

Distance and similarity (I)

Hongxin Zhang
zhx@cad.zju.edu.cn

State Key Lab of CAD&CG, ZJU
2015-03-24





In the last lesson

- Data driven decomposition:
 - Data driven curve fitting
 - Singular value decomposition (the power of orthogonal basis)
 - A sort of spectral analysis
- PCA and its related techniques are very useful



In the last lesson

- Mathematical concepts and techniques
 - Least squares (LSQ)
 - Curve fitting
 - Norm (范数) and inner product (内积)
 - Singular value decomposition
 - Eigen vectors and eigen-values
 - Low rank matrix approximation and decomposition



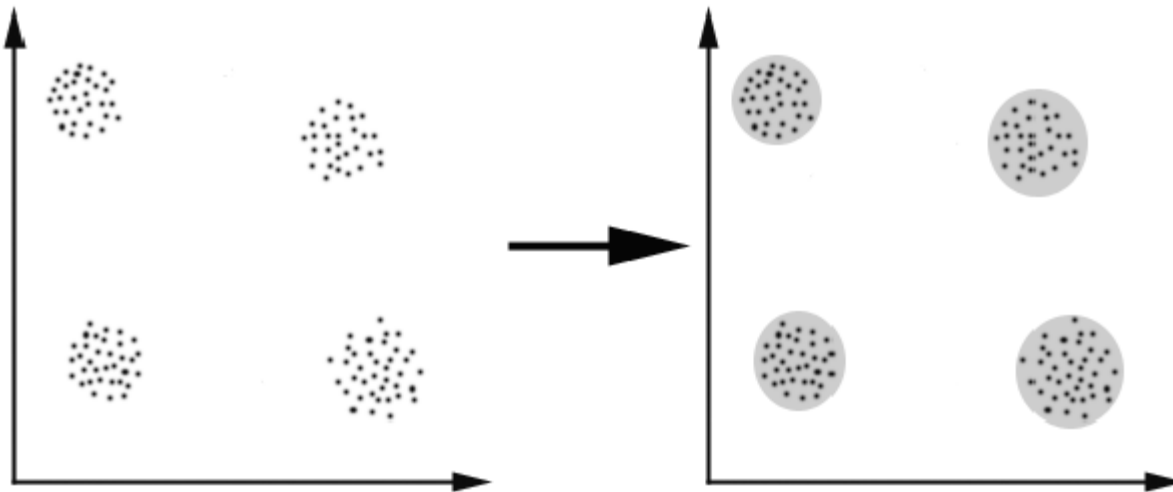
Today's Talk

- Pre-processing: Distance / metric learning
 - ISO-map
 - LLE
- What is similarity? How to Clustering
 - Spectral based
 - E-M based
 - MOG and K-means
 - Mean shift

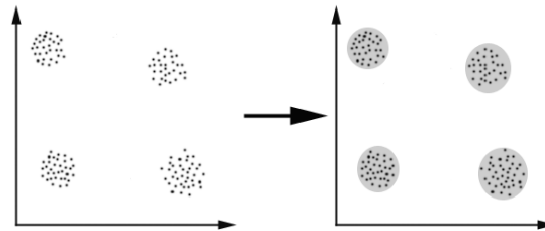


Clustering

- Given set of data points, group them, find the overall structure
- **Unsupervised** learning
- Learn the similarity. Which patient are similar? (or customers, faces, earthquakes, ...)



Distance



- Given n -dimensional vector \mathbf{x} , \mathbf{y}
- Euclidian (L^2 distance)

$$\text{dist}(\mathbf{x}, \mathbf{y}; 2) = \left(\sum_{i=1}^n (x_i - y_i)^2 \right)^{1/2}$$

- L^1 distance

$$\text{dist}(\mathbf{x}, \mathbf{y}; 1) = \sum_{i=1}^n |x_i - y_i|$$

- L^p distance (Minkowsky)

$$\text{dist}(\mathbf{x}, \mathbf{y}; p) = \left(\sum_{i=1}^n (x_i - y_i)^p \right)^{1/p}$$

$$\text{dist}(\mathbf{x}, \mathbf{y}; \infty) = \max_i |x_i - y_i|$$

切比雪夫

Distance, Norm and inner product



- Distance

$$dist(\mathbf{x}, \mathbf{y}; 2) = \left(\sum_{i=1}^n (x_i - y_i)^2 \right)^{1/2}$$

- Norm

$$norm(\mathbf{x}; 2) := \|\mathbf{x}\|_2 = \left(\sum_{i=1}^n x_i^2 \right)^{1/2}$$

- Inner product

$$\mathbf{x} \cdot \mathbf{y} = \sum_{i=1}^n x_i y_i$$

$$norm(\mathbf{x} - \mathbf{y}; 2) = dist(\mathbf{x}, \mathbf{y}; 2) \quad dist(x, y; 2) = (\mathbf{x} \cdot \mathbf{x} + \mathbf{y} \cdot \mathbf{y} - 2\mathbf{x} \cdot \mathbf{y})^{1/2}$$
$$= \left(\|\mathbf{x}\|_2^2 + \|\mathbf{y}\|_2^2 - 2\mathbf{x} \cdot \mathbf{y} \right)^{1/2}$$



Dimensional aware distances

- Along dimension j :

$$\bar{x}_j = \frac{1}{N} \sum_{i=1}^N x_{i,j}$$

$$R_j = \max_i x_{i,j} - \min_i x_{i,j}$$

$$S_j = \left(\frac{1}{N-1} \sum_{i=1}^N (x_{i,j} - \bar{x}_j)^2 \right)^{1/2}$$

- Normalized data:

$$x'_{i,j} = \frac{x_{i,j} - \bar{x}_j}{R_j}$$

$$x'_{i,j} = \frac{x_{i,j} - \bar{x}_j}{S_j}$$



M-distance

- Consider the dependency of different dimensions

$$\text{dist}(\mathbf{x}, \mathbf{y}; \mathbf{M}) = (\mathbf{x} - \mathbf{y})^T \mathbf{M}^{-1} (\mathbf{x} - \mathbf{y})$$

