

Part 1: PCA & MDS

COS 323

Last Time

- How do we solve least-squares...
 - without incurring condition-squaring effect of normal equations ($A^T A x = A^T b$)
 - when A is singular, “fat”, or otherwise poorly-specified?
- QR Factorization
 - Householder method
- Singular Value Decomposition
- Total least squares

Dimensionality Reduction

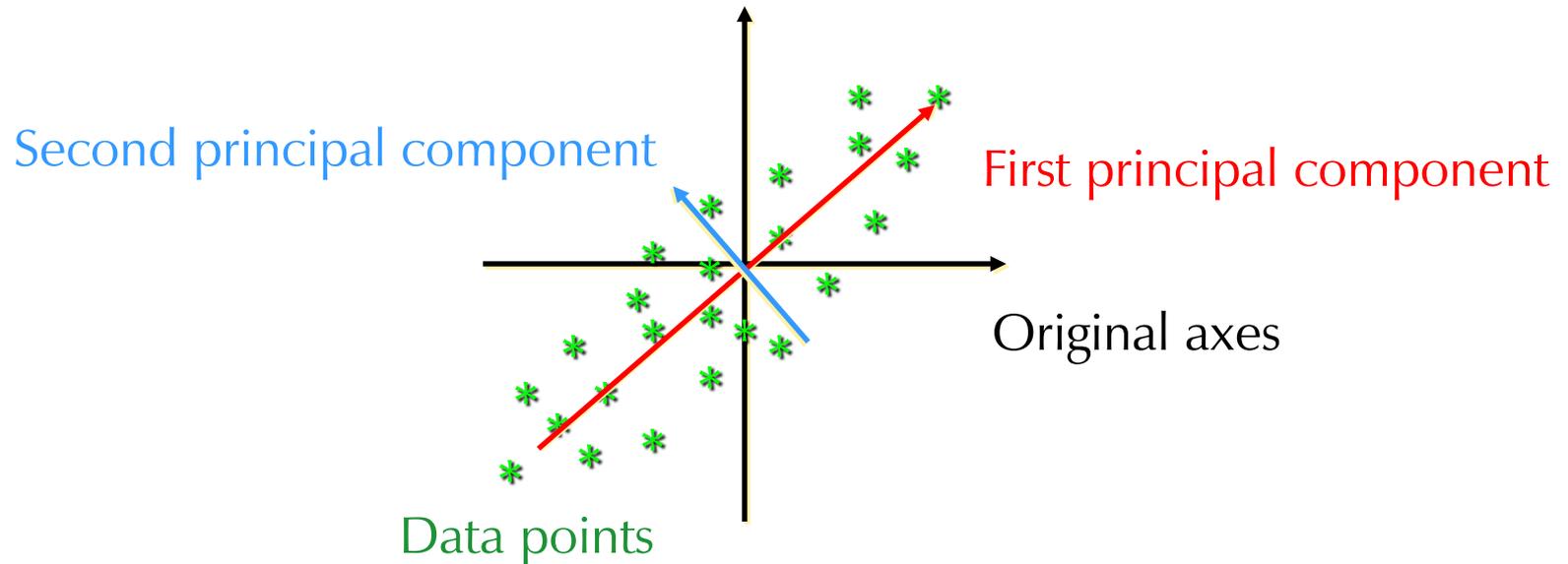
- Map points in high-dimensional space to lower number of dimensions
- Preserve structure: pairwise distances, etc.
- Useful for further processing:
 - Less computation, fewer parameters
 - Easier to understand, visualize

SVD for rank- k approximation

- \mathbf{A} is $m \times n$ matrix of rank $> k$
- Suppose you want to find best rank- k approximation to \mathbf{A}
- Take SVD: $\mathbf{A} = \mathbf{U}\mathbf{W}\mathbf{V}^T$
- Set all but the largest k singular values of \mathbf{W} to 0
- Can form compact representation by eliminating columns of \mathbf{U} and \mathbf{V} corresponding to zeroed w_i

Principal Components Analysis (PCA)

