

# Dimensionality Reduction

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Machine Learning 10-701/15-781

Apr 5, 2010

Slides Courtesy: Tom Mitchell, Eric Xing, Lawrence Saul

The logo consists of the letters 'ML' in a bold, black, sans-serif font. A thick red horizontal line is positioned directly beneath the 'L'.

**MACHINE LEARNING** DEPARTMENT

The logo features the text 'Carnegie Mellon.' in a red serif font, with 'School of Computer Science' in a smaller black sans-serif font below it. To the left of the text is a decorative graphic of a grid of dots that tapers to the right.

# High-Dimensional data

- High-Dimensions = Lot of Features

## Document classification

Features per document =  
thousands of words/unigrams  
millions of bigrams, contextual  
information



## Surveys - Netflix

480189 users x 17770 movies

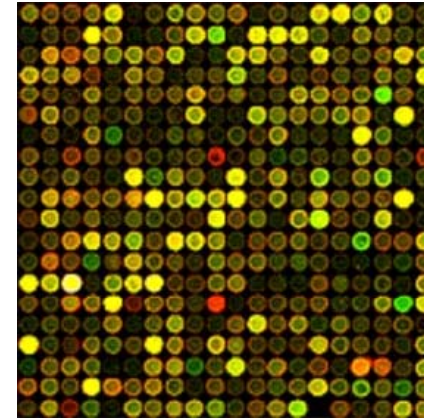
	movie 1	movie 2	movie 3	movie 4	movie 5	movie 6
Tom	5	?	?	1	3	?
George	?	?	3	1	2	5
Susan	4	3	1	?	5	1
Beth	4	3	?	2	4	2

# High-Dimensional data

- High-Dimensions = Lot of Features

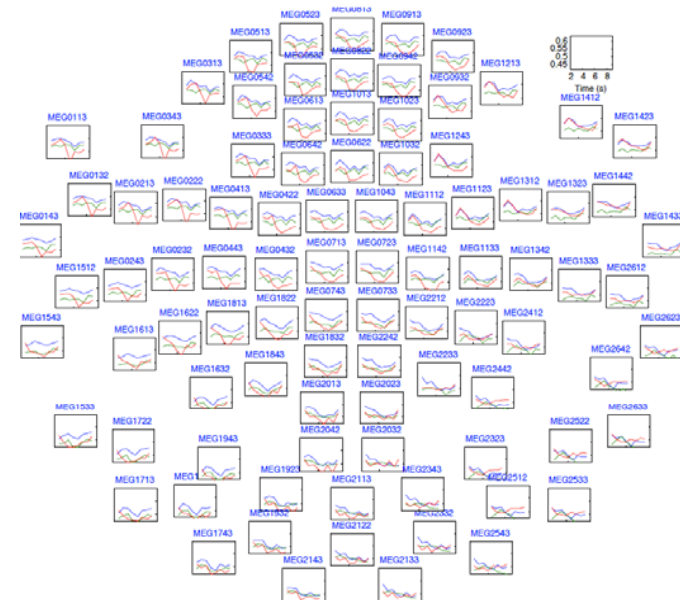
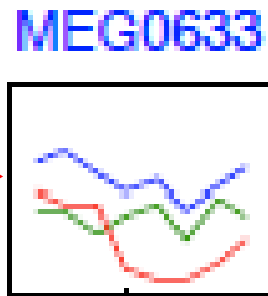
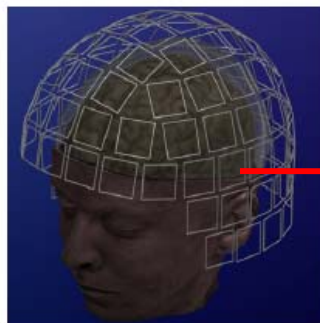
## Discovering gene networks

10,000 genes x 1000 drugs  
x several species



## MEG Brain Imaging

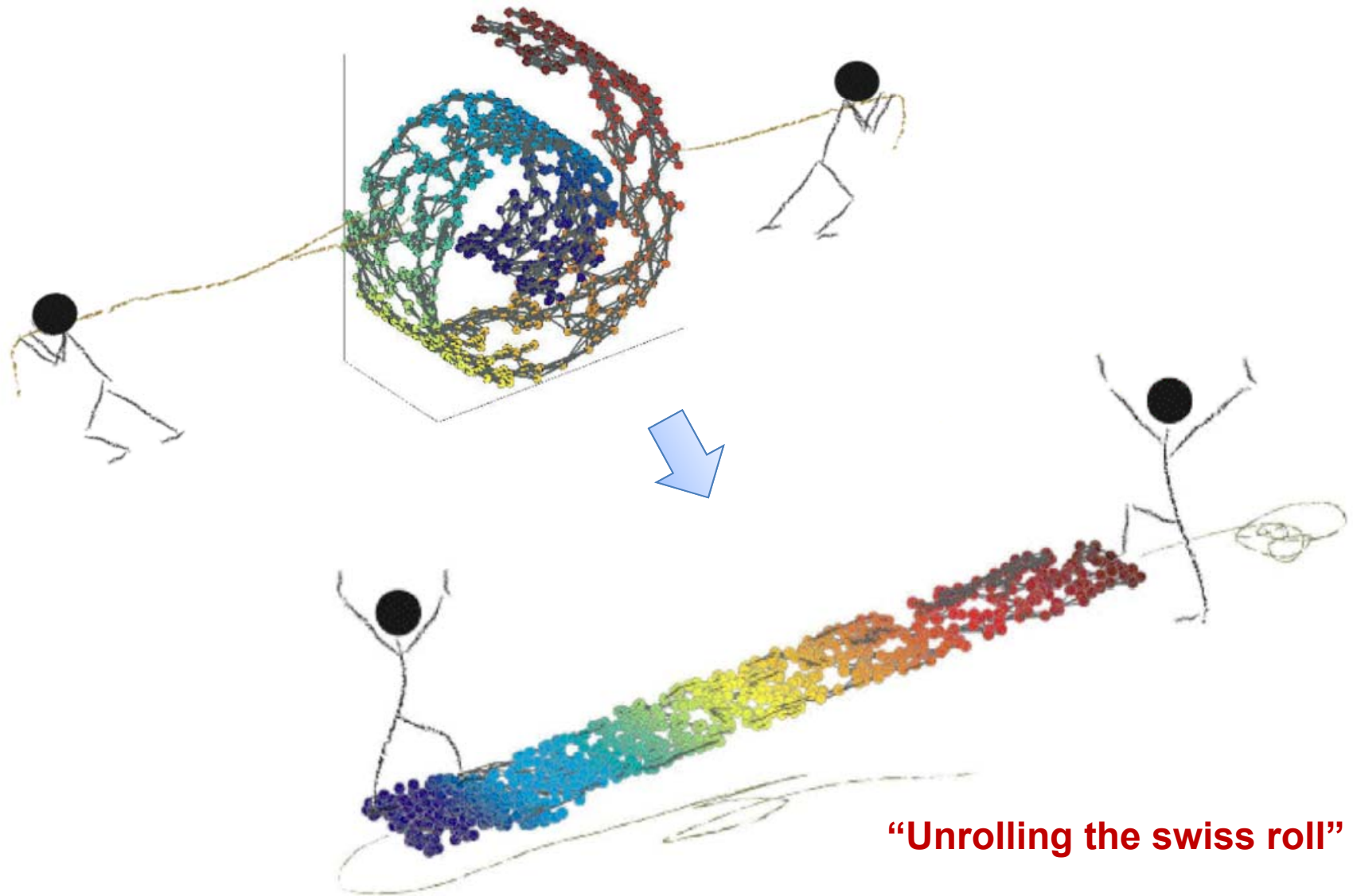
120 locations x 500 time points  
x 20 objects



# Curse of Dimensionality

- Why are more features bad?
  - Redundant features (not all words are useful to classify a document)  
more noise added than signal
  - Hard to interpret and visualize
  - Hard to store and process data (computationally challenging)
  - Complexity of decision rule tends to grow with # features. Hard to learn complex rules as VC dimension increases (statistically challenging)