- M is the covariance matrix of data
- Transform invariance



Data whitening

设有均值为零的随机信号向量 x ，其自相关矩阵为

$$R_x = E[xx^T] \neq I$$

很明显， R_x 是对称矩阵，且是非负定的（所有特征值都大于或等于0）。

现在，寻找一个线性变换 B 对 x 进行变换，即 $y = Bx$ ，使得

$$R_y = BE[xx^T]B^T = I$$

上式的含义是： y 的各分量是不相关的，即 $E[y_i y_j] = \delta_{ij}$ 。通常将这个过程称为“空间解相关”、“空间白化”或“球化”。 B 称为空间解相关矩阵（空间白化矩阵、球化矩阵）。

由 R_x 的性质可知，其存在特征值分解：

$$R_x = Q\Sigma Q^T$$

Q 是正交矩阵， Σ 是对角矩阵，其对角元素是 R_x 特征值。

令

$$B = \Sigma^{-1/2} Q^T \tag{1}$$

$$R_y = BE[xx^T]B^T = I$$

D 上式的含义是： y 的各分量是不相关的，即 $E[y_i y_j] = \delta_{ij}$ 。通常将这个过程称为“空间解相关”、“空间白化”或“球化”。 B 称为空间解相关矩阵（空间白化矩阵、球化矩阵）。

由 R_x 的性质可知，其存在特征值分解：

$$R_x = Q\Sigma Q^T$$

Q 是正交矩阵， Σ 是对角矩阵，其对角元素是 R_x 特征值。

令

$$B = \Sigma^{-1/2} Q^T \quad (1)$$

则有

$$R_y = (\Sigma^{-1/2} Q^T) Q \Sigma Q^T (\Sigma^{-1/2} Q^T)^T = I$$

因此，通过矩阵 B 线性变换后， y 的各个分量变得不相关了。

对于 R_x 来说，特征值分解和奇异值分解是等价的，而奇异值分解的数值算法比特征值分解的数值算法具有更好的稳定性，因此一般都用奇异值分解来构造空间解相关矩阵 B 。

应该注意到，“空间解相关”不能保证各分量信号之间的“独立性”，但它能够简化盲分离算法或改善分离算法的性能。

More complex method for distance computing



- PCA ...
- structure aware
 - Main idea:
 - Find a suitable mapping
 - Compute distance in mapped space
 - Available techniques
 - MDS + global geodesic distance: ISO-MAP
 - Local distance approximation: LLE
 - ...

Classical Multi-dimensional Scaling



- MDS: 多维标度法
- Main idea:
 - Compute (match) distance between samples
 - Use SVD to find similarity

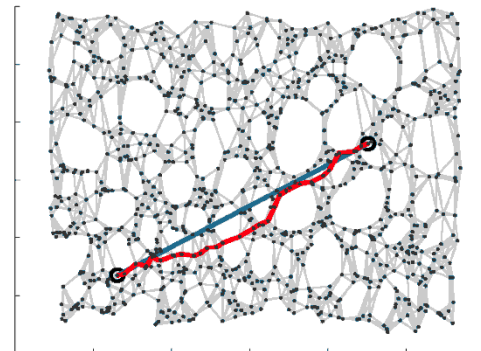
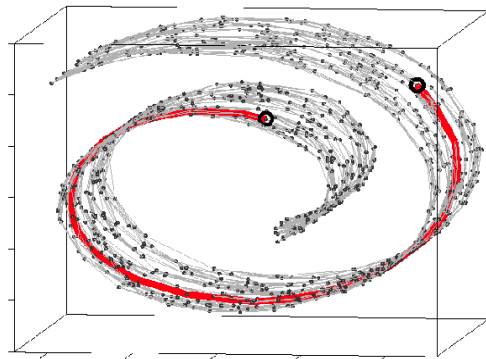
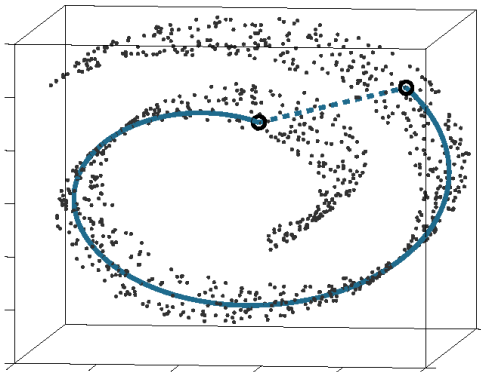
Isomap: (Science 2001)

Isometric feature mapping



- Preserve the **intrinsic geometry** of the data.
- Use the **geodesic distances** on manifold between all pairs.

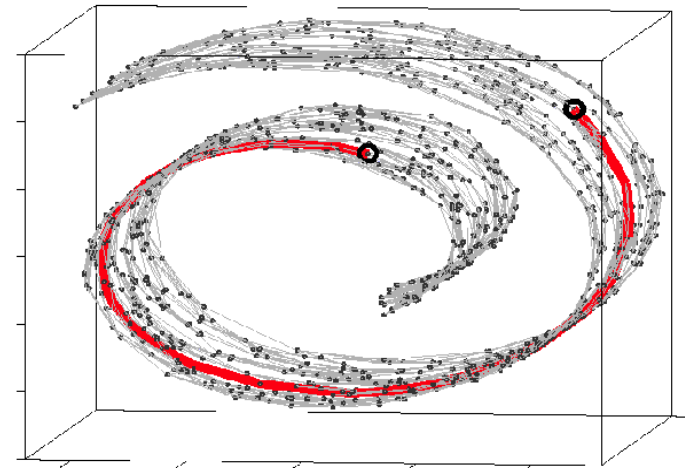
Three steps algorithm



Isomap: Construct Neighborhood Graph



- Determine which points are neighbors, based on the distances $d(i,j)$.
 - K nearest neighbors
 - ϵ -radius



- Create a graph G , with edges between neighbors and distance weights.