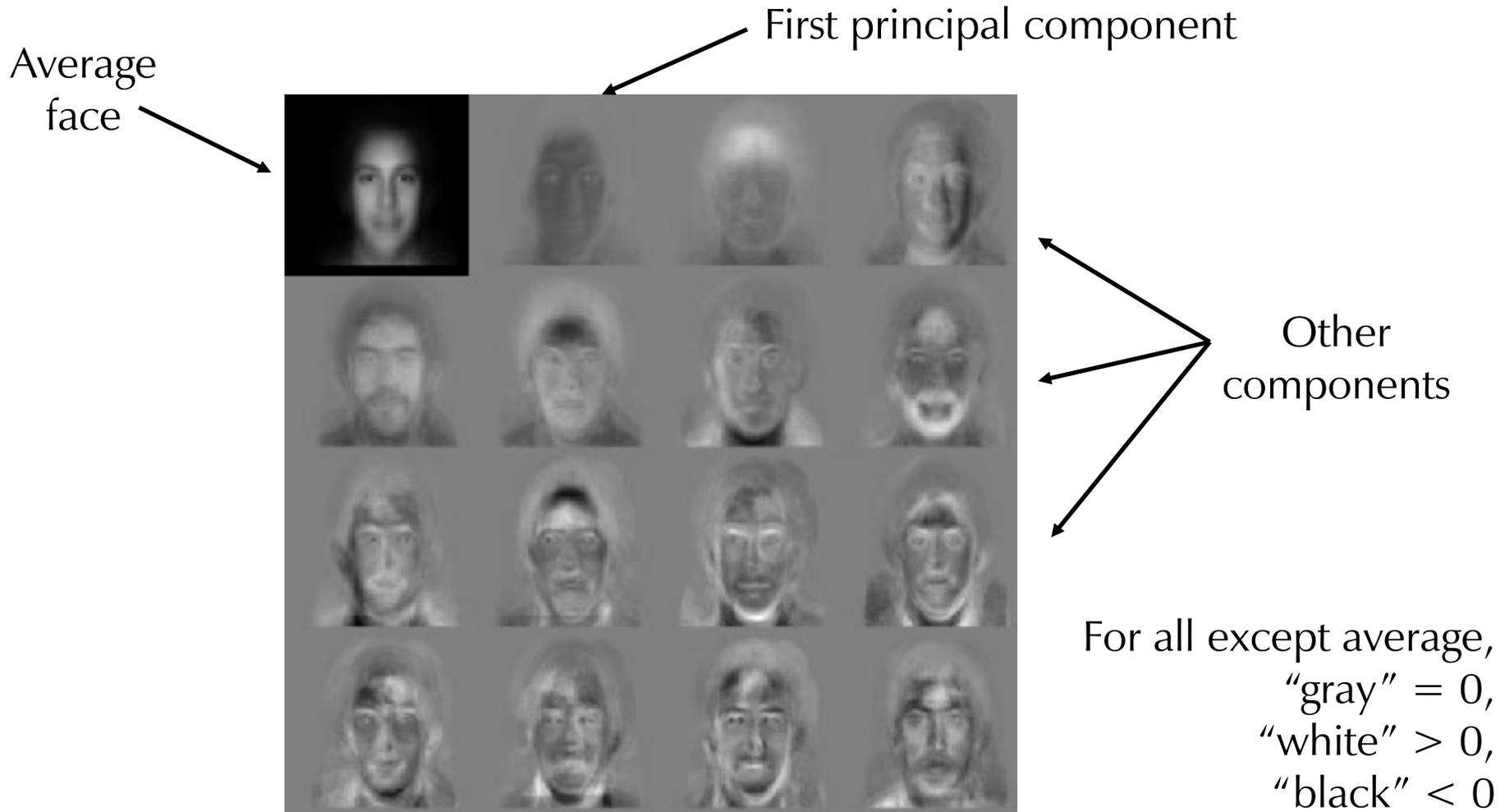
- Approximating a high-dimensional data set with a lower-dimensional linear subspace
- Also converts possibly-correlated attributes into uncorrelated attributes



SVD and PCA

- Data matrix with points/examples as rows
- Center data by subtracting mean (“whitening”)
- Compute SVD
- Columns of \mathbf{V}_k are principal components
- Value of w_i gives importance of each component

PCA on Faces: “Eigenfaces”



Uses of PCA

- Compression: each new image can be approximated by projection onto first few principal components
- Recognition: for a new image, project onto first few principal components, match feature vectors
- Generation: Adjust contributions of a few principal components to generate new plausible data points