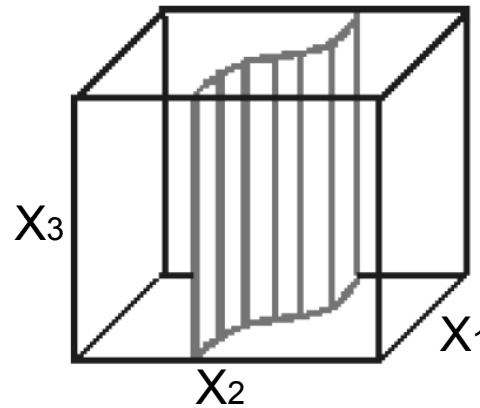
# Dimensionality Reduction



**“Unrolling the swiss roll”**

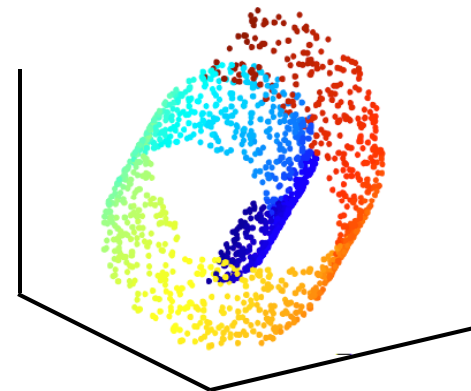
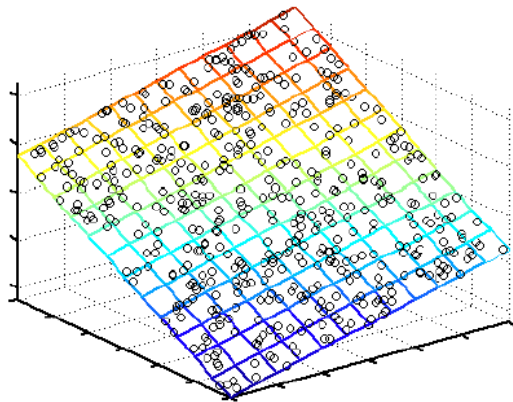
# Dimensionality Reduction

- **Feature Selection** – Only a few features are relevant to the learning task



$X_3$  - Irrelevant

- **Latent features** – Some linear/nonlinear combination of features provides a more efficient representation than observed features



# Feature Selection

- Approach 1: Score each feature and extract a subset

Common scoring methods:

- Training or cross-validated accuracy of single-feature classifiers  $f_i: X_i \rightarrow Y$

- Estimated mutual information between  $X_i$  and  $Y$ :

$$\hat{I}(X_i, Y) = \sum_k \sum_y \hat{P}(X_i = k, Y = y) \log \frac{\hat{P}(X_i = k, Y = y)}{\hat{P}(X_i = k) \hat{P}(Y = y)}$$

- $\chi^2$  statistic to measure independence between  $X_i$  and  $Y$
- Domain specific criteria
  - Text: Score “stop” words (“the”, “of”, ...) as zero
  - fMRI: Score voxel by T-test for activation versus rest condition
  - ...

# Feature Selection

- Approach 1: **Score each feature and extract a subset**

Common subset selection methods:

- One step: Choose  $d$  highest scoring features
- Iterative:
  - Choose single highest scoring feature  $X_k$
  - Rescore all features, conditioned on the set of already-selected features
    - E.g.,  $\text{Score}(X_i | X_k) = I(X_i, Y | X_k)$
    - E.g.,  $\text{Score}(X_i | X_k) = \text{Accuracy}(\text{predicting } Y \text{ from } X_i \text{ and } X_k)$
  - Repeat, calculating new scores on each iteration, conditioning on set of selected features



# Feature Selection: Text Classification

Approximately  $10^5$  words in English

[Rogati&Yang, 2002]

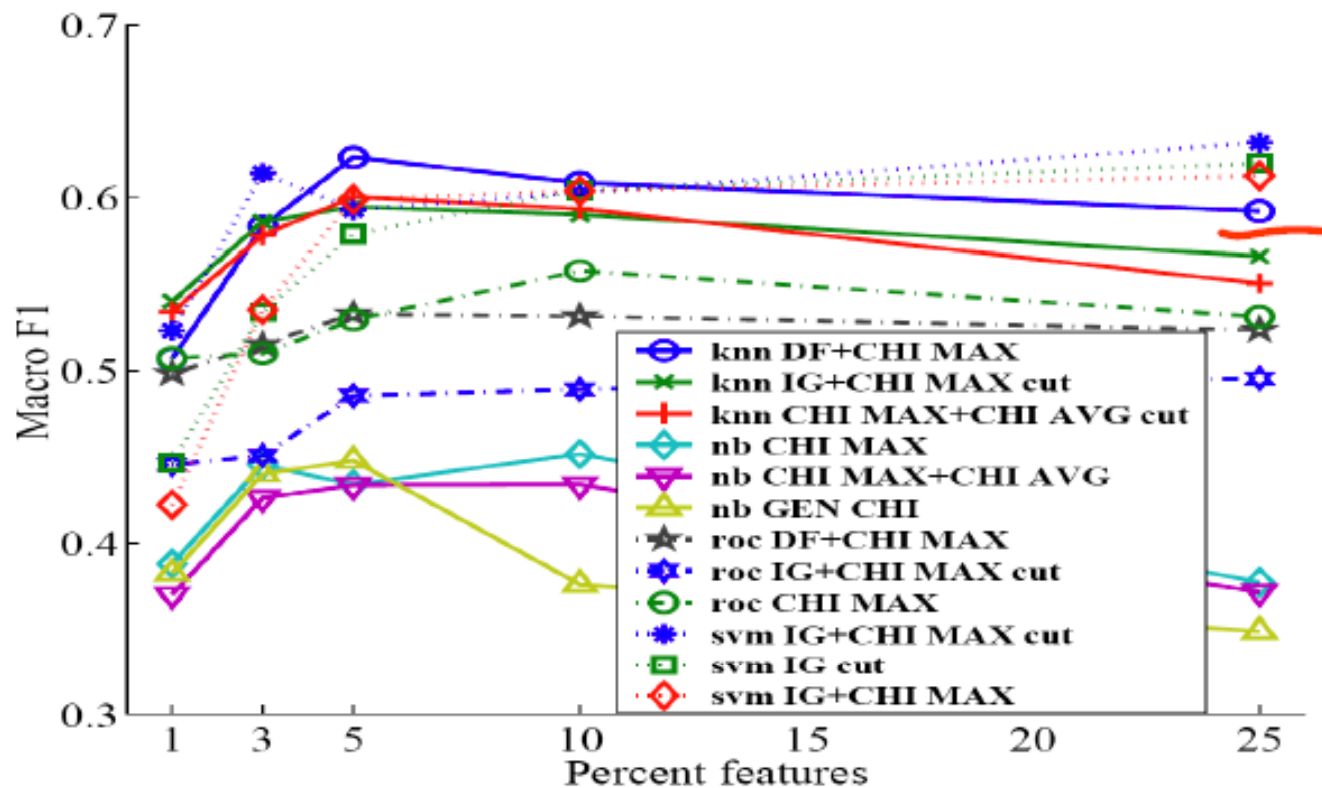


Figure 2: Top 3 feature selection methods for Reuters-21578 (Macro F1)

IG=information gain, chi=  $\chi^2$ , DF=doc frequency,

# Impact of Feature Selection on Classification of fMRI Data

[Pereira et al., 2005]

Accuracy classifying  
category of word read  
by subject

#voxels	mean	subjects							
		233B	329B	332B	424B	474B	496B	77B	86B
50	0.735	0.783	0.817	0.55	0.783	0.75	0.8	0.65	0.75
100	0.742	0.767	0.8	0.533	0.817	0.85	0.783	0.6	0.783
200	0.737	0.783	0.783	0.517	0.817	0.883	0.75	0.583	0.783
<b>300</b>	<b>0.75</b>	<b>0.8</b>	<b>0.817</b>	<b>0.567</b>	<b>0.833</b>	<b>0.883</b>	<b>0.75</b>	<b>0.583</b>	<b>0.767</b>
400	0.742	0.8	0.783	0.583	0.85	0.833	0.75	0.583	0.75
800	0.735	0.833	0.817	0.567	0.833	0.833	0.7	0.55	0.75
1600	0.698	0.8	0.817	0.45	0.783	0.833	0.633	0.5	0.75
all (~2500)	0.638	0.767	0.767	0.25	0.75	0.833	0.567	0.433	0.733

Table 1: Average accuracy across all pairs of categories, restricting the procedure to use a certain number of voxels for each subject. The highlighted line corresponds to the best mean accuracy, obtained using 300 voxels.