Isomap: Compute Shortest Paths



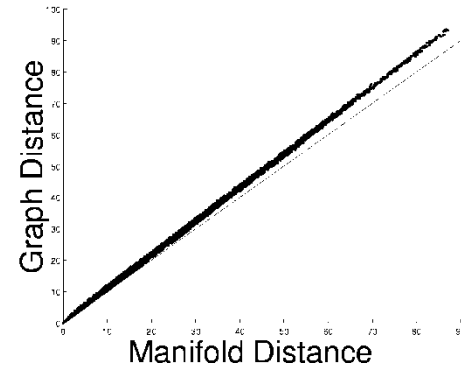
- Estimate the geodesic distances.
- Compute all-pairs shortest paths in G .
- Can be done using Floyd's algorithm, $O(N^2 \ln N)$.

$$d_G(i, j) = d(i, j) \text{ neighborin g } i, j$$

$$d_G(i, j) = \infty \quad \text{otherwise}$$

for $k = 1, 2, \dots, N$

$$d_G(i, j) = \min\{ d_G(i, j), d_G(i, k) + d_G(k, j) \}$$



Isomap: Construct d-dimensional Embedding



Classical **MDS** with $d_G(i,j)$,
minimize the cost function:

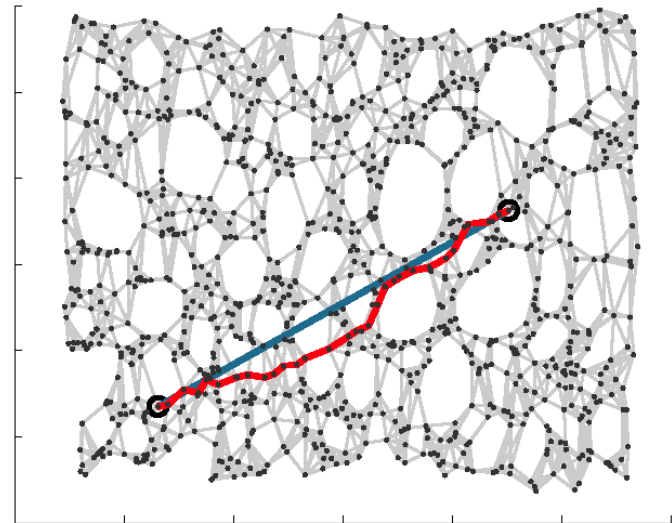
$$E = \left\| \tau(D_G) - \tau(D_Y) \right\|_{L^2}$$

$$\text{where } D_Y(i, j) = \|y_i - y_j\|$$

$$D_G(i, j) = d_G(i, j)$$

and

$$\tau(D) = \frac{-1}{2} \left(I - \frac{1}{N} \right) D^2 \left(I - \frac{1}{N} \right)$$



Solution: take top d
eigenvectors of the
matrix $\tau(D_G)$

Isomap: Classical Multi-dimensional Scaling



$$\mathbf{X}'\mathbf{X} = -\frac{1}{2}\mathbf{J}\mathbf{E}\mathbf{J} \quad \text{E: Euclidian distance matrix}$$

$$\mathbf{B} = -\frac{1}{2}\mathbf{J}\mathbf{M}\mathbf{J} \quad \text{M: Manifold distance matrix}$$

$$L(\hat{\mathbf{X}}) = \left\| -\frac{1}{2}\mathbf{J}(\mathbf{E} - \mathbf{M})\mathbf{J} \right\|$$

$$= \left\| \hat{\mathbf{X}}\hat{\mathbf{X}}' - \mathbf{B} \right\|.$$

$$\mathbf{B} = \mathbf{Q}\mathbf{\Lambda}\mathbf{Q}' \quad \hat{\mathbf{X}} = \mathbf{Q}_+ \mathbf{\Lambda}_+^{\frac{1}{2}}$$

$$c_i = \sum_{a=1}^m x_{ia}^2$$

$$d_{ij}^2 = \sum_{a=1}^m (x_{ia} - x_{ja})^2$$

$$\mathbf{E} = \mathbf{c}\mathbf{1}' + \mathbf{1}\mathbf{c}' - 2\mathbf{X}\mathbf{X}'$$

$$\mathbf{J} = \mathbf{I} - \frac{1}{n}\mathbf{1}\mathbf{1}'$$

$$\mathbf{B} = -\frac{1}{2}\mathbf{J}(\mathbf{c}\mathbf{1}' + \mathbf{1}\mathbf{c}' - 2\mathbf{X}\mathbf{X}')\mathbf{J}$$

$$= -\frac{1}{2}\mathbf{J}\mathbf{c}\mathbf{0}' - \frac{1}{2}\mathbf{0}\mathbf{c}'\mathbf{J} + \mathbf{J}\mathbf{X}\mathbf{X}'\mathbf{J}$$