PCA for Relighting

- Images under different illumination



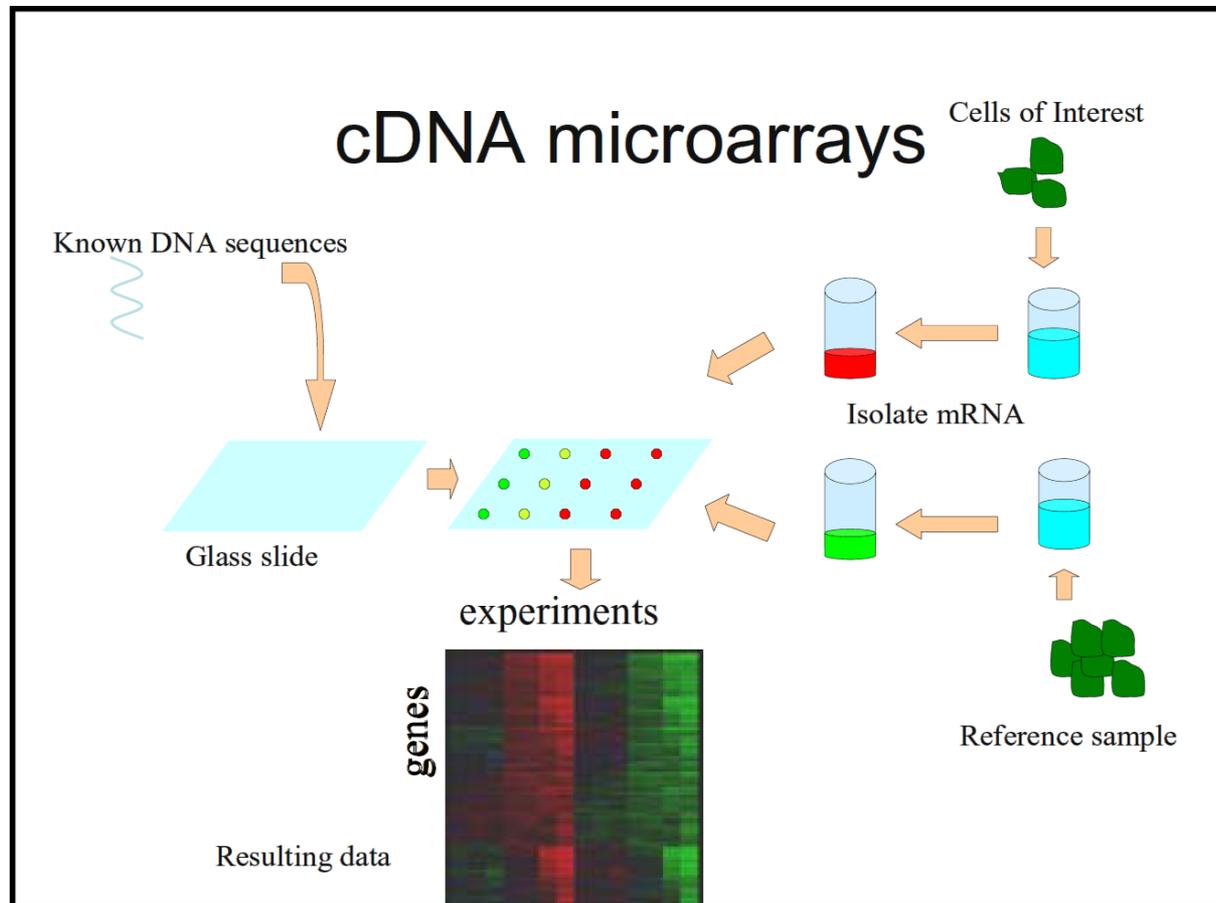
PCA for Relighting

- Images under different illumination
- Most variation captured by first 5 principal components – can re-illuminate by combining only a few images



