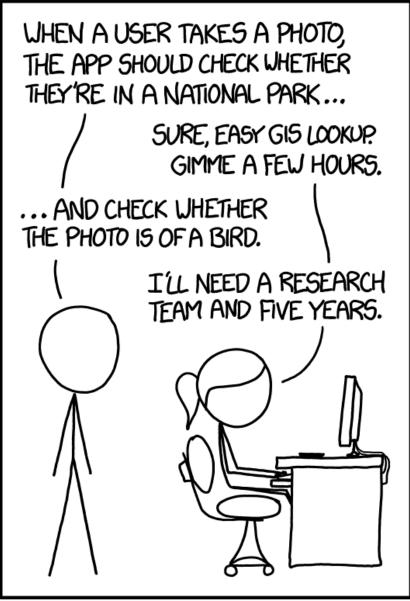
INFO 251: Applied Machine Learning

### **Dimensionality Reduction**



IN CS, IT CAN BE HARD TO EXPLAIN THE DIFFERENCE BETWEEN THE EASY AND THE VIRTUALLY IMPOSSIBLE.

### **Course Outline**

- Causal Inference and Research Design
  - Experimental methods
  - Non-experiment methods
- Machine Learning
  - Design of Machine Learning Experiments
  - Linear Models and Gradient Descent
  - Non-linear models
  - Neural models
  - Fairness and Bias
  - Practicalities and summary
  - Unsupervised Learning
- Special topics

#### Announcements

- Upcoming lectures
  - Today: Dimensionality reduction
  - Tuesday (April 19): No lecture
  - Thursday (April 21): Remote lecture: recommender systems
  - Tuesday (April 26): Applied ML, start to finish
  - Thursday (April 28): Final quiz
- Final problem set
  - Available on bCourses
  - Due May 2 (2+ weeks)
  - Warning: it's long and some problems require significant compute, so start early!

### Last Lecture: Key concepts

- Supervised vs. unsupervised learning
- Why cluster?
- k-Means clustering
- Hierarchical clustering
- Distance and linkage functions

### Outline

#### Dimensionality Reduction

- Motivation & Intuition
- Principal Component Analysis
- Example: Eigenfaces
- Other approaches to dimensionality reduction

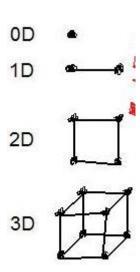
#### Motivation

- Most data is relatively low dimensional "large N, small m"
  - Housing price prediction (features are house characteristics)
  - Wage prediction (features are socio-demographics)
  - Firm performance analysis (features are...?)
  - Development metrics for countries, etc.
- Other types of problems involve high-dimensional data
  - Genome classification (features?)
  - Document analysis (features?)
  - Image recognition (features?)
  - Customer segmentation (features?)
  - Product recommendations (features?)

# Why Dimensionality Reduction?

#### Formal Reasons

- Handling the curse of dimensionality
- Reducing degrees of freedom
- Guarding against overfitting



#### Data Visualization

- In practice, visualizing and interpreting high-dimensional data is also very difficult
- We can plot data in  $\mathbb{R}^1$ ,  $\mathbb{R}^2$ , and  $\mathbb{R}^3$ , beyond that?

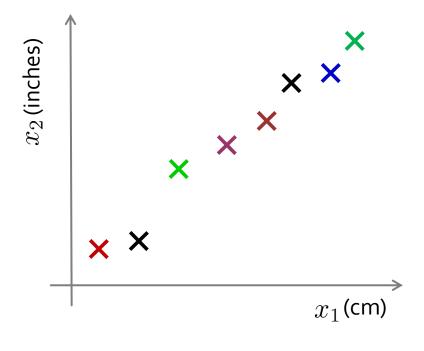
# Why Dimensionality Reduction?