Each feature  $X_i$  is a voxel, scored by error in regression to predict  $X_i$  from  $Y$

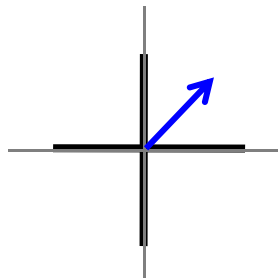
# Feature Selection

- Approach 2: **Regularization (MAP)**

Integrate feature selection into learning objective by penalizing number of features with non-zero weights

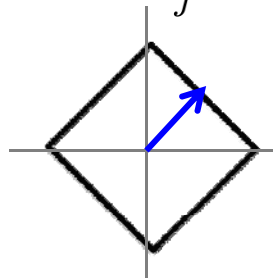
$$\widehat{W} = \arg \max_W \underbrace{\sum_{i=1}^n \log P(Y_i | X_i; W)}_{\text{log likelihood}} + \underbrace{\lambda \|W\|}_{\text{penalty}}$$

$$\|W\|_0 = \#\{W_j > 0\}$$



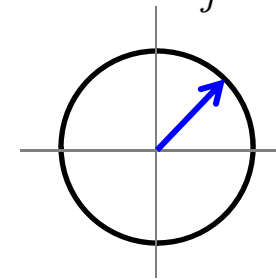
Minimizes # features chosen

$$\|W\|_1 = \sum_j |W_j|$$



Convex compromise

$$\|W\|_2 = \sum_j W_j^2$$



Small weights of features chosen

# Latent Feature Extraction

Combinations of observed features provide more efficient representation, and capture underlying relations that govern the data

E.g. Ego, personality and intelligence are hidden attributes that characterize human behavior instead of survey questions

Topics (sports, science, news, etc.) instead of documents

Often may not have physical meaning

- Linear

Principal Component Analysis (PCA)

Factor Analysis

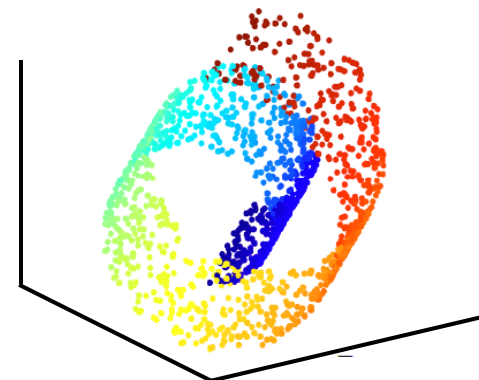
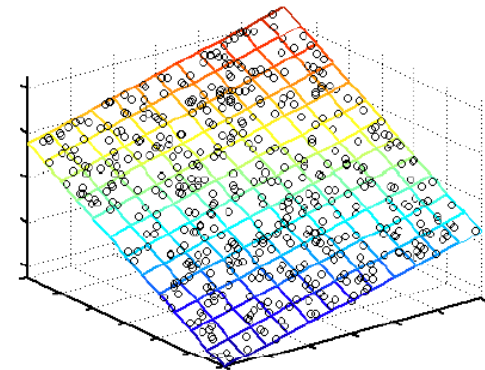
Independent Component Analysis (ICA)

- Nonlinear

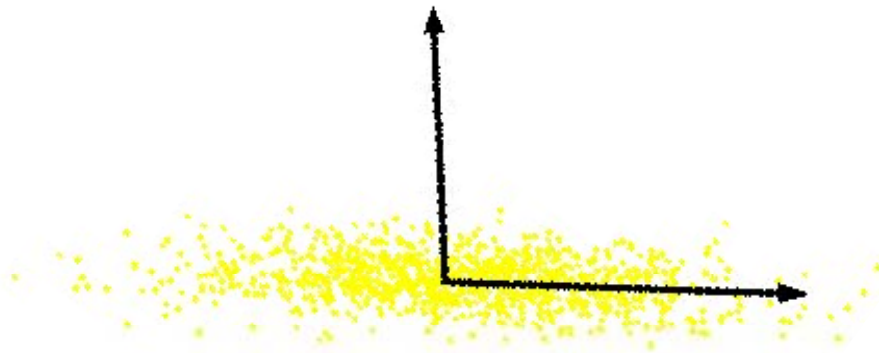
ISOMAP

Local Linear Embedding (LLE)

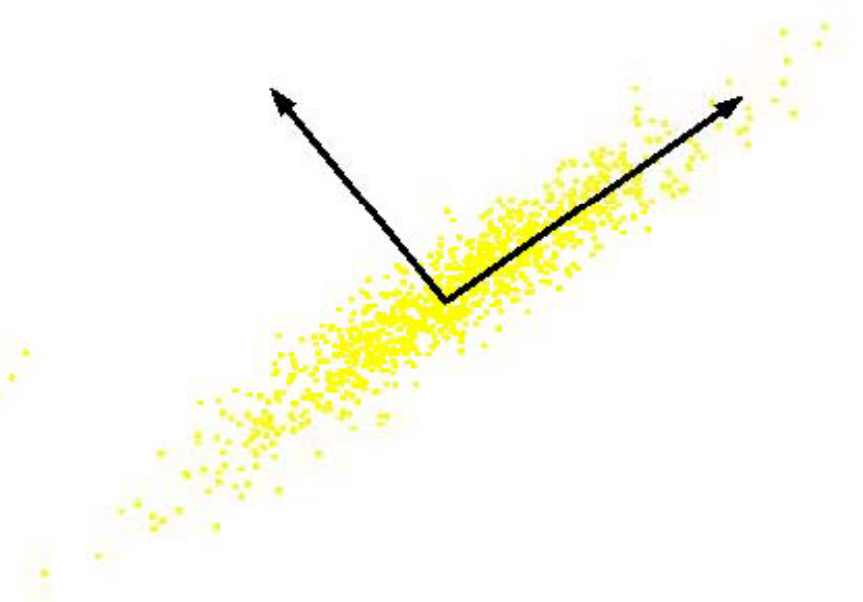
Laplacian Eigenmaps



# Principal Component Analysis (PCA)



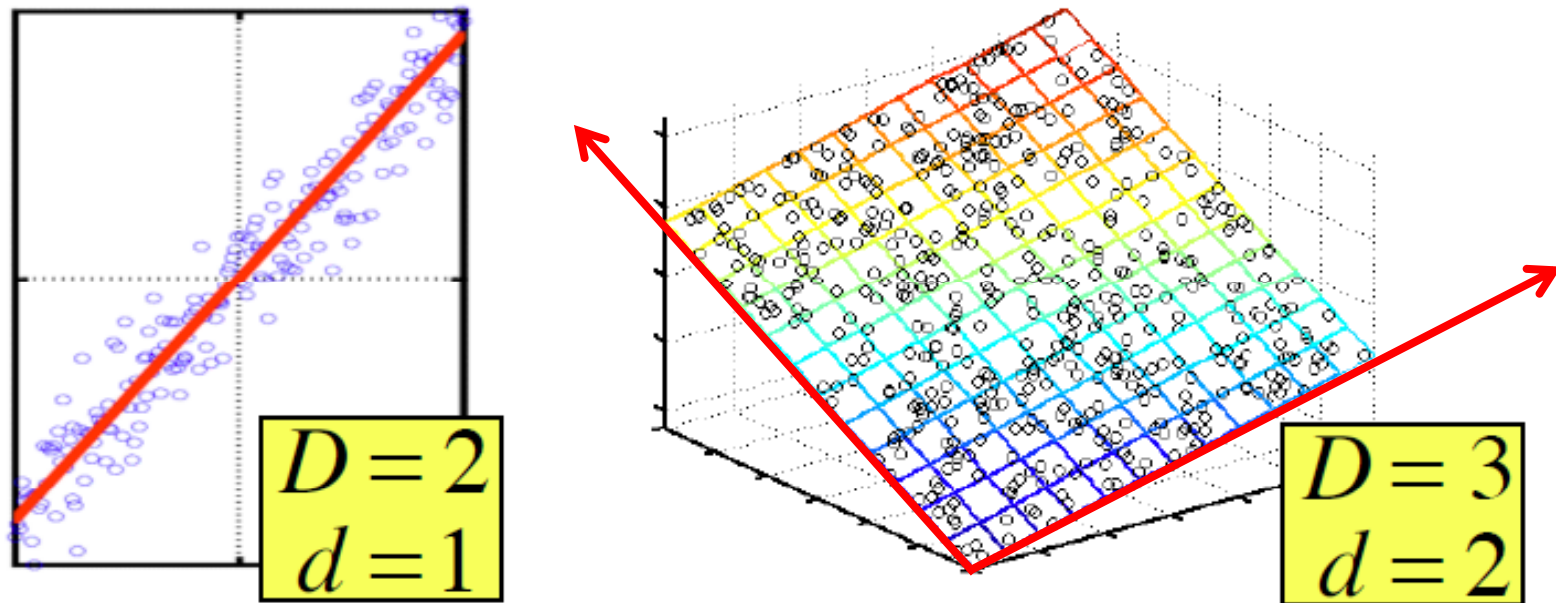
Only one relevant feature



Both features become relevant

Can we transform the features so that we only need to preserve one latent feature? Find linear projection so that projected data is uncorrelated.