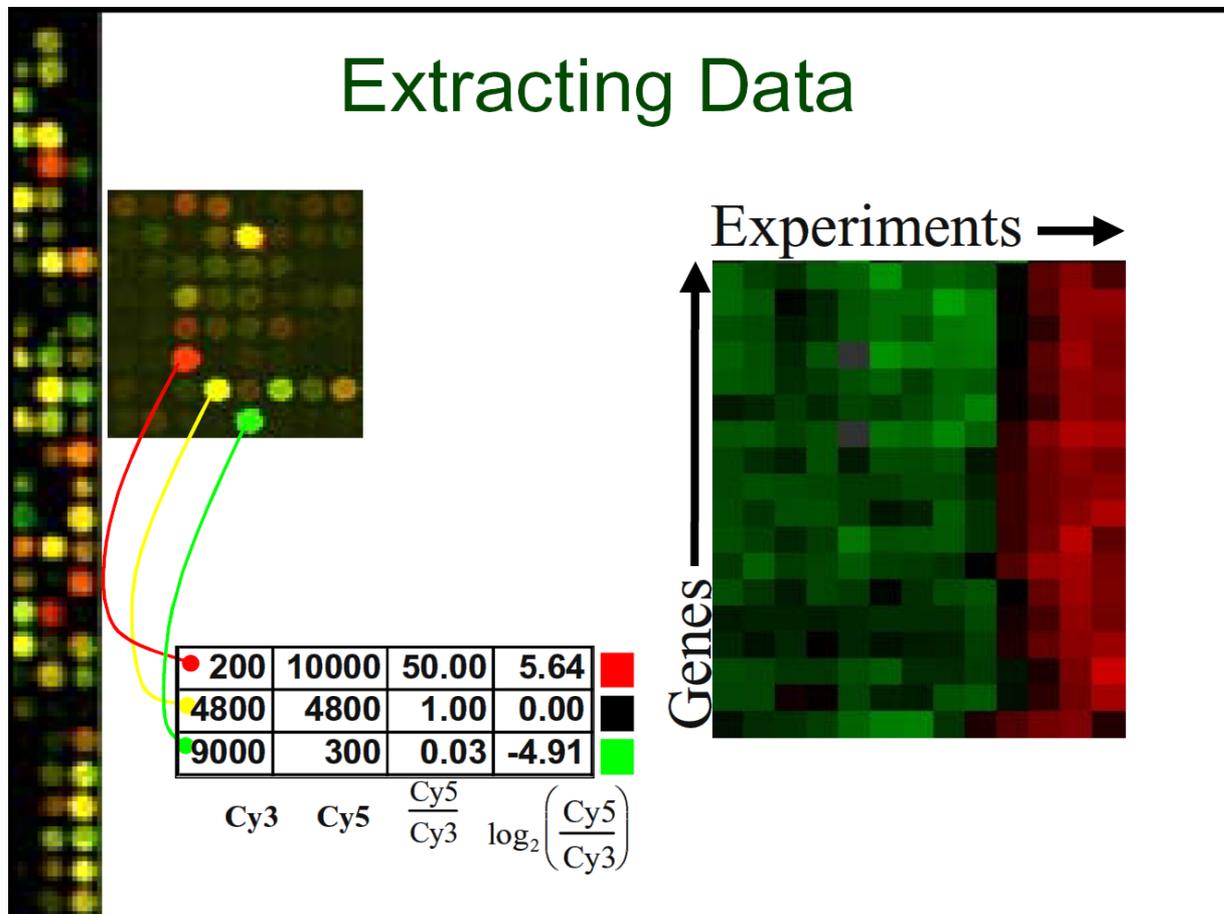
PCA for DNA Microarrays

- Measure gene activation under different conditions



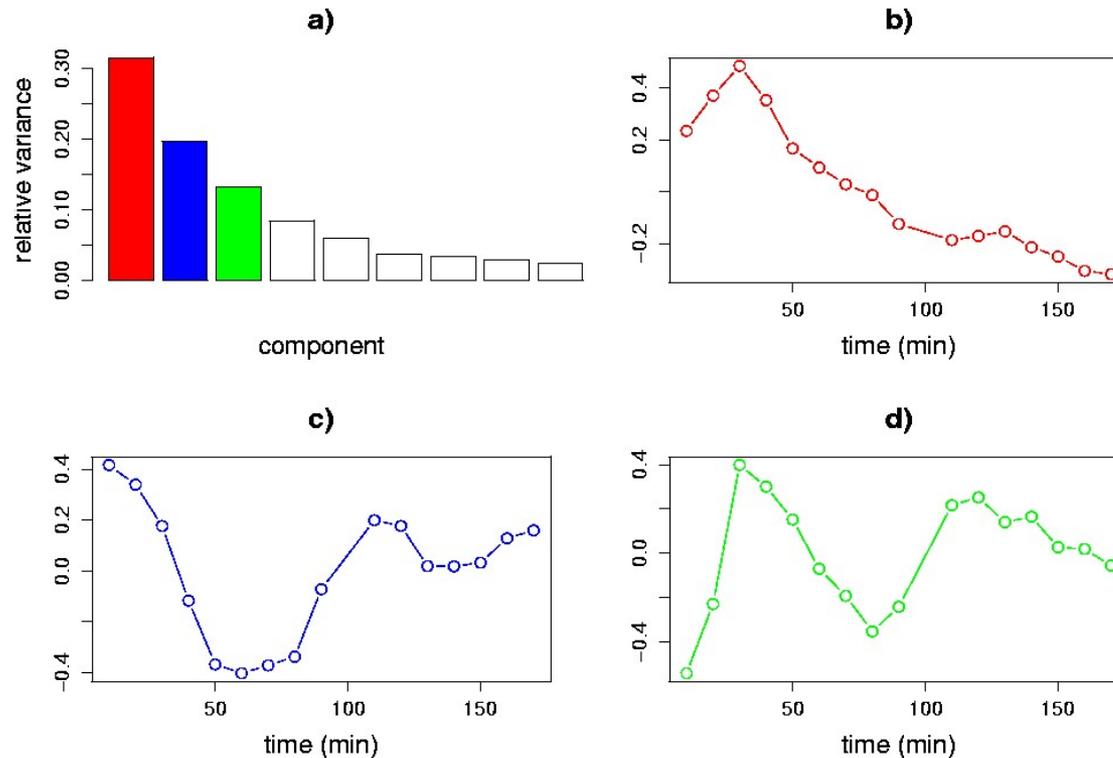
PCA for DNA Microarrays