$$= \mathbf{X}\mathbf{X}'.$$

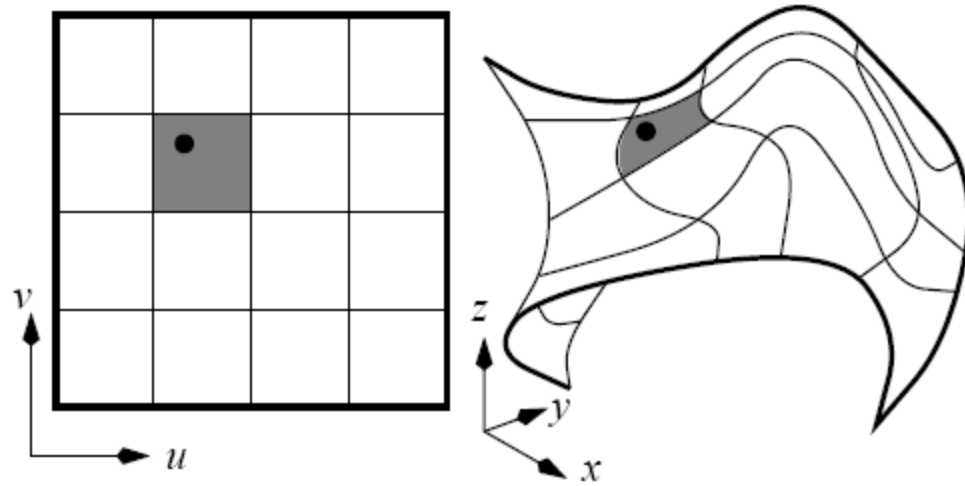
Eigen-structure analysis again

Isomap: Classical Multi-dimensional Scaling (2D)



```
J = eye(n) - ones(n)./n;  
B = -0.5 * J * M * J;  
      % Find largest eigenvalues+their eigenvectors:  
[Q, L] = eigs(B, 2, 'LM');  
      % Extract the coordinates:  
newy = sqrt(L(1, 1)). * Q(:, 1);  
newx = sqrt(L(2, 2)). * Q(:, 2);
```

Isomap: application texture mapping



(a)



(b)



(a)



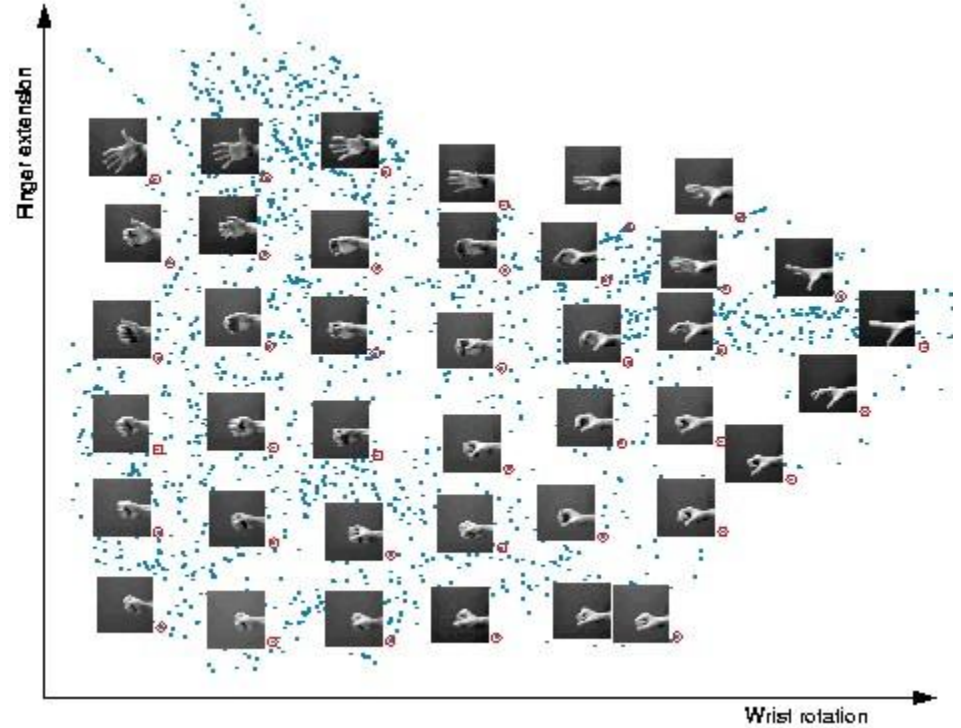
(b)

Fig. 3. An example of a face flattening. (a) A 3D reconstruction of a face. (b) The flattened texture image of the face.

Isomap: Examples



- $N=2000$ images
64x64 pixels $K=6$



Isomap: More Results

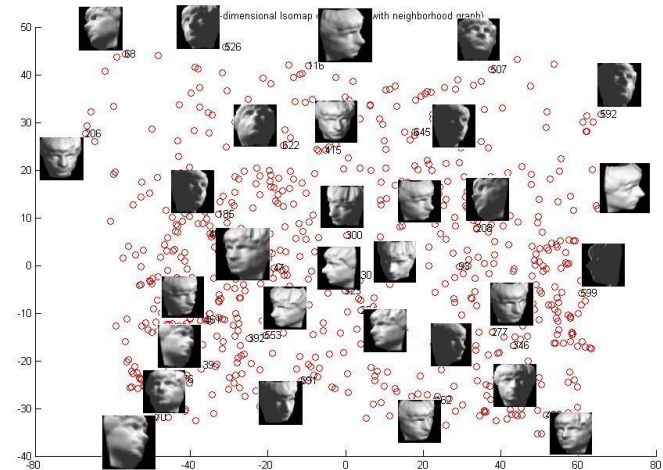
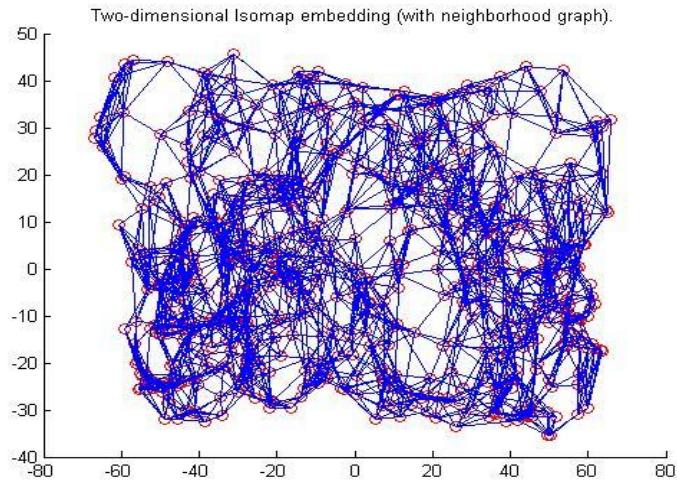