- Data Interpretation
  - Often there is fundamental structure in high-dimensional data that can be represented in a small number of dimensions
    - (height, weight) => size (age, education) => maturity
    - Topics in documents
    - Authorship styles
    - People's personalities
    - Movie genres

# Why Dimensionality Reduction?

#### Data Compression

- Suppose we take measurements  $(x_{i1}, x_{i2})$  on a set of n objects  $\{x_1, ..., x_n\}$ , and our data look like the following:
- Such data are very redundant
- Why not reduce data to a single dimension where each object has one dimension  $(x_{i1})$ ?
- Seem contrived? Lots of cases where several features are highly correlated!
  - Height/weight=>size
  - GPA and GREW => school performance
- Collapsing the data can greatly increase computational efficiency



### **Summary: Motivation**

- Statistical: fewer dimensions can lead to better generalization, improve tractability
- Visualization/Interpretation: Visualize and understand structure of data

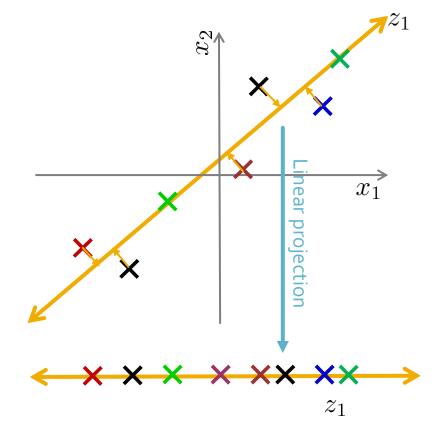
Computational: compress data to increase time/space efficiency

### **Dimensionality Reduction: Goal**

- Idea: Reduce each object from m dimensions to a corresponding object with k dimensions
  - k<m</p>
  - Do this by "collapsing" dimensions (features)
  - $x_i \in \mathbb{R}^m \Rightarrow z_i \in \mathbb{R}^k$
- Goal: Minimal "loss of information"
- Intuition: find new dimensions that still accurately represent the variation in the original data

### Dimensionality Reduction: Idea

- "Linear projection"
  - Example: Transform points from R<sup>2</sup> to R<sup>1</sup>
     with minimal loss of information
  - Idea: find common component of "length" that captures both x₁ and x₂



### Methods for Dim. Reduction

- Linear methods: PCA
  - Workhorse method of dimensionality reduction
- Nonlinear methods
  - CCA, MDS, IsoMap, Autoencoders
- Feature selection

### Outline

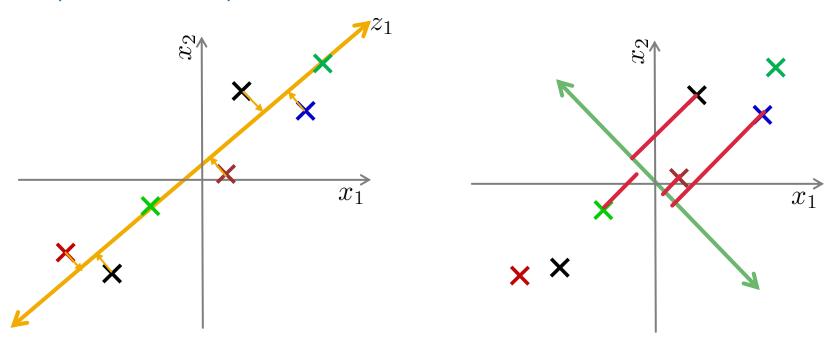
- Dimensionality Reduction
  - Motivation & Intuition
  - Principal Component Analysis
    - Intuition
    - Algorithm
    - Implementation
  - Example: Eigenfaces
  - Other approaches to dimensionality reduction

### **Principal Component Analysis**

- Most common form of dimensionality reduction
- Idea: find a lower dimensional surface onto which to project the original data so as to minimize the error

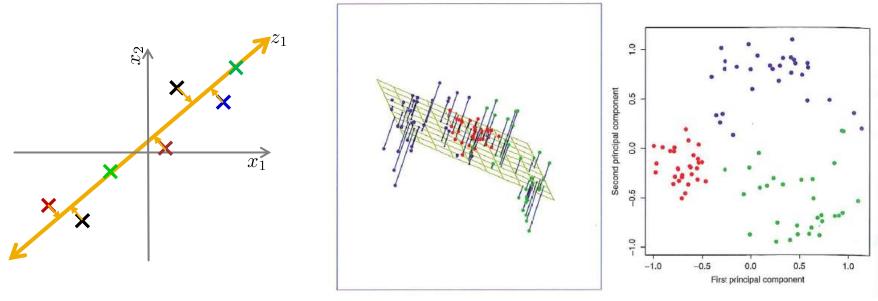
### **Principal Component Analysis**

- Graphical intuition: Minimize projection error
- What if we chose a different line?
  - 1. "Projection error" is much larger
  - 2. Projection captures less variance
  - Formally, we can prove these are equivalent statements