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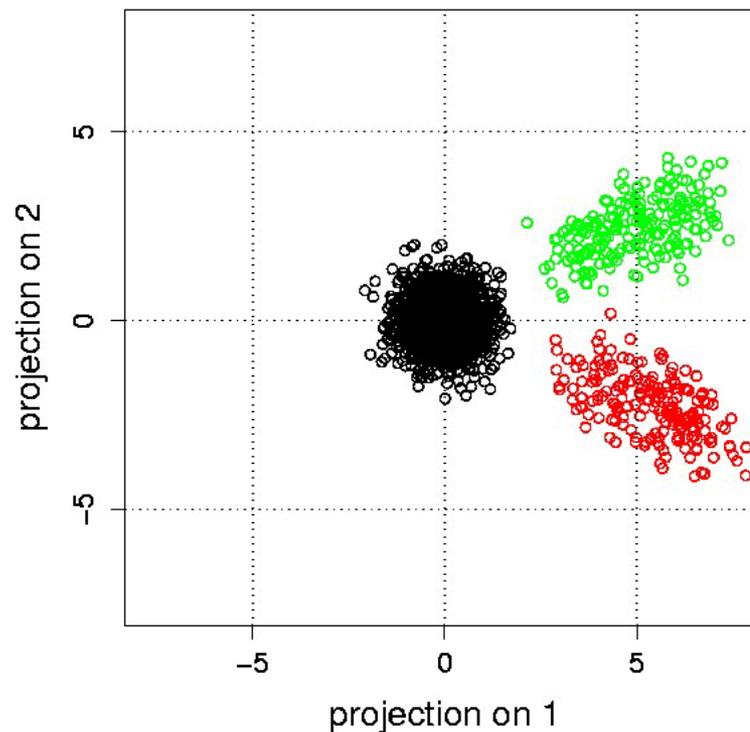
PCA for DNA Microarrays

- PCA shows patterns of correlated activation
 - Genes with same pattern might have similar function