Input: 698
images of 64x64

$K=7, d=2$



Outputs:

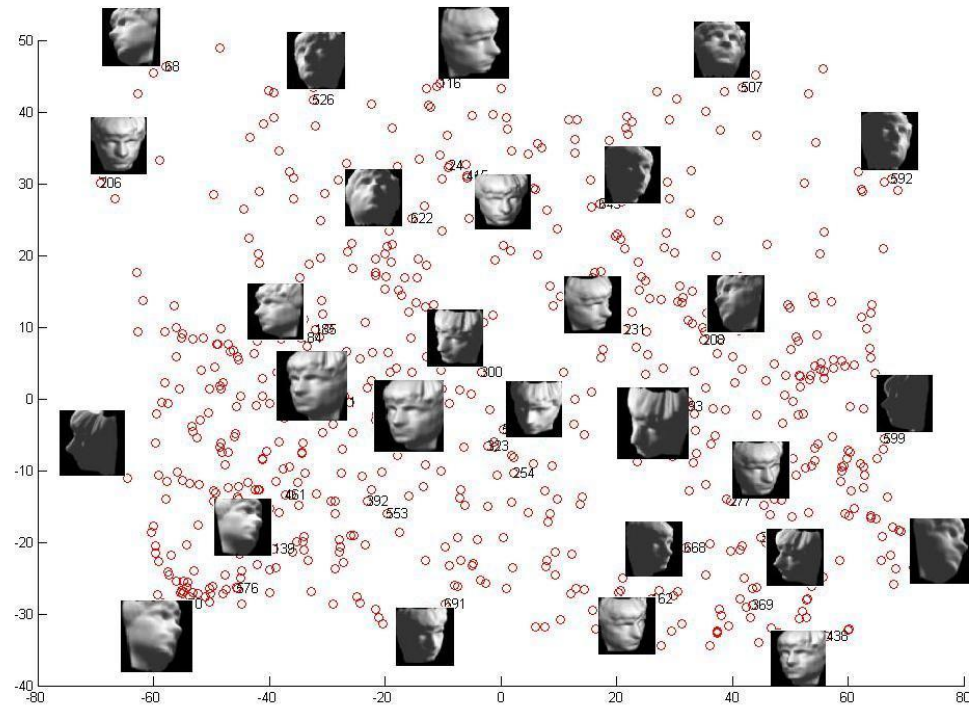


Isomap: More Results



- Same inputs, but this time with $d=3$

698 images of 64×64 $K=7$



Locally Linear Embedding (LLE)

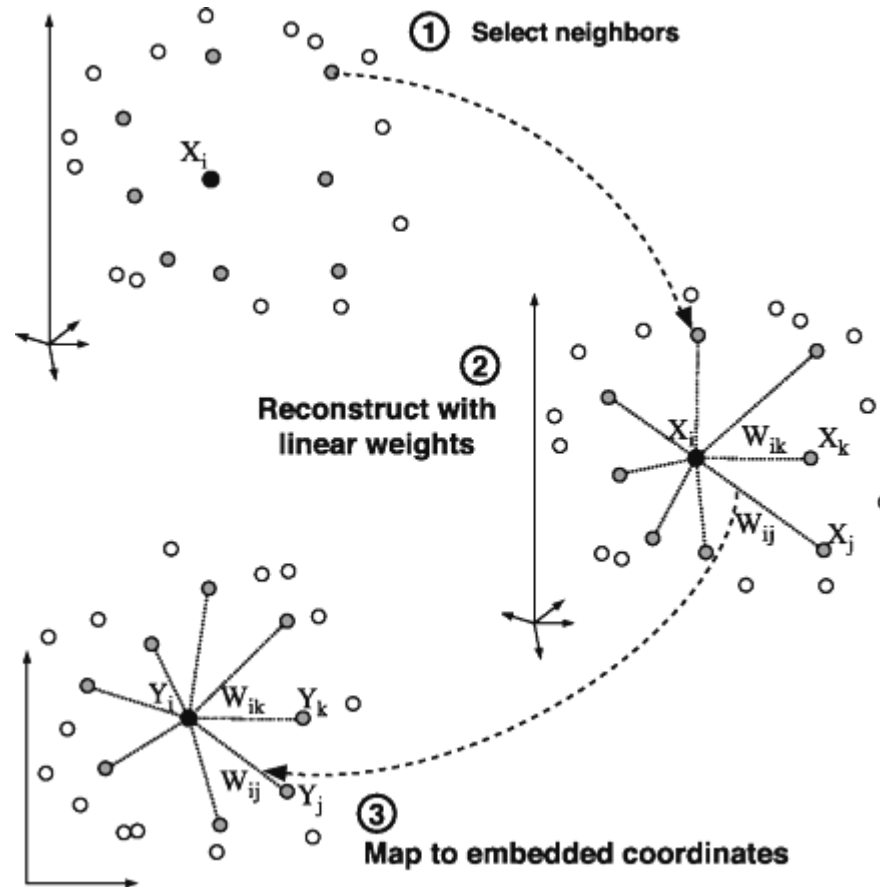


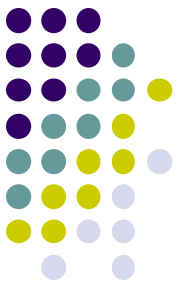
- Recovers global nonlinear structure from locally linear fits.
- Each data point and it's neighbors is expected to lie on or close to a locally linear patch.
- Each data point is constructed by it's neighbors:

$$\vec{\hat{X}}_i = \sum_j w_{ij} \vec{X}_j$$

$$w_{ij} = 0 \text{ if } \vec{X}_j \text{ is not a neighbor of } \vec{X}_i$$

LLE – main idea





LLE: Getting the Reconstruction Weights

- We want to minimize the error function:

$$\mathcal{E}(W) = \sum_i \left| \vec{X}_i - \sum_j W_{ij} \vec{X}_j \right|^2$$

- With the constraints:

$$W_{ij} = 0 \quad \text{if } \vec{X}_j \text{ is not a neighbor of } \vec{X}_i$$

$$\sum_j W_{ij} = 1$$

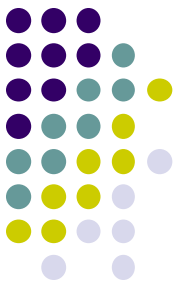
- Solution (using Lagrange multipliers):

$$W_j = \sum_k C_{jk}^{-1} (\vec{X} \vec{\eta}_k + \lambda)$$

$$\lambda = 1 - \frac{\sum_{jk} C_{jk}^{-1} (\vec{X} \vec{\eta}_k)}{\sum_{jk} C_{jk}^{-1}}$$

LLE:

Find Embedded Coordinates



- Choose d-dimensional coordinates, Y , to minimize:

$$\phi(Y) = \sum_i \left| \vec{Y}_i - \sum_j W_{ij} \vec{Y}_j \right|^2$$

$$\text{Under: } \sum_i \vec{Y}_i = \vec{0}, \quad \frac{1}{N} \sum_i \vec{Y}_i \vec{Y}_i^T = I$$

Quadratic form:
$$\phi(Y) = \sum_{ij} M_{ij} (\vec{Y}_i \vec{Y}_j)$$

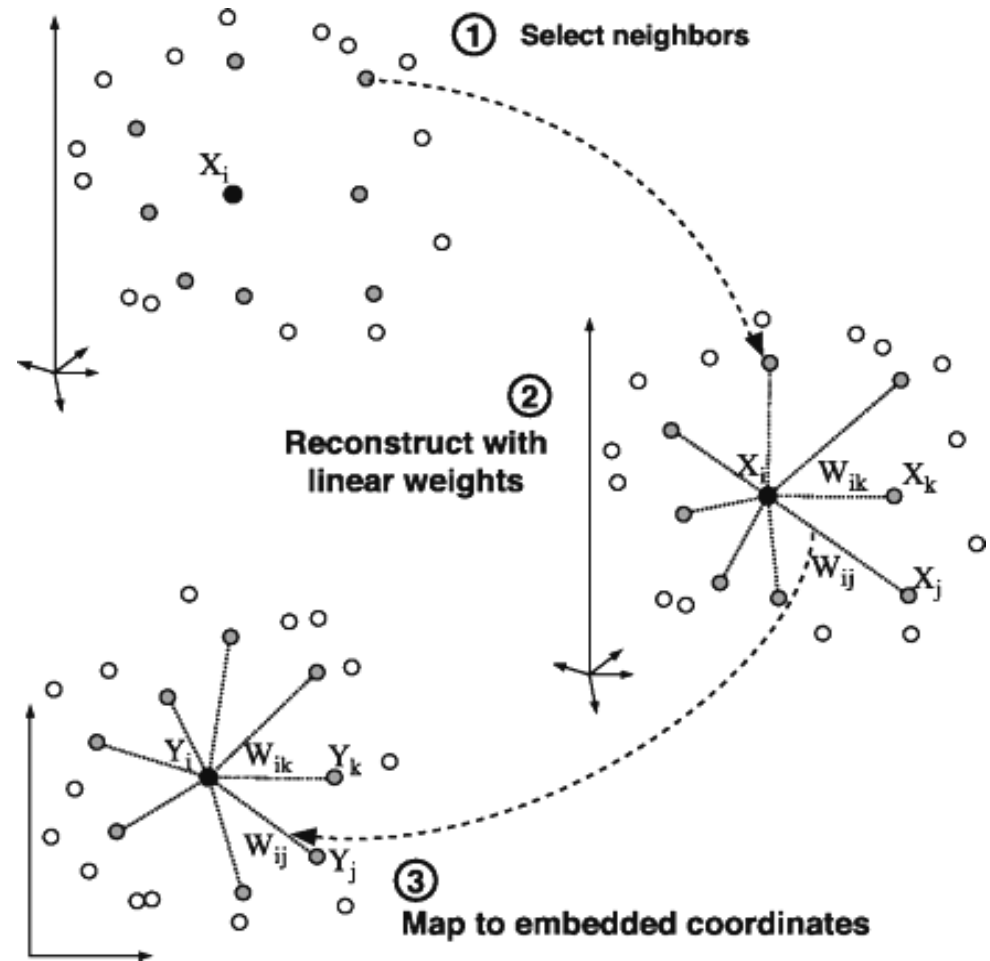
where:
$$M = (I - W)^T (I - W)$$

- Solution: compute bottom d+1 eigenvectors of M . (discard the last one)

LLE: Summary



- Input: N data items in D dimension (X).
- Output: $d < D$ dimensional embedding coordinates (Y) for the input points.