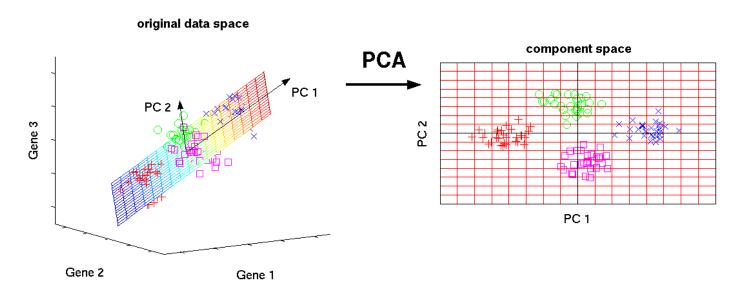
### **Principal Component Analysis**

- Example thus far: 2D->1D
- General case:
  - Find k vectors onto which the original m dimensions will be projected
  - Think of these as the k "directions" in the data
- Formally: project onto linear subspace spanned by k vectors



### PCA Example: Gene expression

- Example from Genome Sciences
  - Samples (patients) colored by treatment status
  - Original "gene space" transformed to 2-D space



### Outline

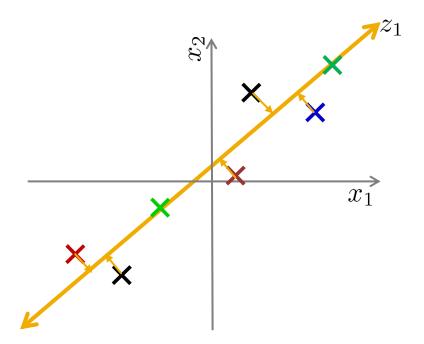
- Dimensionality Reduction
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#### **How does PCA work?**

- Goal: find vectors (principal components) and compute/project old data onto new space
  - Project m-dimensional data (in  $\mathbb{R}^m$ ) to k dimensions (in  $\mathbb{R}^k$ )
  - $x_i = (x_{11}, ..., x_{1m}) \Rightarrow (z_{11}, ..., z_{1k}), with k \le m$
  - Generally, z=f(x)
  - In *linear* dimensionality reduction (matrix notation)  $z = U^T x$
- How to choose transformation (U)?
  - 1. Intuitive, inefficient way
  - 2. Efficient, analytic solution

## PCA Algorithm (the wrong way)

- Intuitive, brute force algorithm:
- 1. Fit a squared-distance minimizing line to the data
- 2. Compute residuals normal to the line
- 3. Iterate



### PCA Algorithm (intuition)

#### Just like OLS:

- Gradient descent (MLE) yields "correct" solution
- Analytic solution is far faster: (X'X)<sup>-1</sup>(X'Y)

#### With PCA

- Analytic solution uses linear algebra on X (not Y!)
- Formal derivation is tricky, but: the principal components of a set of data can be found from the (first k) eigenvalues and eigenvectors of the covariance matrix of the data

#### Summary:

- Compute covariance matrix of data
- 2. Compute **eigenvectors** of covariance matrix
- 3. Compute **projections**

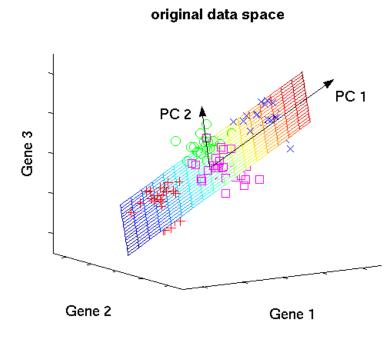
## PCA algorithm

- Compute covariance matrix of data X
  - Mean and covariance of data
  - Mean of data tells you "where" the cloud is
  - Covariance tells you "which way it's pointing"

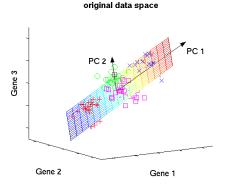
$$Q = \frac{1}{n} \sum_{i=1}^{n} (x_i)(x_i)^T$$

$$q_{j,m} = \frac{1}{n-1} \sum_{i=1}^{n} (x_{i,j} - \bar{x}_j)(x_{i,m} - \bar{x}_m)$$