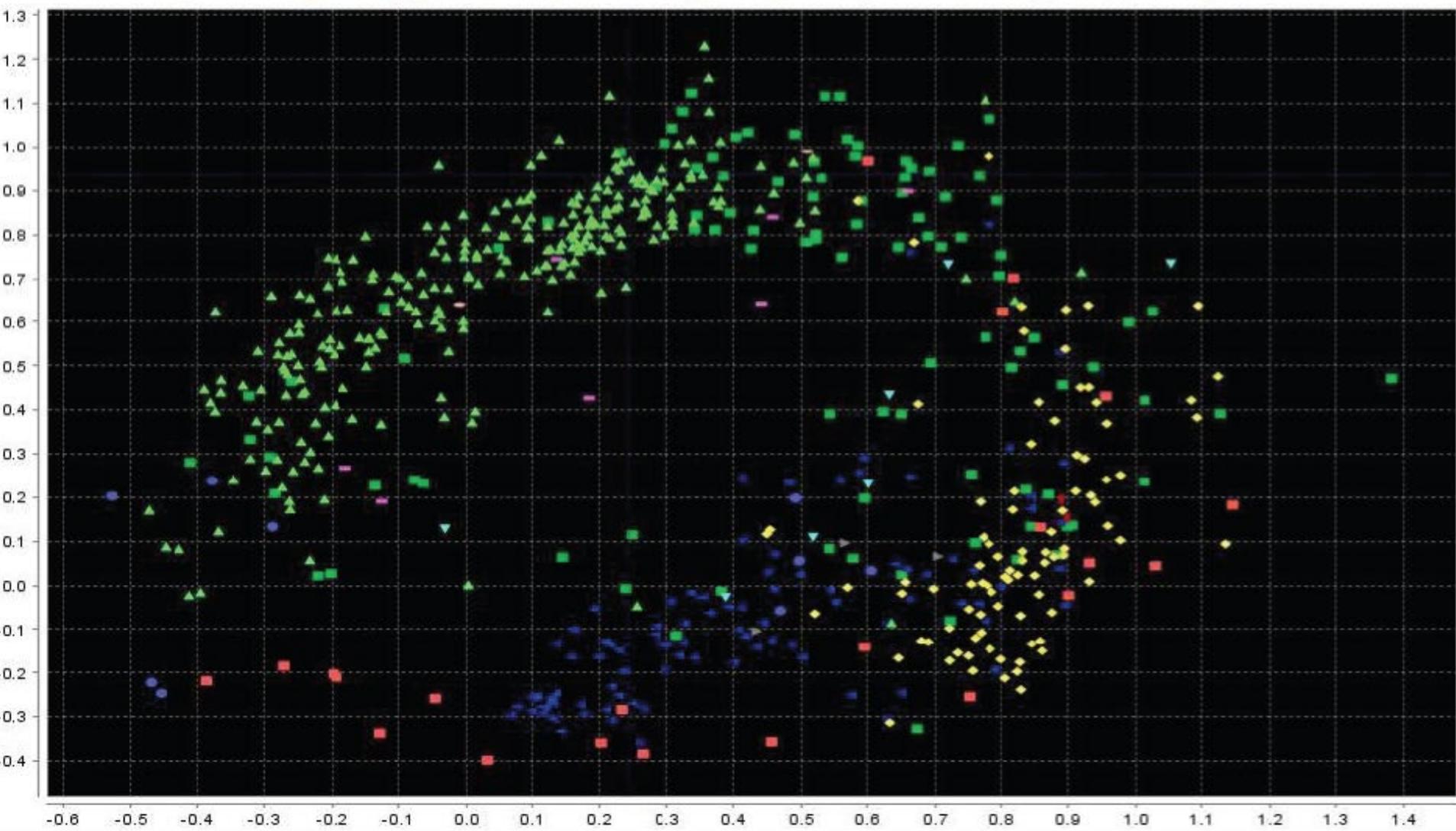
PCA for DNA Microarrays

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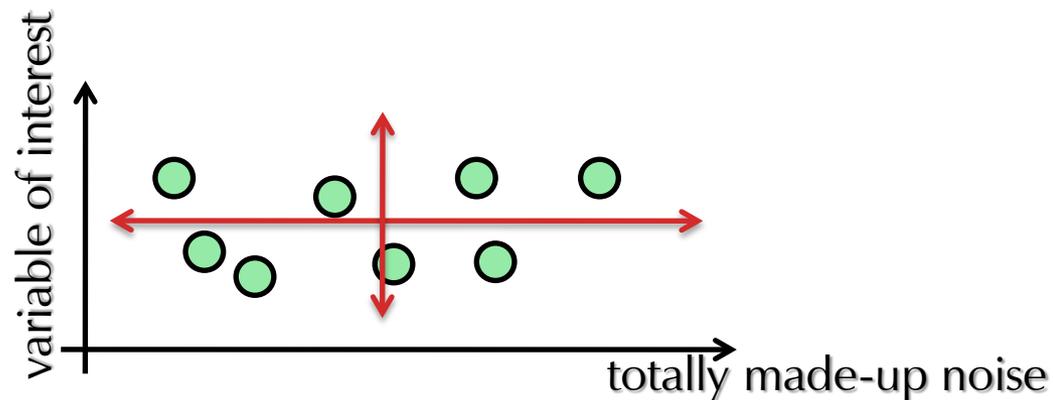
Music Map

■ ambient ● blues ▲ classical ◆ electronica ◆ folk ▲ jazz ◆ other ▶ pop ■ rap ◀ rock ■ world



Practical Considerations for PCA

- Sensitive to scale of each attribute (column)
 - In practice, may scale each attribute to have unit variance
- Sensitive to noisy attributes
 - Just because a dimension is highly weighted by PCA doesn't mean it's relevant, informative, etc.



Multidimensional Scaling

Multidimensional Scaling

- In some experiments, can only measure similarity or dissimilarity
 - e.g., is response to stimuli similar or different?
 - Frequent in psychophysical experiments, preference surveys, etc.
- Want to recover absolute positions in k -dimensional space

Multidimensional Scaling

- Example: given pairwise distances between cities

	Atl	Chi	Den	Hou	LA	Mia	NYC	SF	Sea	DC
Atlanta	0									
Chicago	587	0								
Denver	1212	920	0							
Houston	701	940	879	0						
LA	1936	1745	831	1374	0					
Miami	604	1188	1726	968	2339	0				
NYC	748	713	1631	1420	2451	1092	0			
SF	2139	1858	949	1645	347	2594	2571	0		
Seattle	2182	1737	1021	1891	959	2734	2406	678	0	
DC	543	597	1494	1220	2300	923	205	2442	2329	0

– Want to recover locations

Euclidean MDS

- Formally, let's say we have $n \times n$ matrix D consisting of squared distances $d_{ij} = (x_i - x_j)^2$
- Want to recover $n \times k$ matrix X of positions in k -dimensional space

$$D = \begin{pmatrix} 0 & (x_1 - x_2)^2 & (x_1 - x_3)^2 & \dots \\ (x_1 - x_2)^2 & 0 & (x_2 - x_3)^2 & \dots \\ (x_1 - x_3)^2 & (x_2 - x_3)^2 & 0 & \dots \\ \vdots & \vdots & \vdots & \ddots \end{pmatrix}$$

$$X = \begin{pmatrix} (\dots x_1 \dots) \\ (\dots x_2 \dots) \\ \vdots \end{pmatrix}$$