LLE: Algorithm Pseudocode (I)



Find neighbors in X space

For $i=1:N$

 compute the distance from X_i to every other point X_j

 find the K smallest distances

 assign the corresponding points to be neighbors of X_i

end

<http://www.cs.toronto.edu/~roweis/lle/algorithm.html>

LLE:

Algorithm Pseudocode (II)



Solve for reconstruction weights W .

for $i=1:N$

 create matrix Z consisting of all neighbors of X_i

 subtract X_i from every column of Z

 compute the local covariance $C=Z'^*Z$

 solve linear system $C*w = 1$ for w

 set $W_{ij}=0$ if j is not a neighbor of i

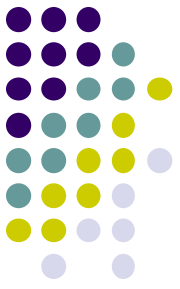
 set the remaining elements in the i th row of W equal to

$w/\text{sum}(w)$;

end

LLE:

Algorithm Pseudocode (III)



Compute embedding coordinates Y using weights W .

create sparse matrix $M = (I-W)'*(I-W)$

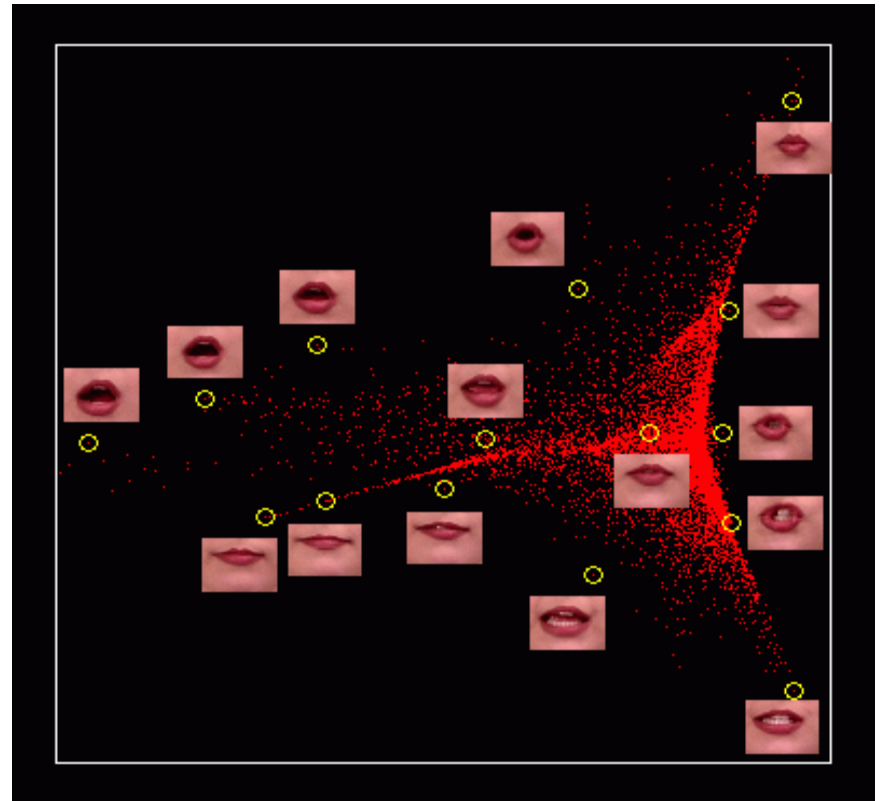
find bottom $d+1$ eigenvectors of M (corresponding to the $d+1$ smallest eigenvalues)

set the q -th ROW of Y to be the $q+1$ smallest eigenvector (discard the bottom eigenvector $[1,1,1,1\dots]$ with eigenvalue zero)

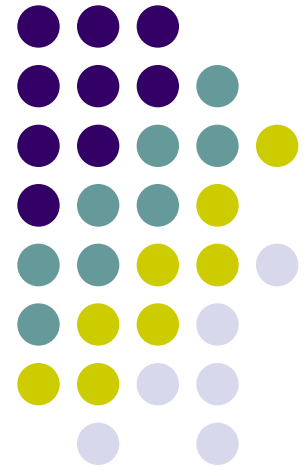
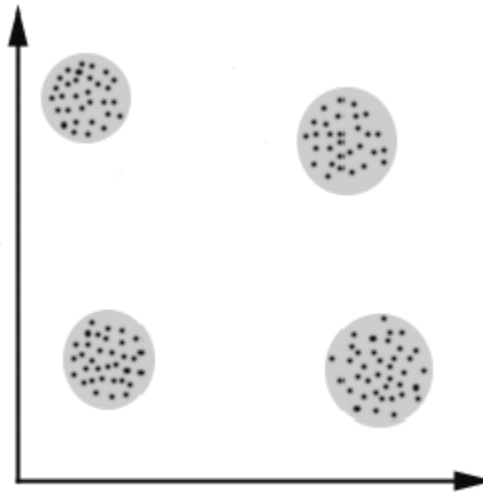
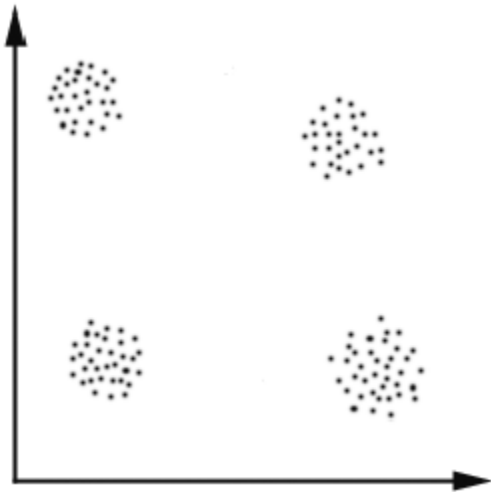
LLE: Example



- $N=8588$ (RGB) images of lips of size 108×84 .
 $D=27216$
- Num of neighbors $K=16$



Clustering



Spectral clustering



- The spectral clustering method we define relies on a random walk representation over the points. We construct this in three steps