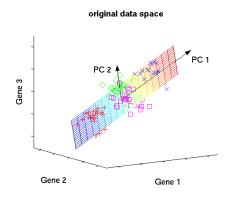
- Compute eigenvectors
- 3. Compute **projections**



- Compute covariance matrix of data
- 2. Compute **eigenvectors** of covariance matrix
  - Singular value decomposition (SVD) of (m x n) matrix Q is:  $Q=U\sum V$ 
    - U is m x m unitary matrix;  $\sum$  is m x n diagonal matrix of singular values; V is n x n unitary matrix
  - Columns of U are the eigenvectors (of  $QQ^T$ )
    - Columns of V are eigenvectors of Q<sup>T</sup>Q
  - ullet is diagonal matrix, with eigenvalues on the diagonal
  - Eigenvectors represent the directions of greatest variation of the data
  - **Eigenvalues** tell you how much variation is captured by the corresponding eigenvector
- 3. Compute **projections**



- 1. Compute **covariance matrix** of data
- 2. Compute **eigenvectors** of covariance matrix
  - Singular value decomposition (SVD) of (m x n) matrix Q is: Q=U∑V
  - Columns of U are the eigenvectors (of QQ<sup>T</sup>)
- 3. Compute **projections** 
  - Denote by  $U_k$  the matrix comprising the first k columns of U
  - The projection of X is  $XU_k$  (we call this "Z")
  - In other words, multiply the original data by the matrix of eigenvectors, this maps it to k-space



- Summary:
  - 1. Compute **covariance matrix** of data
  - 2. Compute **eigenvectors**
  - 3. Compute **projections**

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  - Motivation & Intuition
  - Principal Component Analysis
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    - Algorithm
    - Implementation
  - Example: Eigenfaces
  - Other approaches to dimensionality reduction

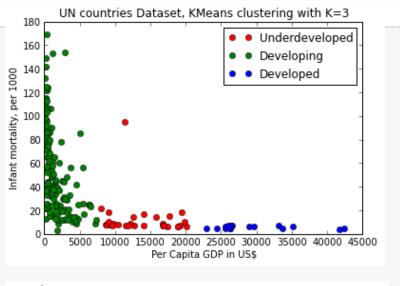
### **PCA Implementation & Use**

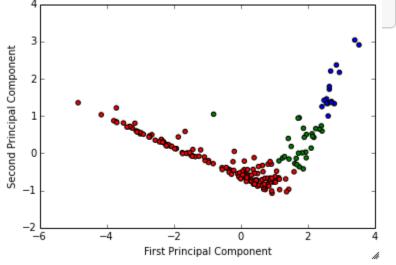
- Before starting: Standardize your features!
  - Replace each  $x_{im}$  with  $\frac{x_{im}-\mu_m}{s_m}$  , where  $\mu_m=\frac{1}{k}\sum_{i=1}^k x_{im}$

### **PCA** in Python

```
from sklearn.decomposition import PCA
estimator = PCA(n_components=2)
X_pca = estimator.fit_transform(normalize(X))

colors = ['red', 'blue', 'green']
for i in xrange(len(colors)):
    px = X_pca[:, 0][c == i]
    py = X_pca[:, 1][c == i]
    plt.scatter(px, py, c=colors[i])
    plt.xlabel('First Principal Component')
    plt.ylabel('Second Principal Component')
```



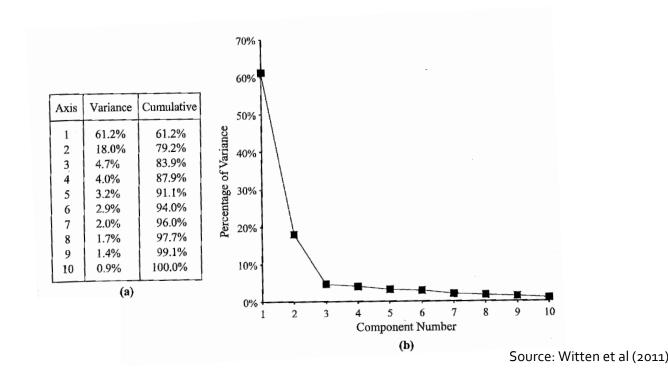


### How many components?

- How many components?
  - Some art and some science
  - Often there is a "natural" breaking point
- Depends on amount of variation you want to explain!
- "Retained variance"
  - Average squared projection error / total variation
  - Call  $\hat{x}_i$  our reconstructed  $x_i$  (like predicted values from a regression)
    - Think of this as "inflation" of projected/reduced  $z_i \Rightarrow U z_i$ , where U are eigenvectors of the covariance matrix Q
    - Variance lost:  $\frac{\frac{1}{n}\sum_{i=1}^{n}|x_i-\hat{x}_i|^2}{\frac{1}{n}\sum_{i=1}^{n}|x_i|}$  (typically between 0.01 and 0.10)
  - Rule of thumb: retain 99% (or 95% or 90%) of variance