# Principal Component Analysis (PCA)

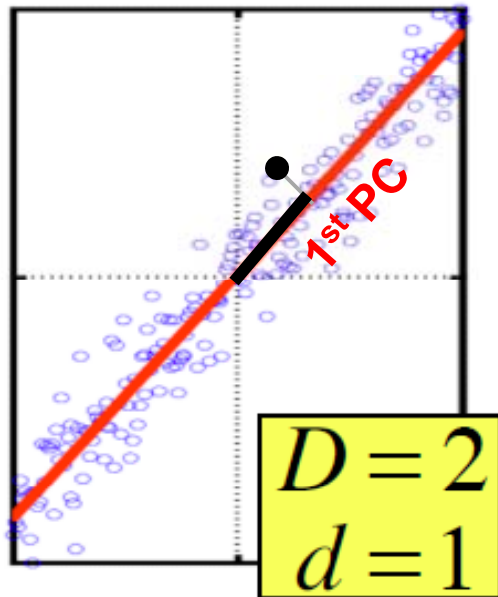


**Assumption:** Data lies on or near a low  $d$ -dimensional linear subspace.

Axes of this subspace are an effective representation of the data

Identifying the axes is known as Principal Components Analysis, and can be obtained by Eigen or Singular value decomposition

# Principal Component Analysis (PCA)



Principal Components (PC) are orthogonal directions that capture most of the variance in the data

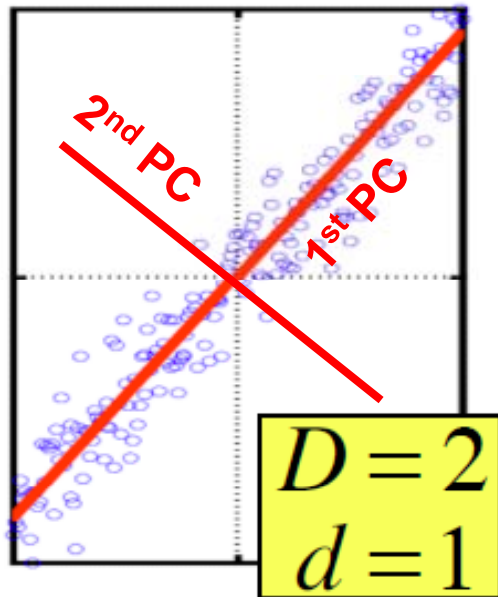
1<sup>st</sup> PC – direction of greatest variability in data

Projection of data points along 1<sup>st</sup> PC discriminate the data most along any one direction

Take a data point  $x_i$  (D-dimensional vector)

Projection of  $x_i$  onto the 1<sup>st</sup> PC  $v$  is  $v^T x_i$

# Principal Component Analysis (PCA)



Principal Components (PC) are orthogonal directions that capture most of the variance in the data

1<sup>st</sup> PC – direction of greatest variability in data

2<sup>nd</sup> PC – Next orthogonal (uncorrelated) direction of greatest variability

(remove all variability in first direction, then find next direction of greatest variability)

And so on ...



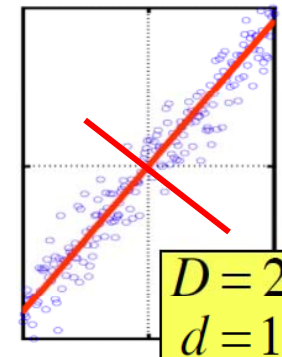
# Principal Component Analysis (PCA)

Let  $v_1, v_2, \dots, v_d$  denote the principal components

Orthogonal and unit norm  $v_i^T v_j = 0 \quad i \neq j$

$$v_i^T v_i = 1$$

Find vector that maximizes sample variance of projection



$$\frac{1}{n} \sum_{i=1}^n (v^T x_i)^2 = v^T \mathbf{X} \mathbf{X}^T v$$

Assume data are centered  
Data points  $X = [x_1 \ x_2 \ \dots \ x_n]$

$$\max_v v^T \mathbf{X} \mathbf{X}^T v \quad \text{s.t.} \quad v^T v = 1$$

Lagrangian:  $\max_v v^T \mathbf{X} \mathbf{X}^T v - \lambda v^T v$

Wrap constraints into the objective function

$$\partial / \partial v = 0 \quad (\mathbf{X} \mathbf{X}^T - \lambda \mathbf{I}) v = 0 \quad \Rightarrow \quad \boxed{(\mathbf{X} \mathbf{X}^T) v = \lambda v}$$

# Principal Component Analysis (PCA)

$$(\mathbf{X}\mathbf{X}^T)\mathbf{v} = \lambda\mathbf{v}$$

Therefore,  $\mathbf{v}$  is the eigenvector of  $\mathbf{X}\mathbf{X}^T$  with eigenvalue  $\lambda$

Sample variance of projection =  $\mathbf{v}^T \mathbf{X}\mathbf{X}^T \mathbf{v} = \lambda \mathbf{v}^T \mathbf{v} = \lambda$

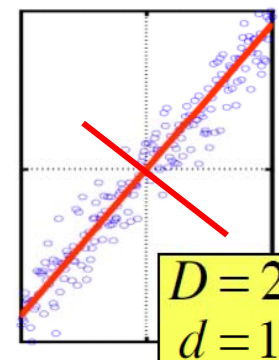
Thus, the eigenvalue  $\lambda$  denotes the amount of variability captured along that dimension.

Eigenvalues  $\lambda_1 > \lambda_2 > \lambda_3 > \dots$

The 1<sup>st</sup> Principal component  $\mathbf{v}_1$  is the eigenvector of the sample covariance matrix  $\mathbf{X}\mathbf{X}^T$  associated with the largest eigenvalue  $\lambda_1$

The 2<sup>nd</sup> Principal component  $\mathbf{v}_2$  is the eigenvector of the sample covariance matrix  $\mathbf{X}\mathbf{X}^T$  associated with the second largest eigenvalue  $\lambda_2$

And so on ...



# Computing the PCs

Eigenvectors are solutions of the following equation:

$$(\mathbf{X}\mathbf{X}^T)\mathbf{v} = \lambda\mathbf{v} \quad (\mathbf{X}\mathbf{X}^T - \lambda\mathbf{I})\mathbf{v} = 0$$

Non-zero solution  $\mathbf{v} \neq 0$  possible only if

$$\det(\mathbf{X}\mathbf{X}^T - \lambda\mathbf{I}) = 0 \quad \text{Characteristic Equation}$$

This is a  $D^{\text{th}}$  order equation in  $\lambda$ , can have at most  $D$  distinct solutions (roots of the characteristic equation)

Once eigenvalues are computed, solve for eigenvectors (Principal Components) using

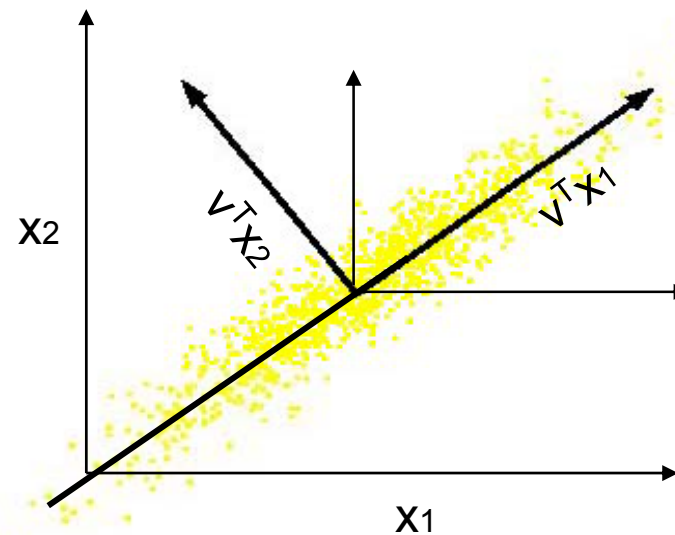
$$(\mathbf{X}\mathbf{X}^T - \lambda\mathbf{I})\mathbf{v} = 0$$

For symmetric matrices, eigenvectors for distinct eigenvalues are orthogonal.

# Principal Component Analysis (PCA)

So, the new axes are the eigenvectors of the matrix of sample correlations  $XX^T$  of the data, which capture the similarities of the original features based on how data samples project to the new axes.

Transformed features are uncorrelated.



- Geometrically: centering followed by rotation
  - Linear transformation

# Another interpretation

**Maximum Variance Subspace:** PCA finds vectors  $\mathbf{v}$  such that projections on to the vectors capture maximum variance in the data

$$\frac{1}{n} \sum_{i=1}^n (\mathbf{v}^T \mathbf{x}_i)^2 = \mathbf{v}^T \mathbf{X} \mathbf{X}^T \mathbf{v}$$

**Minimum Reconstruction Error:** PCA finds vectors  $\mathbf{v}$  such that projection on to the vectors yields minimum MSE reconstruction

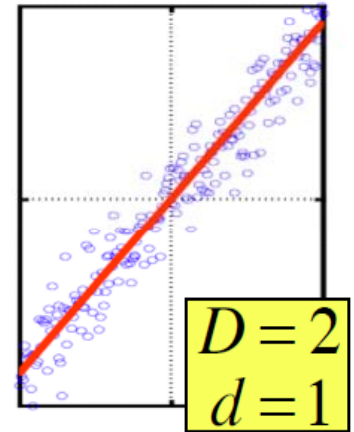
$$\frac{1}{n} \sum_{i=1}^n \|\mathbf{x}_i - \mathbf{v}^T \mathbf{x}_i\|^2$$

# Dimensionality Reduction using PCA

The eigenvalue  $\lambda$  denotes the amount of variability captured along that dimension.