Euclidean MDS

- Observe that

$$d_{ij}^2 = (x_i - x_j)^2 = x_i^2 - 2x_i x_j + x_j^2$$

- Strategy: convert matrix D of d_{ij}^2 into matrix B of $x_i x_j$
 - “Centered” distance matrix
 - $B = XX^T$

Euclidean MDS

- Centering:

- Sum of row i of D = sum of column i of D =

$$\begin{aligned} s_i &= \sum_j d_{ij}^2 = \sum_j x_i^2 - 2x_i x_j + x_j^2 \\ &= nx_i^2 - 2x_i \sum_j x_j + \sum_j x_j^2 \end{aligned}$$

- Sum of all entries in D =

$$s = \sum_i s_i = 2n \sum_i x_i^2 - 2 \left(\sum_i x_i \right)^2$$

Euclidean MDS

- Choose $\sum x_i = 0$
 - Solution will have average position at origin

$$s_i = nx_i^2 + \sum_j x_j^2, \quad s = 2n \sum_j x_j^2$$

- Then,

$$d_{ij}^2 - \frac{1}{n}s_i - \frac{1}{n}s_j + \frac{1}{n^2}s = -2x_i x_j$$

- So, to get B :
 - compute row (or column) sums
 - compute sum of sums
 - apply above formula to each entry of D
 - Divide by -2

Factoring $B = XX^T$ using SVD

- Now have B , want to factor into XX^T
- If X is $n \times k$, B must have rank k
- Take SVD, set all but top k singular values to 0
 - Eliminate corresponding columns of U and V
 - Have $B' = U'W'V'^T$
 - B' is square and symmetric, so $U' = V'$
 - Take $X = U'$ times square root of W'

Multidimensional Scaling

- Result ($k = 2$):