### Example

- Example: Components of a 10-dimensional space
  - Each component is an axis, table indicates share of variance accounted for. 3
     PC's account for roughly 84% of variance of original data



### Interpreting PCA

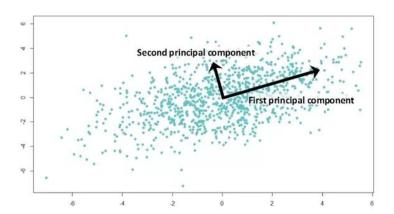
Do the new features (the principal components) have any meaning?

$$PC_1 = U_{11}X_1 + \dots + U_{1m}X_m$$

$$\dots$$

$$PC_k = U_{k1}X_1 + \dots + U_{km}X_m$$

Each component is a linear combination of the original features



### Common applications of PCA

- Computing wealth from assets
- Genetics
  - E.g. compress M=>K, then look at individual contributions to the principal components
- Outlier detection
- "Cocktail party problem"
- Human posture classification
- Data pre-processing and compression

## Another important point

- One of most common uses of PCA is to increase efficiency of other learning algorithms, e.g. PCA then regression
- Important: Best practice is to train PCA algorithm on training data, not on validation/testing data!
  - The transformation matrix (U) is *learned* on the data, and will be different depending on the data used to train it

# Image Compression Example

Eigenvector Number	Cumulative proportion of variance
1	0.5160
2	0.6979
3	0.7931
4	0.8794
5	0.9130
6	0.9378
7	0.9494
8	0.9596
9	0.9678
10	0.9732

Table 2: Cumulative variance accounted for by PCs

102.4:1 compression 2 principal components



17.7:1 compression 14 principal components



6.3:1 compression 40 principal components



2.1:1 compression 120 principal components



39.4:1 compression 6 principal components



12.5:1 compression 20 principal components



4.2:1 compression 60 principal components



1.7:1 compression 150 principal components



24.4:1 compression 10 principal components



8.4:1 compression 30 principal components



2.8:1 compression 90 principal components



1.4:1 compression 180 principal components



Figure 10: The visual effect of retaining principal components

#### Outline

- Dimensionality Reduction
  - Motivation & Intuition
  - Principal Component Analysis
  - Example: Eigenfaces
  - Other methods for dimensionality reduction

## PCA Example: Eigenfaces

- Turk and Pentland (1991), "Eigenfaces for Facial Recognition",
   Journal of Cognitive Neuroscience 3 (1)
- Goal: 2-D Face recognition
  - Efficient, simple, and accurate face recognition in a constrained environment

# PCA Example 2: Eigenfaces

- Initialization (training):
  - Use initial set of facial images as training data
    - 256 pixel X 256 pixel array (65,536 pixels; 8-bit intensity)
  - Calculate PC's (eigenvectors, aka "eigenfaces"), retain components with largest eigenvalues
  - Project original data onto "face space"
- Recognition:
  - Project unknown image onto "face space"
    - Remember this is just the lower-dimensional subspace formed by the principal components of the original data
  - Classify projection as known face, unknown face, or not face

## Eigenfaces: examples

Original Faces and the "average face"





## Eigenfaces

#### Eigenfaces and projections of into "face space"

- Each eigenvector/eigenface is a new dimension of the face space (65536 x 1)
- The eigenvectors are used to transform a new face image into its eigenface components (in this case, a 7-dimensional space)
- In other words: a "real" face is a linear combination of those 7 eigenfaces