Zero eigenvalues indicate no variability along those directions => data lies exactly on a linear subspace

Only keep data projections onto principal components with non-zero eigenvalues, say  $v_1, \dots, v_d$  where  $d = \text{rank}(XX^T)$



## Original Representation

data point

$$x_i = [x_i^1, x_i^2, \dots, x_i^D]$$

(D-dimensional vector)

## Transformed representation

projections

$$[v_1^T x_i, v_2^T x_i, \dots, v_d^T x_i]$$

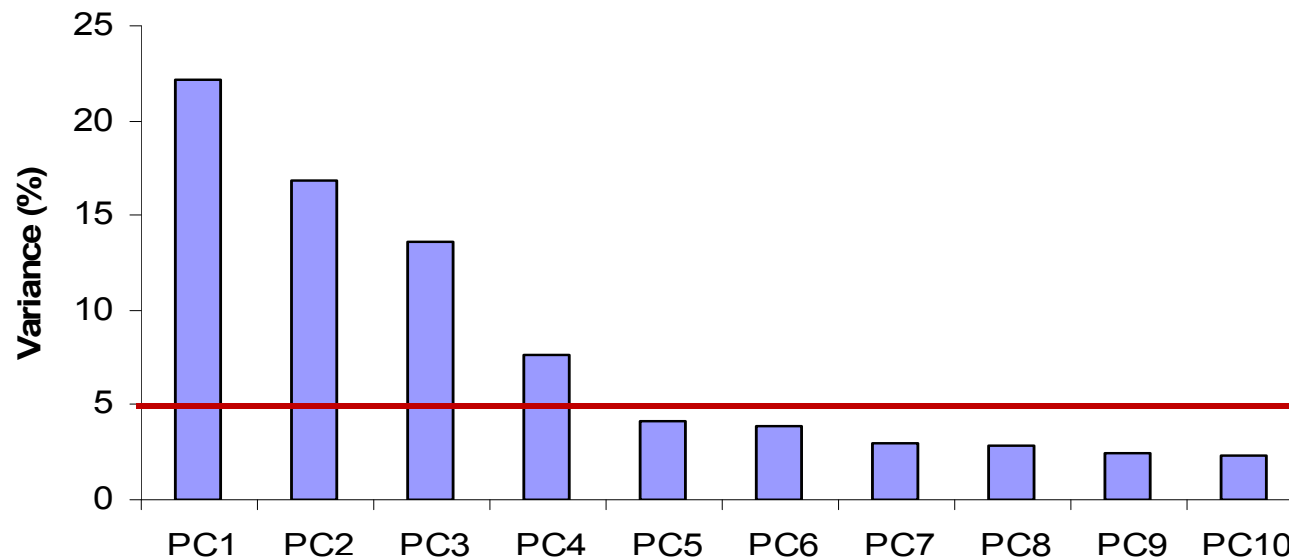
(d-dimensional vector)

# Dimensionality Reduction using PCA

Usually data lies near a linear subspace, as noise introduces small variability

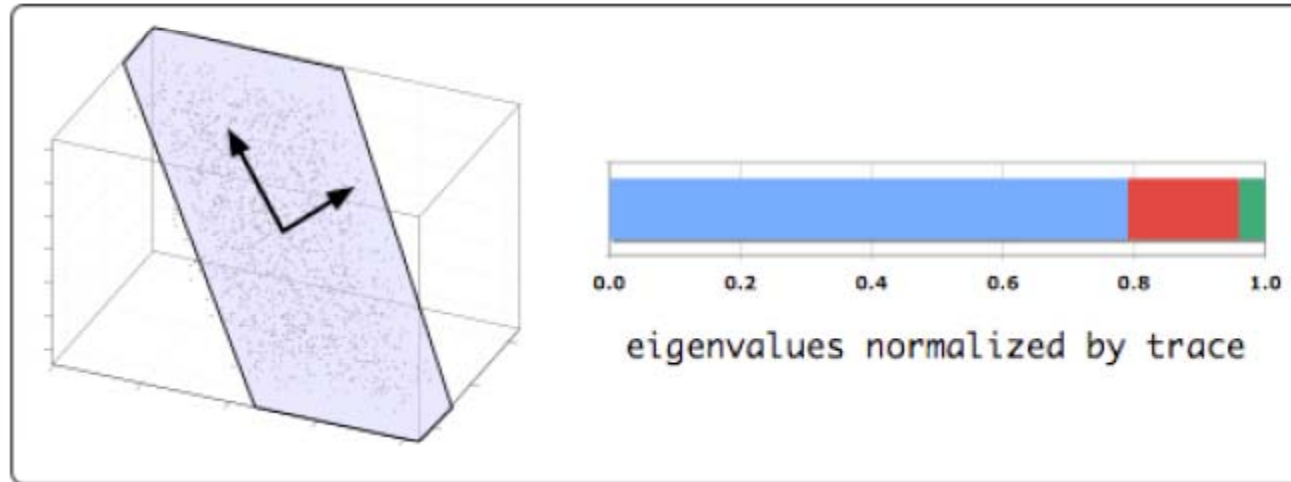
Only keep data projections onto principal components with **large** eigenvalues

Can *ignore the components of lesser significance*.



You might **lose some information**, but if the eigenvalues are small, you don't lose much

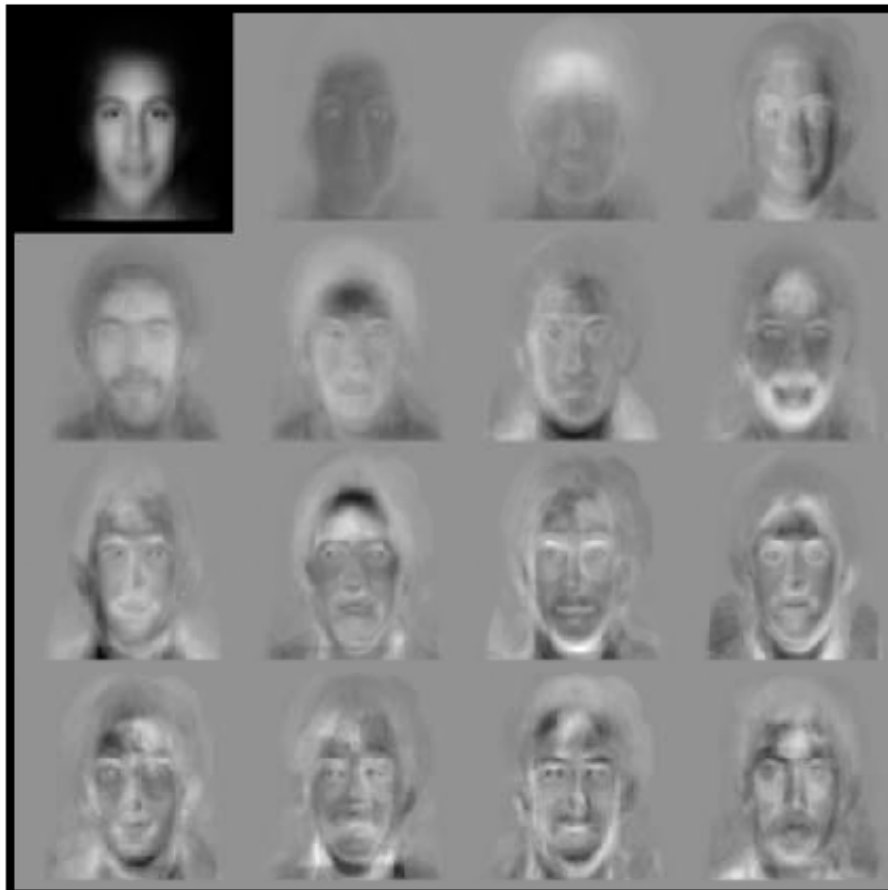
# Example of PCA



**Eigenvectors and eigenvalues of covariance matrix for  $n=1600$  inputs in  $d=3$  dimensions.**



# Example: faces



**Eigenfaces**  
from 7562  
images:

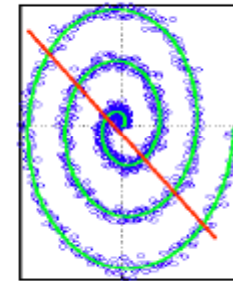
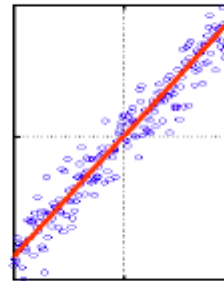
**top left image  
is linear  
combination  
of rest.**

Sirovich & Kirby (1987)  
Turk & Pentland (1991)

