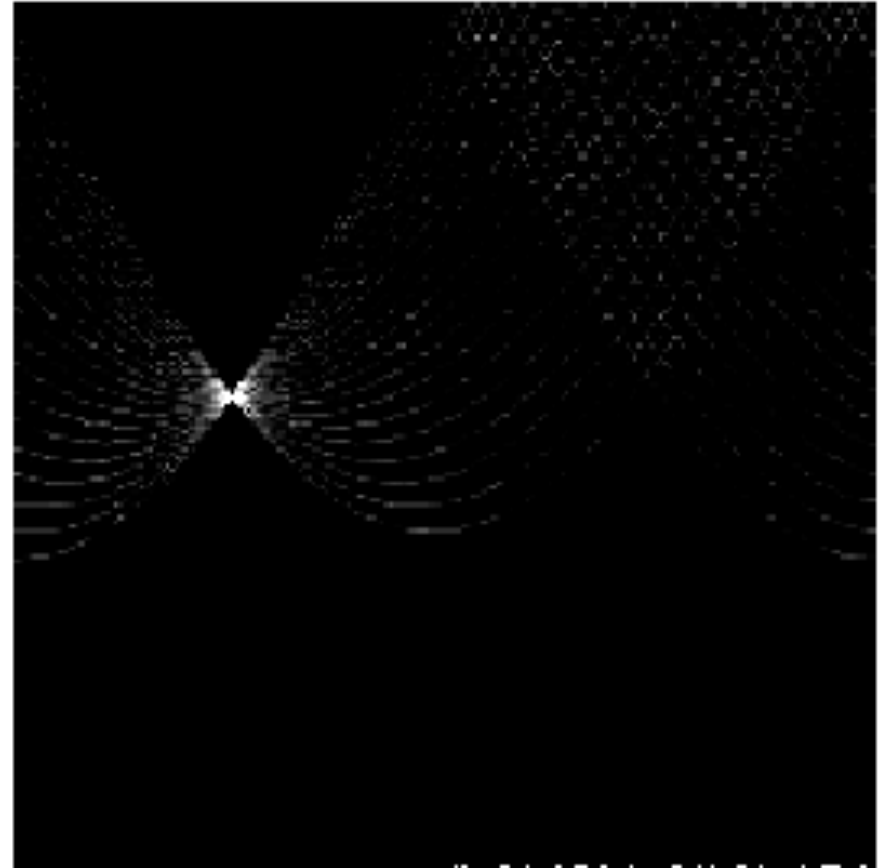
A: Accumulator array



# Basic illustration

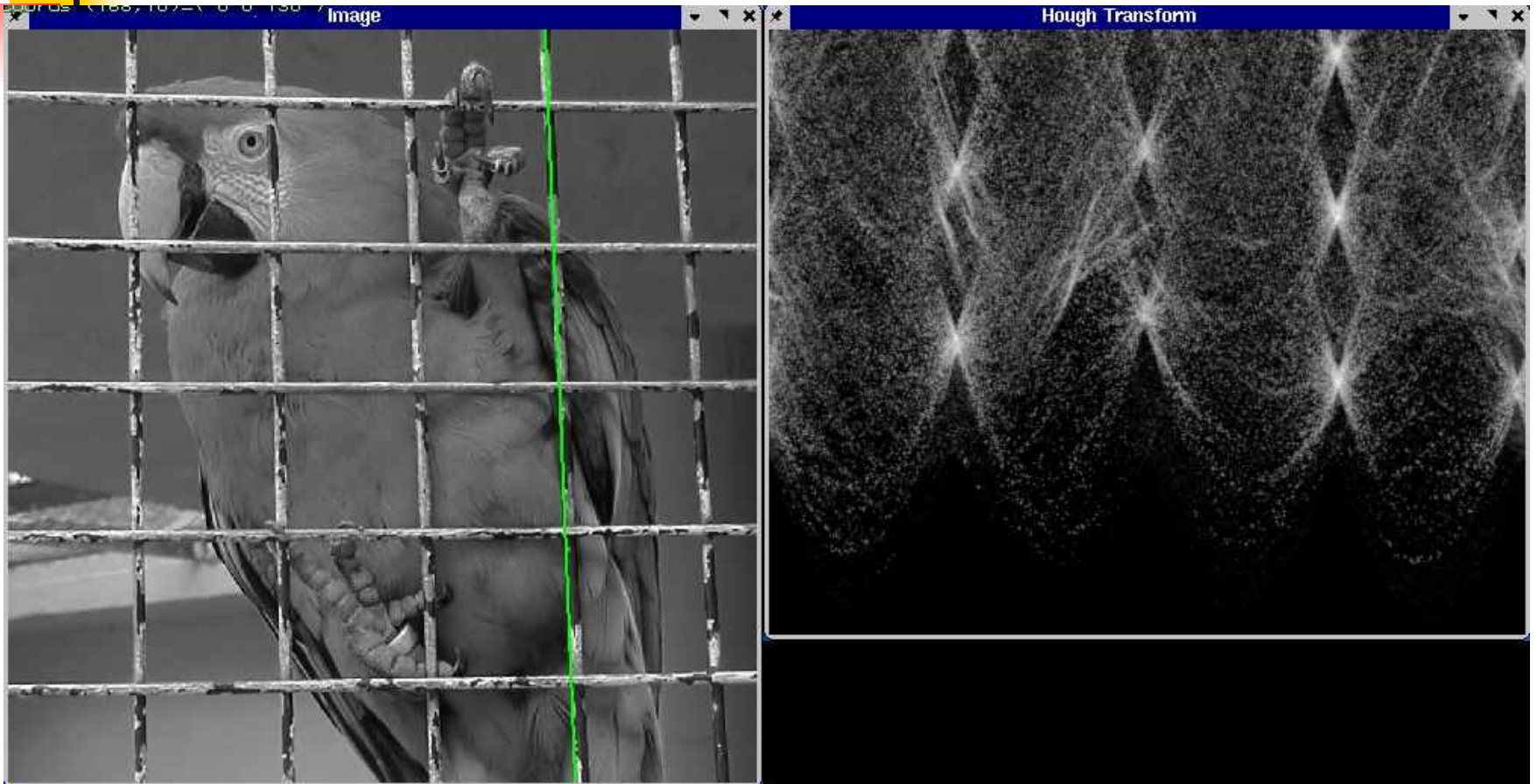


features



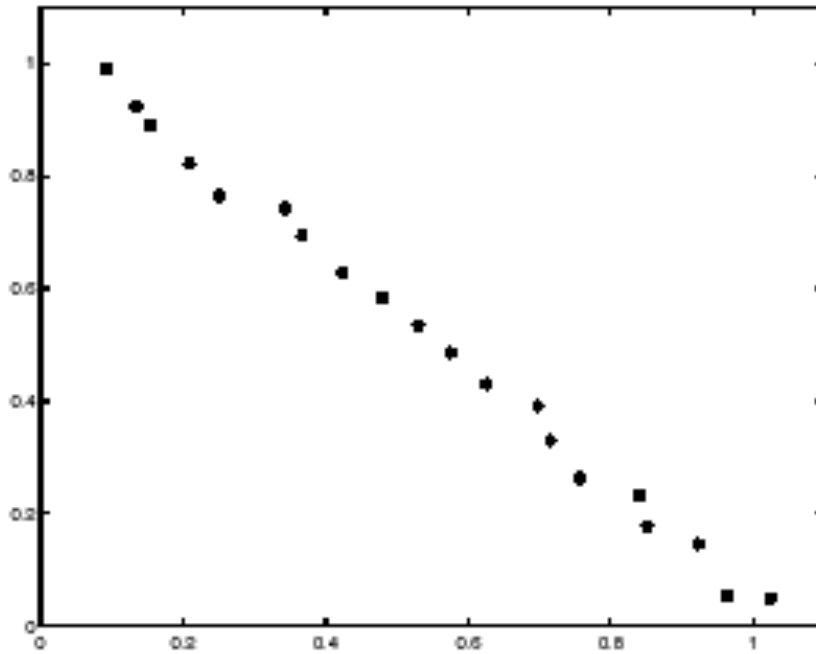
votes

# A more complicated image

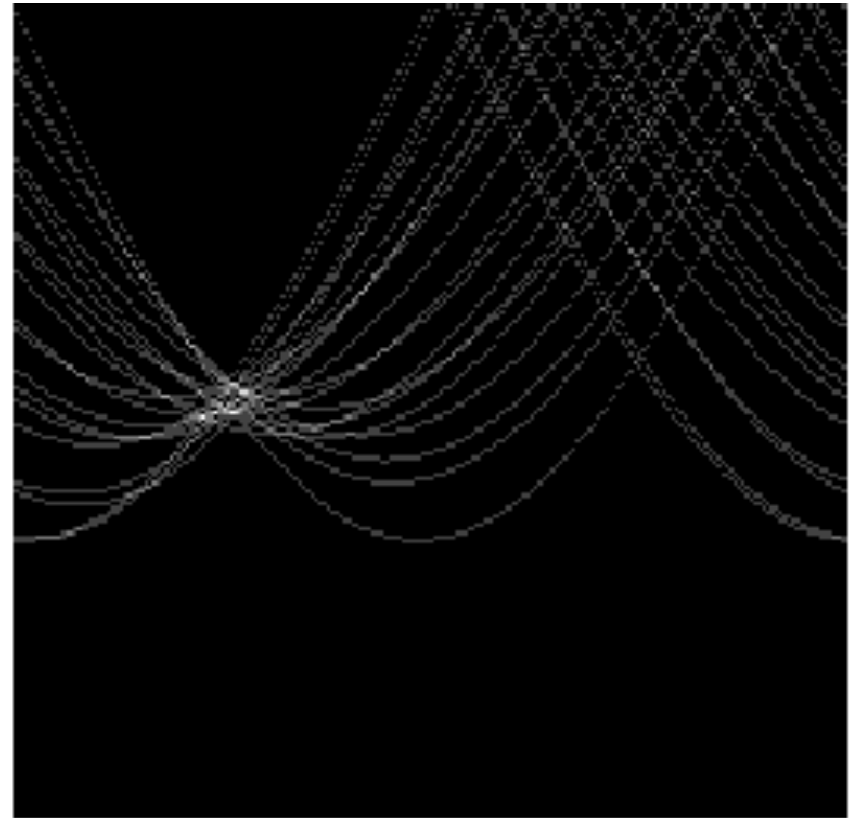




# Effect of noise



features



votes



# Dealing with noise

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- Requires appropriate resolution of discretization of the grid in the parameter space
  - Too coarse:
    - large votes in a cell for accumulating too many different lines correspond to a single bucket
  - Too fine:
    - may miss lines as some points may not be exactly collinear voting different buckets.



# Dealing with noise

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- Smoothing accumulator array.
- Try to get rid of irrelevant features
  - Take only edge points with significant gradient magnitude



# Summary

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- Feature matching.
  - Distance function / Similarity measure.
  - Different policies
    - Use of a threshold value.
    - Nearest Neighbor.
    - Nearest neighbor with distance ratio (NNDR).
  - Use of indexing / hashing for efficient computation.
  - Special distance functions for comparing histograms.
    - Kullback-Leiber Divergence (KLD)
    - Earth Mover's Distance (EMD)



# Summary

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- Model fitting

- Prior knowledge of the model useful.
- Need to consider goodness of fit.
- Different techniques for line fitting
  - Least squares
  - Total least squares.
  - RANSAC
  - Hough Transform.