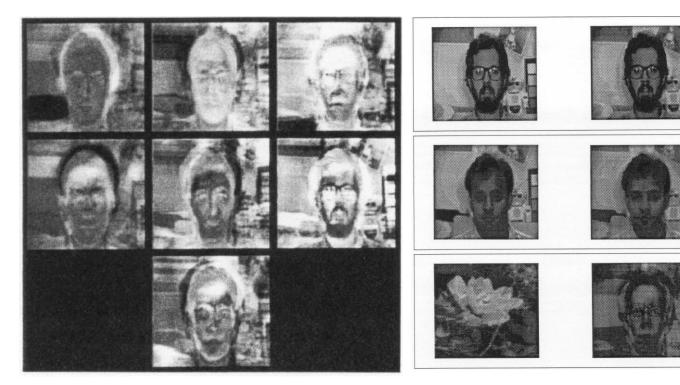


Figure 4. Three images and their projections onto the face space defined by the eigenfaces of Figure 2. The relative measures of distance from face space are (a) 29.8, (b) 58.5, (c) 5217.4. Images (a) and (b) are in the original training set.

## Eigenfaces

- Recognition: Nearest neighbors!
  - Known images projected onto face space
    - Compute "average eigenface" for each person
  - Unknown images projected onto face space
    - Choose "average eigenface" that minimizes distance (in face space!) to projection
- Who cares?
  - Fast and simple: Real-time recognition
  - Facial reconstruction



#### Outline

- Dimensionality Reduction
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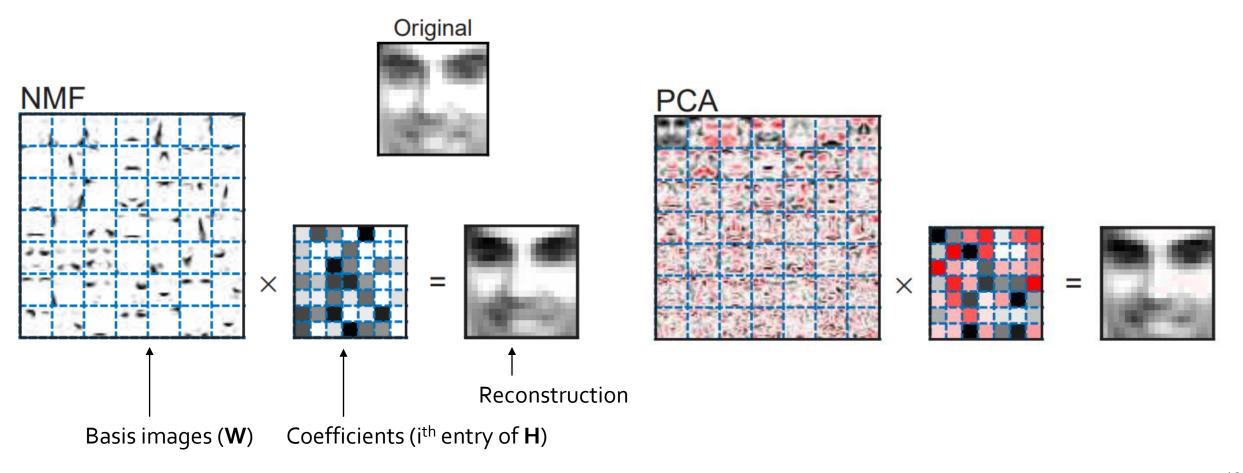
### Other methods for dimensionality reduction

- PCA: Popular, fast, linear
- Review of other common methods
  - Non-negative matrix factorization
  - Multi-dimensional scaling
  - Kernel methods
  - Isomap
  - Autoencoders

## Non-negative matrix factorization

- Non-negative matrix factorization
  - Similar to PCA, but data and components are non-negative
    - Useful for modelling non-negative data, e.g. images
  - Data matrix X approximated by W H
    - W: basis elements
    - H: basis coordinates of data points
  - Basis elements often have interpretation
    - e.g., face features (images), topics (documents)
    - Compare to principal components, which are orthogonal vectors that cumulatively (and greedily) explain variation in data