Another application

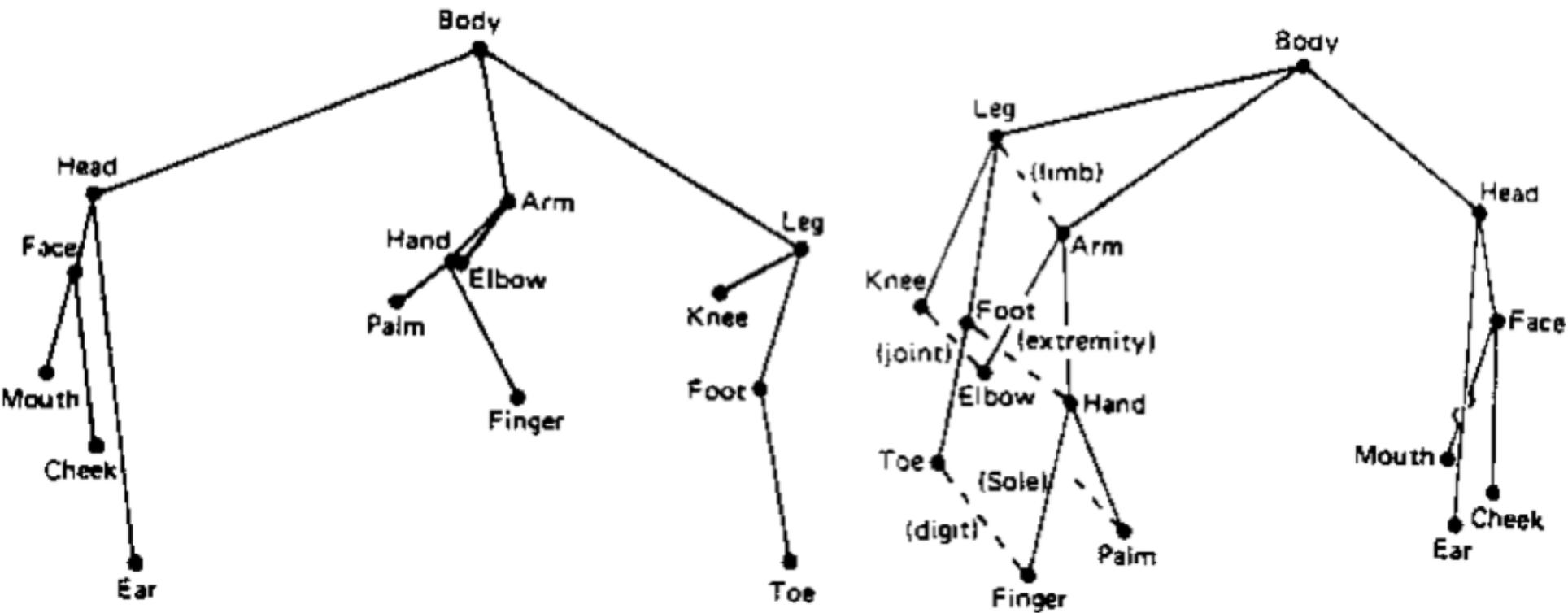
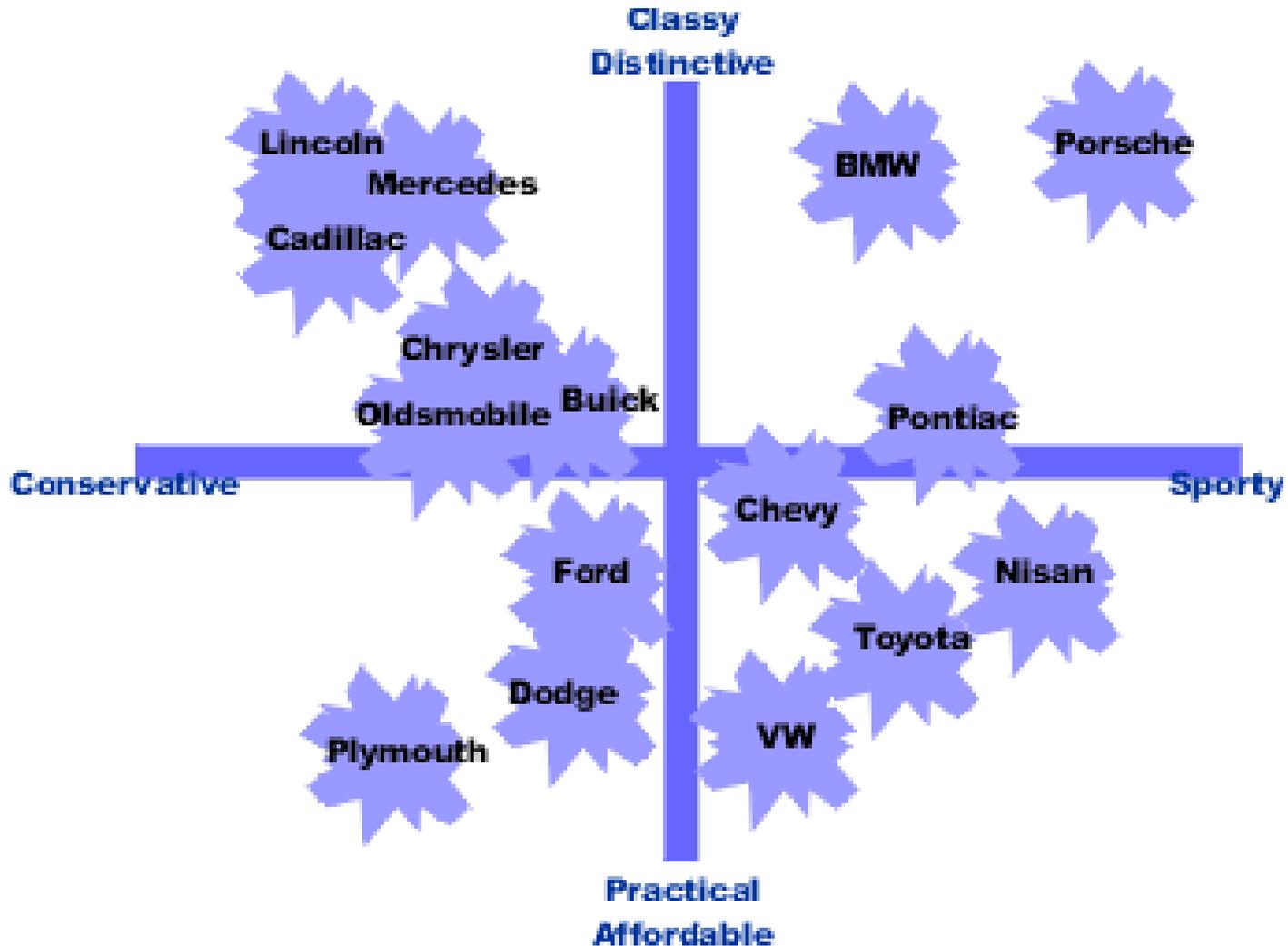


Figure 2 (a) RMDs of children's similarity judgments about 15 body parts: (b) RMDs of adults' similarity judgments about 15 body parts.

Perceptual Mapping for Marketing



Multidimensional Scaling

- Caveat: actual axes, center not necessarily what you want (can't recover them!)
- This is “classical” or “Euclidean” MDS [Torgerson 52]
 - Distance matrix assumed to be actual Euclidean distance
- More sophisticated versions available
 - “Non-metric MDS”: not Euclidean distance, sometimes just *inequalities*
 - Replicated MDS: for multiple data sources (e.g. people)
 - “Weighted MDS”: account for observer bias