# Properties of PCA

- **Strengths**

- Eigenvector method
- No tuning parameters
- Non-iterative
- No local optima

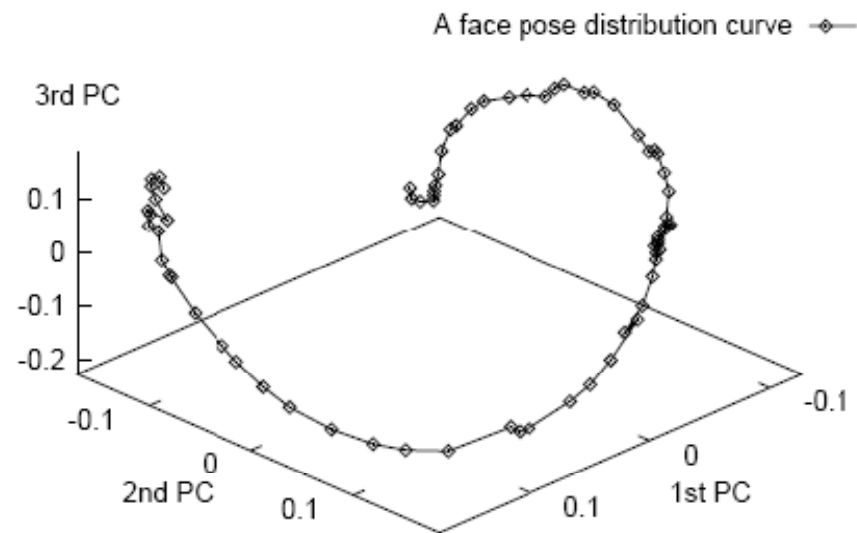


- **Weaknesses**

- Limited to second order statistics
- Limited to linear projections

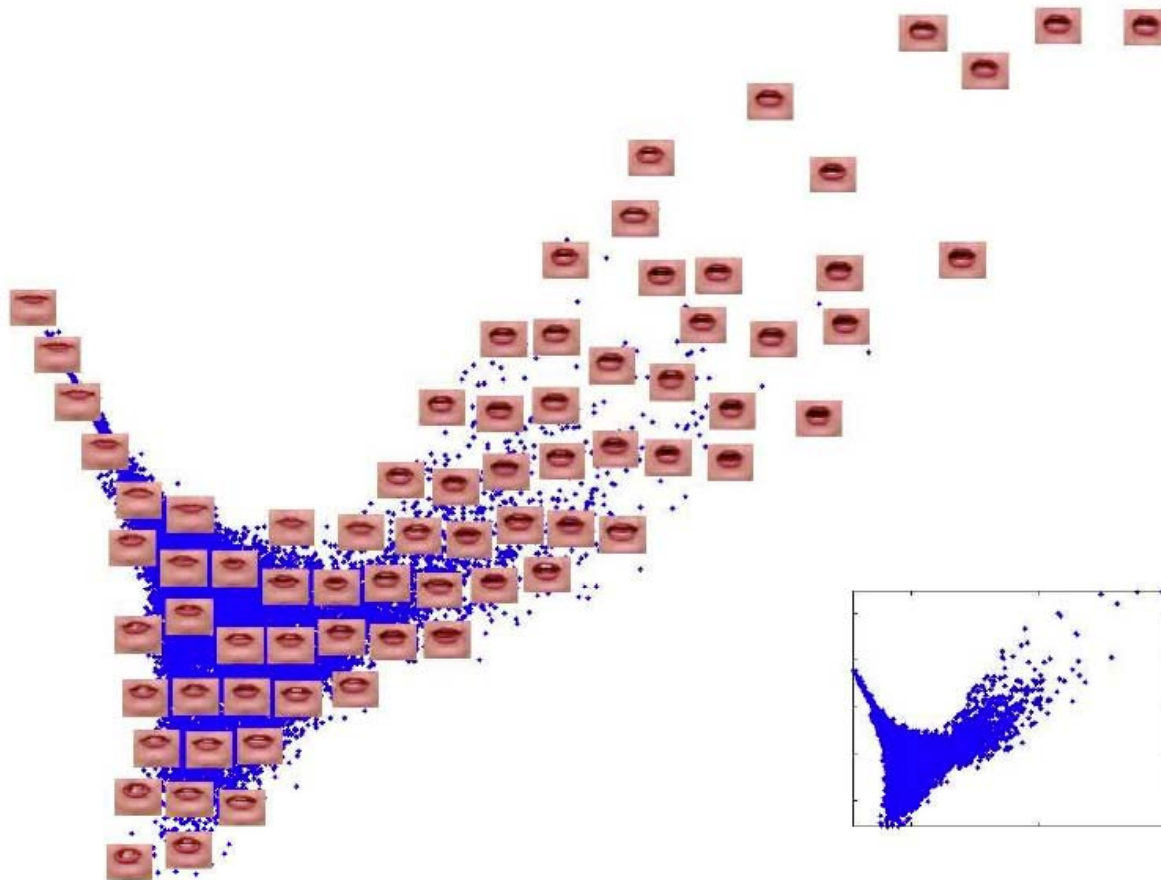
# Nonlinear Methods

Data often lies on or near a nonlinear low-dimensional curve aka manifold.



# Nonlinear Methods

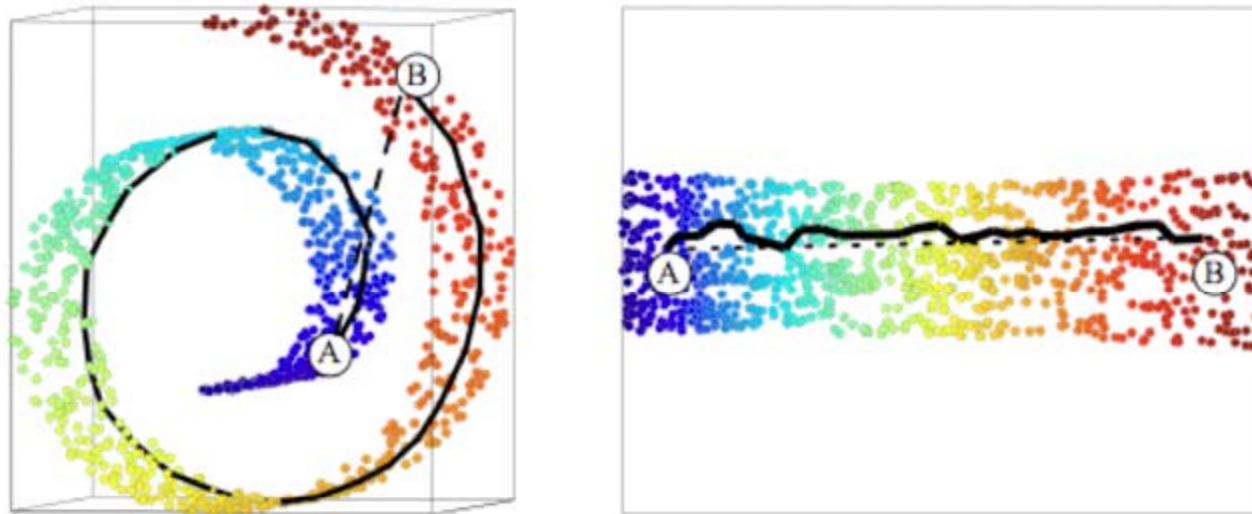
Data often lies on or near a nonlinear low-dimensional curve aka manifold.



# Isomap

Linear methods – Lower-dimensional linear projection that preserves Euclidean distances

ISOMAP basic idea – preserve geodesic distance as measured along the manifold

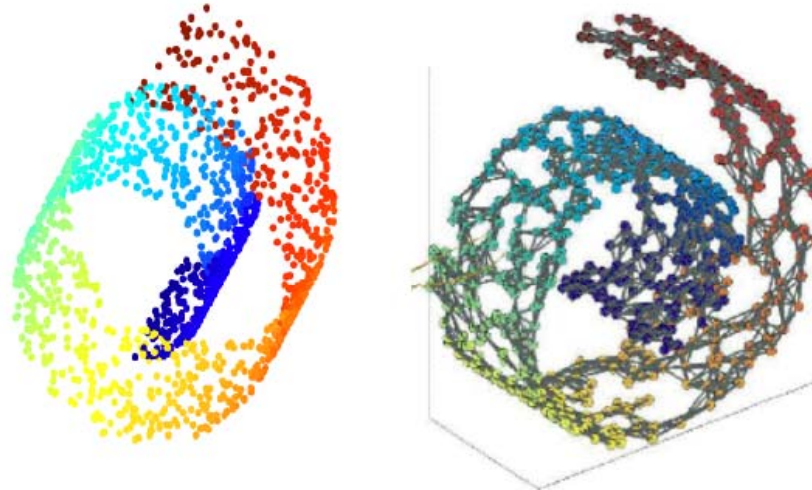


# Isomap

## Step 1. Build Adjacency graph

Vertices = Data points

Undirected edges connect nearest neighbors (k-NN, eps-NN)