# Non-negative matrix factorization



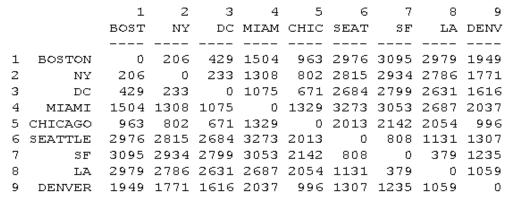
# **Multi-Dimensional Scaling**

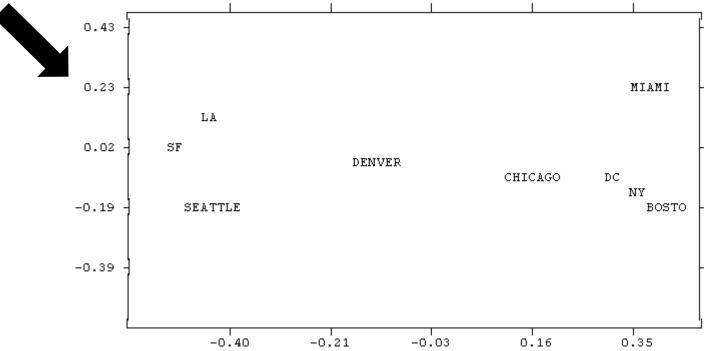
- Multi-Dimensional Scaling (also linear)
  - Idea: preserve inter-point distances, rather than preserving total variance
  - Formally: call  $d_{ij}$  the distance between two points in  $\mathbb{R}^m$  and denote by  $\delta_{ij}$  the distance in  $\mathbb{R}^k$ , goal is to minimize cost:

$$\sum_{i,j} \left( \frac{d_{ij} - \delta_{ij}}{\delta_{ij}} \right)^2$$

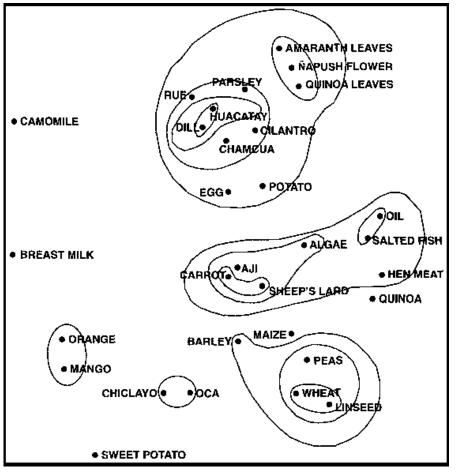
- Useful in situations where similarity and distance is paramount, or when original coordinates of data are unknown
  - e.g. market research where subjects specify similarity between objects.
- Really good for representing/visualizing data in 2D

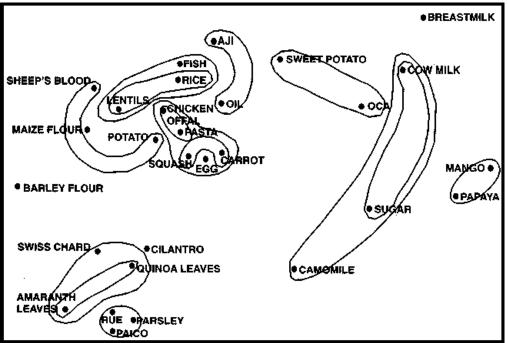
# **Multi-Dimensional Scaling**





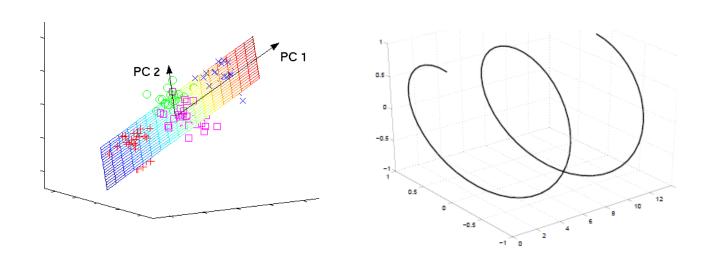
# **Multidimensional Scaling**





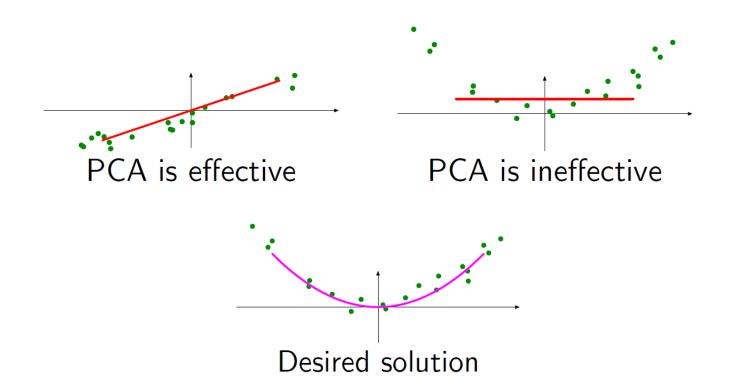
### Non-linear methods

- PCA and MDS are linear
  - Work well when high-D data sit on a lower-D hyperplane
  - Not as effective if data sit on a curved manifold:



### **Kernel Methods**

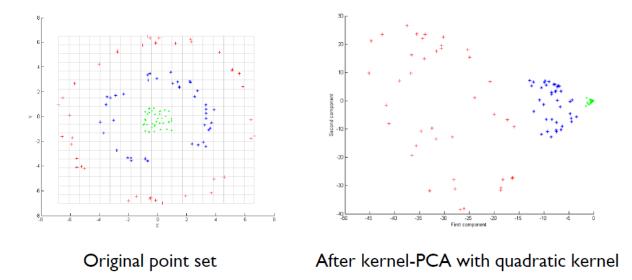
- Simple solution: Kernel
  - Apply kernel transformation to original data, then run PCA on resultant dataset



#### **Kernel Methods**

#### Kernel PCA

- PCA finds linear projections, but sometimes projection space is nonlinear
- Kernel PCA applies kernels to allow for differently shaped subspaces



#### Non-linear methods

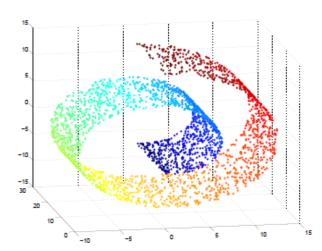
- Other methods
  - Fisher Discriminant Analysis
  - Also locally-linear embedding, Laplacian eigenmaps, self-organizing maps, manifold alignment, curvilinear component analysis, etc.
  - Isomap

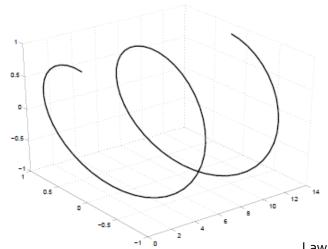
#### Isomap

- Isometric Feature Mapping
  - Example manifold: the "Swiss Roll"
  - PCA finds hyperplane projections

Isomap projects original data onto a non-linear, high-dimensional

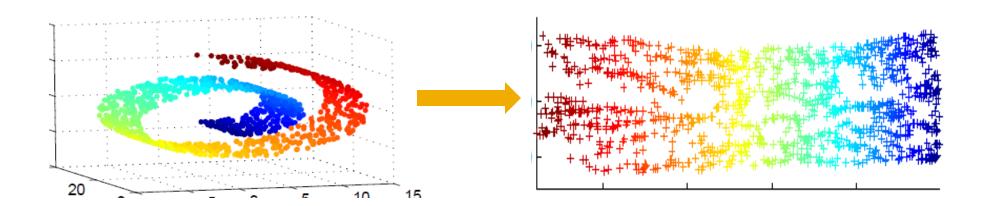
manifold.





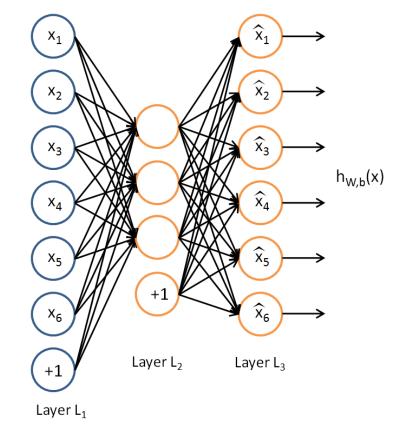
#### Isomap

- Isomap: "MDS with geodesic distance"
  - 1. Construct neighborhood graph G, only add edges if (Euclidean) distance  $< \varepsilon$
  - Compute distance matrix using geodesic (shortest path) distance in G, (using e.g. Floyd's or Djikstra's algorithm)
  - 3. Use MDS on **G** to find points in low-dimensional Euclidean space whose interpoint distances match the distances found in step 1



#### **Autoencoders**

- Auto-encoders map inputs to inputs (i.e., it approximates the identity function):  $\widehat{y_i}(x_i) = x_i$ 
  - (See deep learning lecture notes)



# This lecture: Key Concepts

- Why do dimensionality reduction?
- Principal component analysis nuts and bolts
- Eigenvalues and retained variance
- Eigenfaces
- Multidimensional scaling
- IsoMap ("MDS with geodesic distance")