- Graph is discretized approximation of manifold.
- k or eps chosen so that neighborhoods on graphs represent neighborhoods on the manifold (no “shortcuts” connect different arms of the swiss roll)

# Isomap

**Step 2.** Estimate geodesic distances by graph distances

Weight edges by local distances

Compute shortest path through the graph  $\Delta_{ij}$  (denser sampling => better estimates)

**Step 3.** Find embedding that preserves graph distances  $\Delta_{ij} \sim ||y_i - y_j||$

## MDS (Multi Dimensional Scaling)

Preserve dot products  $G_{ij}$   
(proxy for distances)

$$G_{ij} = \frac{1}{2} \left[ \sum_k (\Delta_{ik}^2 + \Delta_{kj}^2) - \Delta_{ij}^2 - \sum_{kl} \Delta_{kl}^2 \right]$$

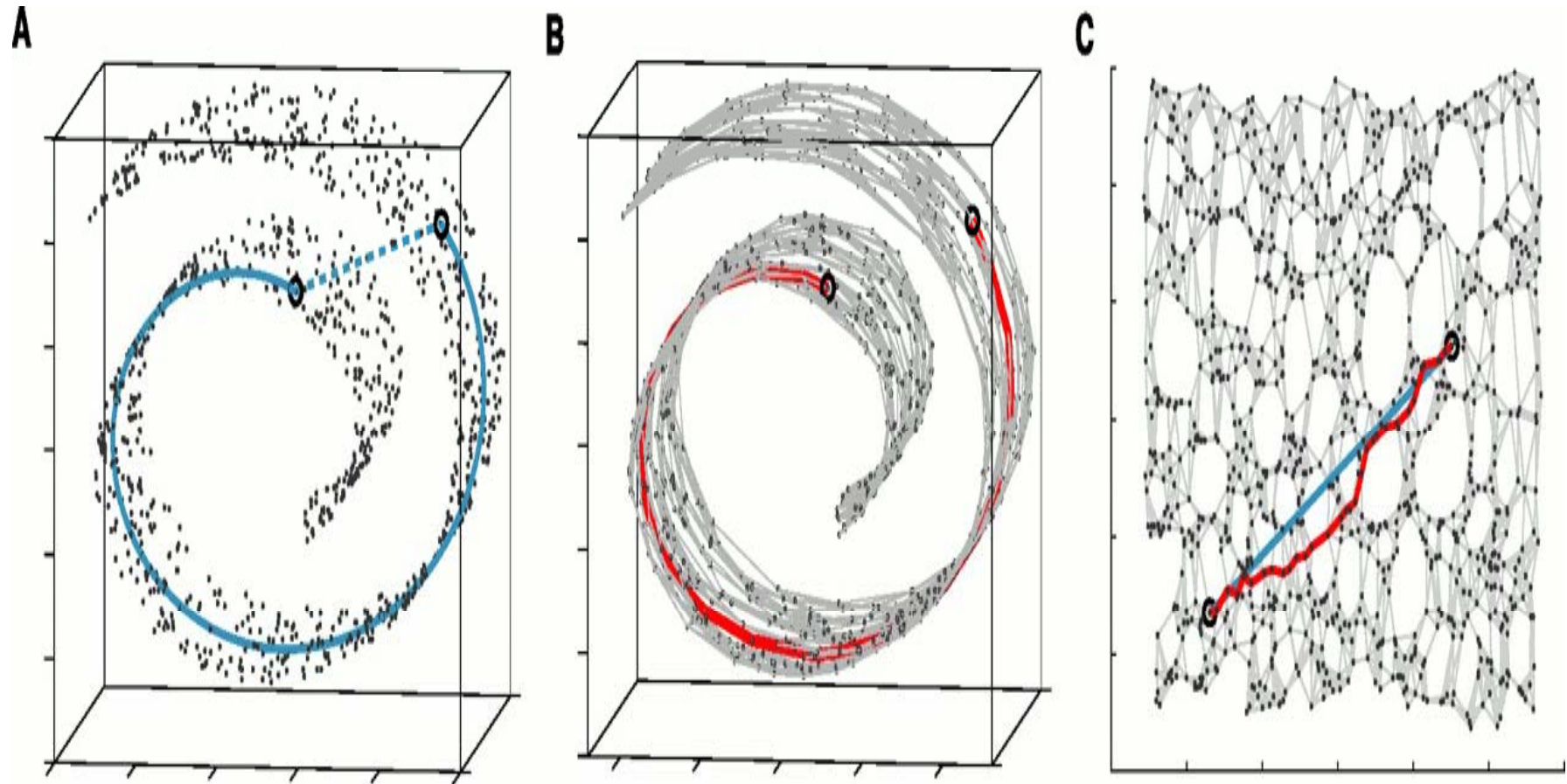
$$\arg \min_{y_1, \dots, y_n} \sum_{i,j} (G_{ij} - y_i^T y_j)$$

**Solution** - Top d eigenvectors of the Gram matrix G

Eigenvalues measure how each dimension contributes to dot product

**Same as PCA if distances are Euclidean**

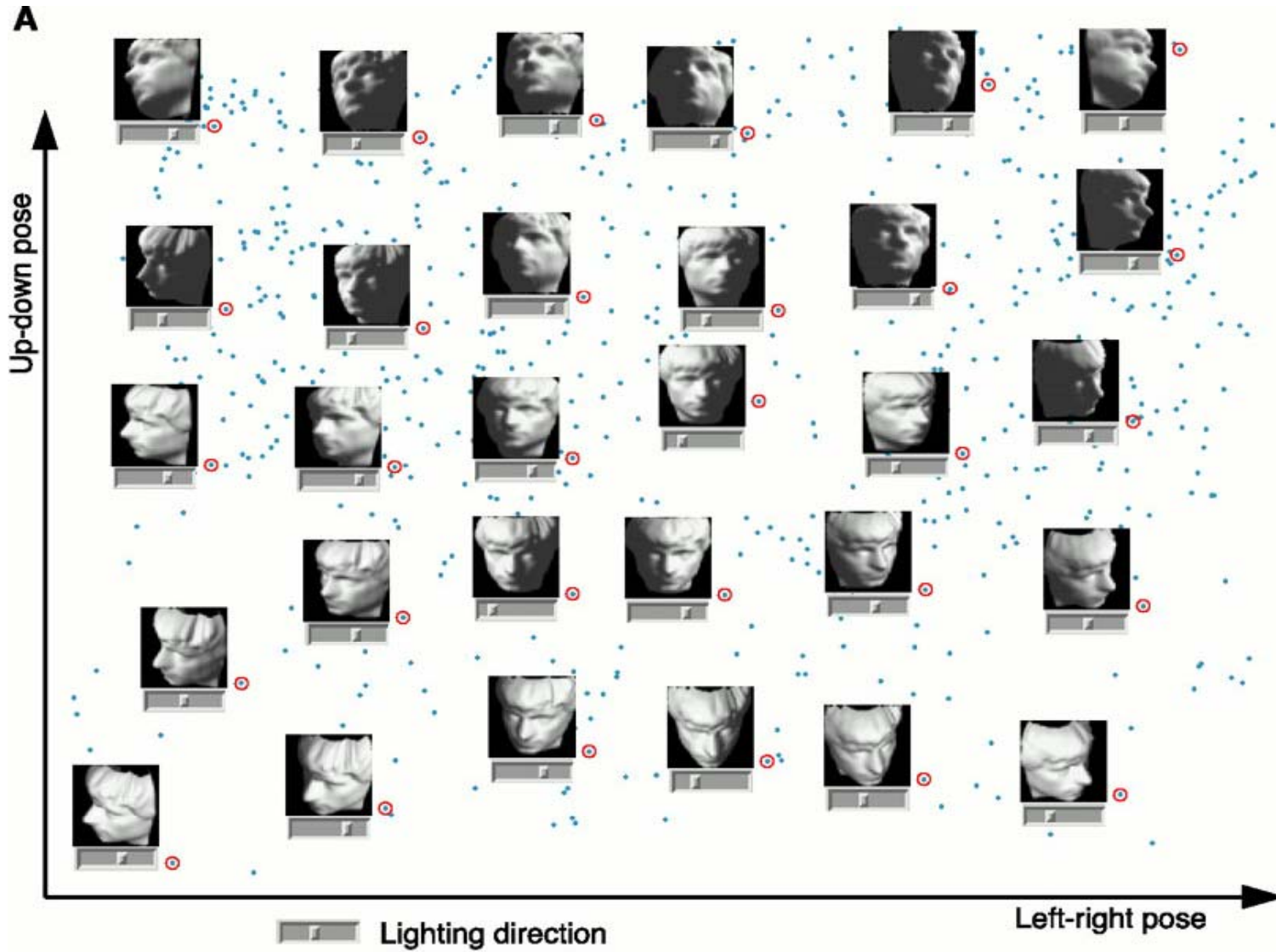
# Isomap



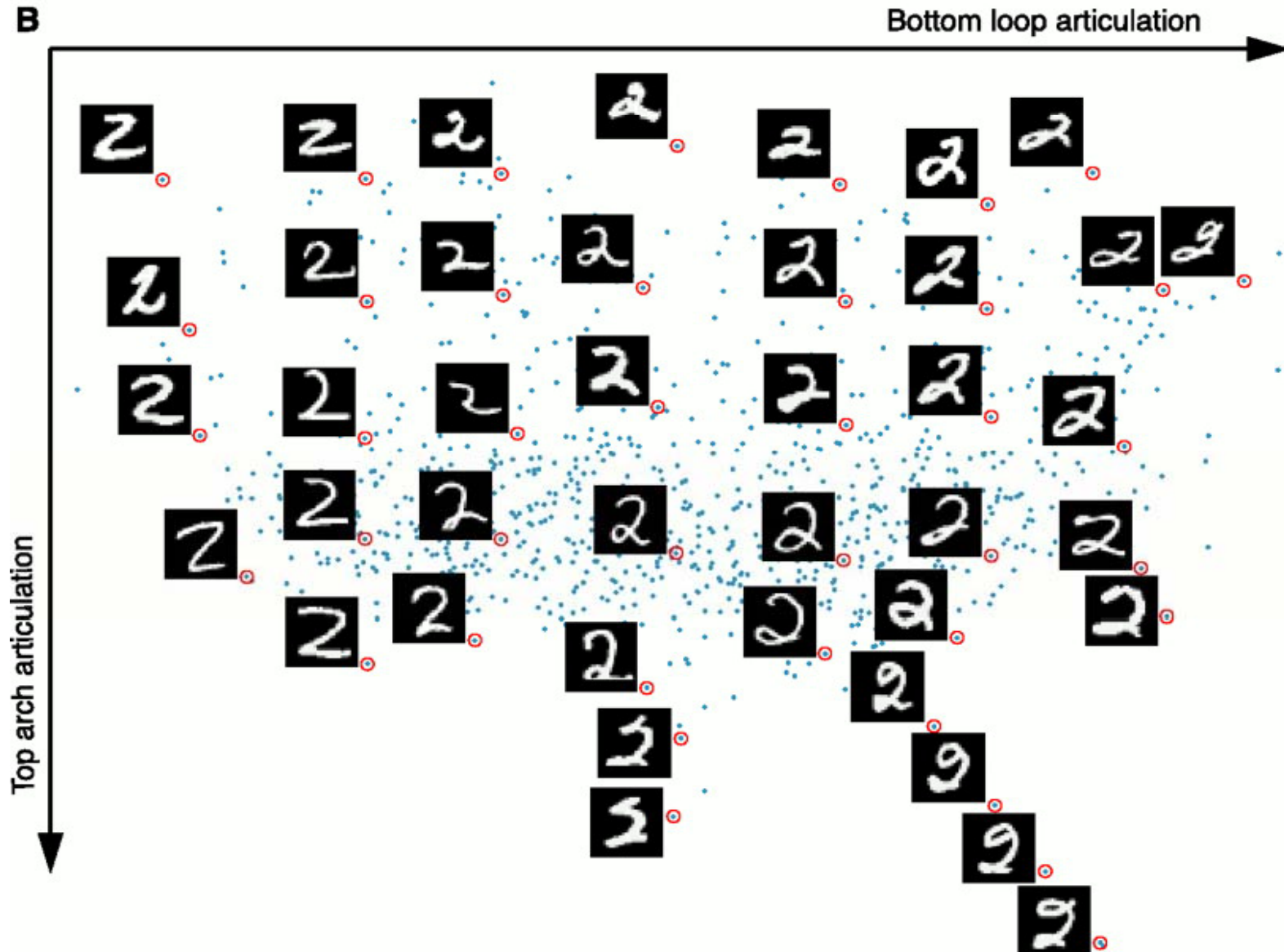
- Theoretically sound, Practically useful



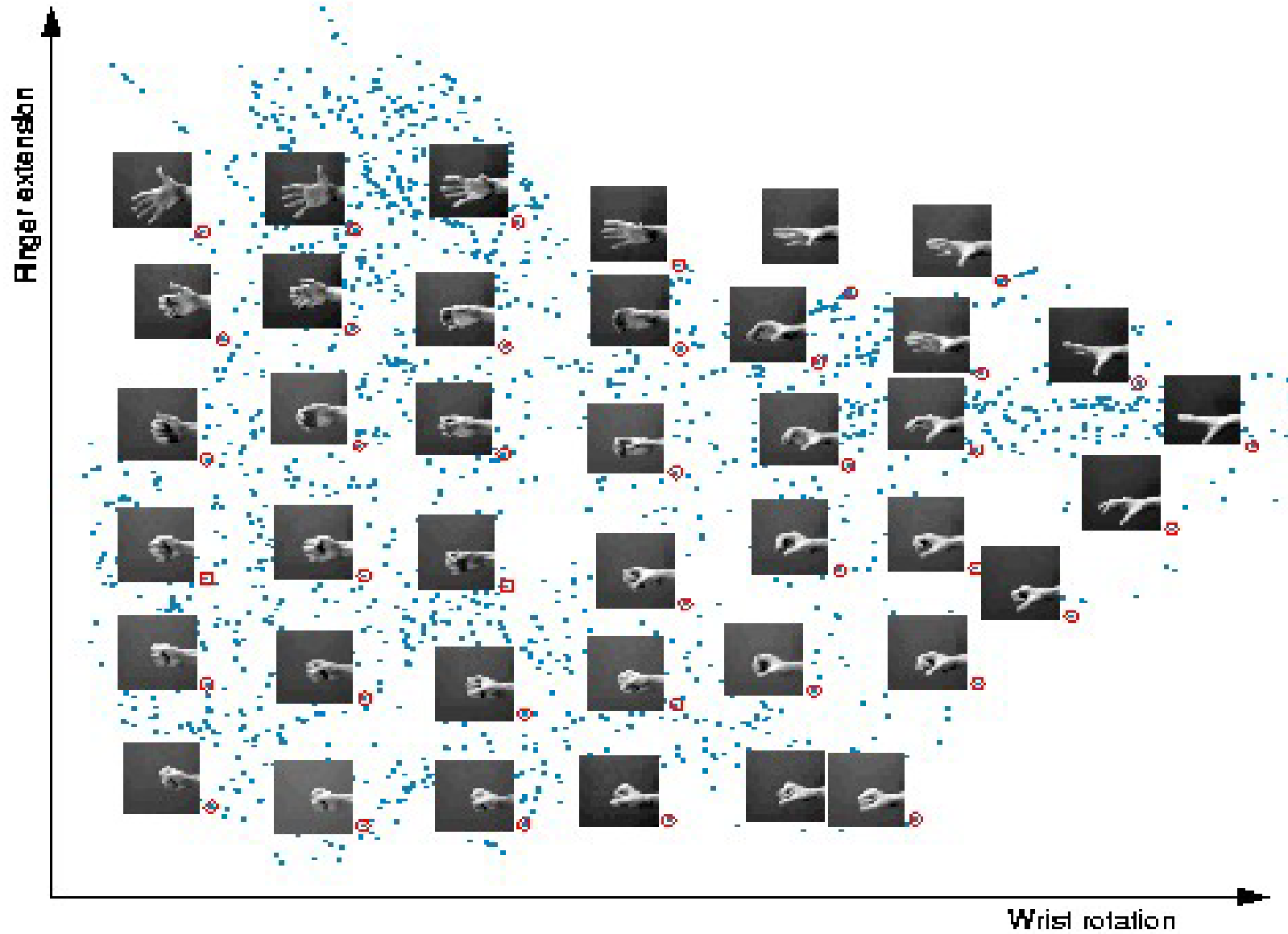
# Isomap Results



# Isomap Results



# Isomap Results



# Dimensionality Reduction Methods

- **Feature Selection** - Only a few features are relevant to the learning task
  - Score features (mutual information, prediction accuracy, domain knowledge)
  - Regularization
- **Latent features** – Some linear/nonlinear combination of features provides a more efficient representation than observed feature
  - Linear: Low-dimensional linear subspace projection
    - PCA (Principal Component Analysis),
    - MDS (Multi Dimensional Scaling),
    - Factor Analysis, ICA (Independent Component Analysis)
  - Nonlinear: Low-dimensional nonlinear projection that preserves geodesic distances along the manifold
    - ISOMAP, Kernel PCA,
    - LLE (Local Linear Embedding), Laplacian Eigenmaps
    - Data-driven linear subspaces (Wavelets)

# Some Homework for next time ...

- Think about all the (classification) algorithms we have discussed so far
  - What loss functions do they optimize?
  - What decision surfaces do they represent?
  - Pros/Cons?