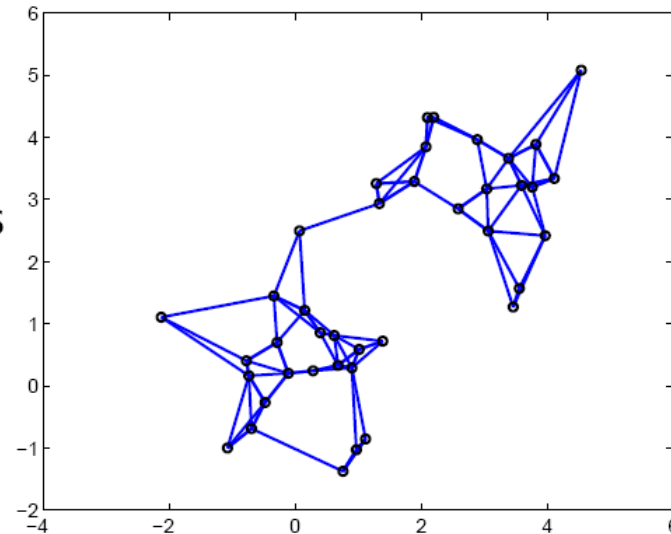
1. a nearest neighbor graph
2. similarity weights on the edges:

$$W_{ij} = \exp\{-\beta\|\mathbf{x}_i - \mathbf{x}_j\|\}$$

where $W_{ii} = 1$ and the weight is zero for non-edges.

3. transition probability matrix

$$P_{ij} = W_{ij} / \sum_{j'} W_{ij'}$$



Properties of the random walk



- If we start from i_0 , the distribution of points i_t that we end up in after t steps is given by

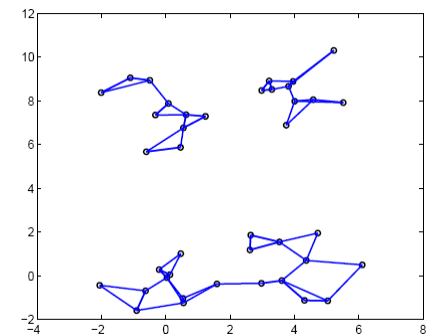
$$i_1 \sim P_{i_0 i_1}, \quad P_{ij} = \frac{W_{ij}}{W_{i\cdot}}, \quad \text{where } W_{i\cdot} = \sum_j W_{ij}$$

$$i_2 \sim \sum_{i_1} P_{i_0, i_1} P_{i_1 i_2} = [P^2]_{i_0 i_2},$$

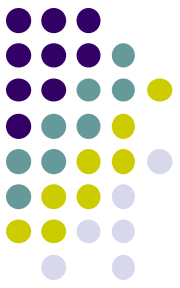
$$i_3 \sim \sum_{i_1} \sum_{i_2} P_{i_0, i_1} P_{i_1 i_2} P_{i_2 i_3} = [P^3]_{i_0 i_3},$$

...

$$i_t \sim [P^t]_{i_0 i_t}$$



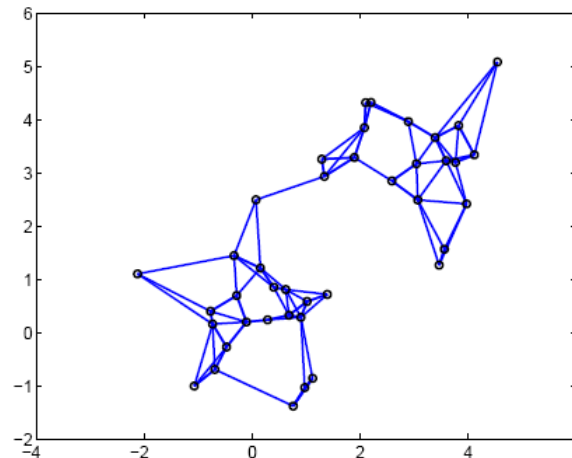
where $P^t = PP \dots P$ (t matrix products) and $[\cdot]_{ij}$ denotes the i, j component of the matrix.



Random walk and clustering

- The distributions of points we end up in after t steps converge as t increases. If the graph is connected, the resulting distribution is independent of the starting point

Even for large t , the transition probabilities $[P^t]_{ij}$ have a slightly higher probability of transitioning within “clusters” than across; we want to recover this effect from eigenvalues/vectors



Eigenvalues/vectors and spectral clustering



- Let W be the matrix with components W_{ij} and D a diagonal matrix such that $D_{ii} = \sum_j W_{ij}$. Then

$$P = D^{-1}W$$

- To find out how P^t behaves for large t it is useful to examine the eigen-decomposition of the following symmetric matrix

$$D^{-\frac{1}{2}}WD^{-\frac{1}{2}} = \lambda_1\mathbf{z}_1\mathbf{z}_1^T + \lambda_2\mathbf{z}_2\mathbf{z}_2^T + \dots + \lambda_n\mathbf{z}_n\mathbf{z}_n^T$$

where the ordering is such that $|\lambda_1| \geq |\lambda_2| \geq \dots \geq |\lambda_n|$.



Eigenvalues/vectors cont'd

- The symmetric matrix is related to P^t since

$$(D^{-\frac{1}{2}}WD^{-\frac{1}{2}}) \dots (D^{-\frac{1}{2}}WD^{-\frac{1}{2}}) = D^{\frac{1}{2}}(P \dots P)D^{-\frac{1}{2}}$$

This allows us to write the t step transition probability matrix in terms of the eigenvalues/vectors of the symmetric matrix

$$\begin{aligned} P^t &= D^{-\frac{1}{2}} \left(D^{-\frac{1}{2}}WD^{-\frac{1}{2}} \right)^t D^{\frac{1}{2}} \\ &= D^{-\frac{1}{2}} \left(\lambda_1^t \mathbf{z}_1 \mathbf{z}_1^T + \lambda_2^t \mathbf{z}_2 \mathbf{z}_2^T + \dots + \lambda_n^t \mathbf{z}_n \mathbf{z}_n^T \right) D^{\frac{1}{2}} \end{aligned}$$

where $\lambda_1 = 1$ and

$$P^\infty = D^{-\frac{1}{2}} \left(\mathbf{z}_1 \mathbf{z}_1^T \right) D^{\frac{1}{2}}$$

Eigenvalues/vectors and spectral clustering



- We are interested in the largest correction to the asymptotic limit

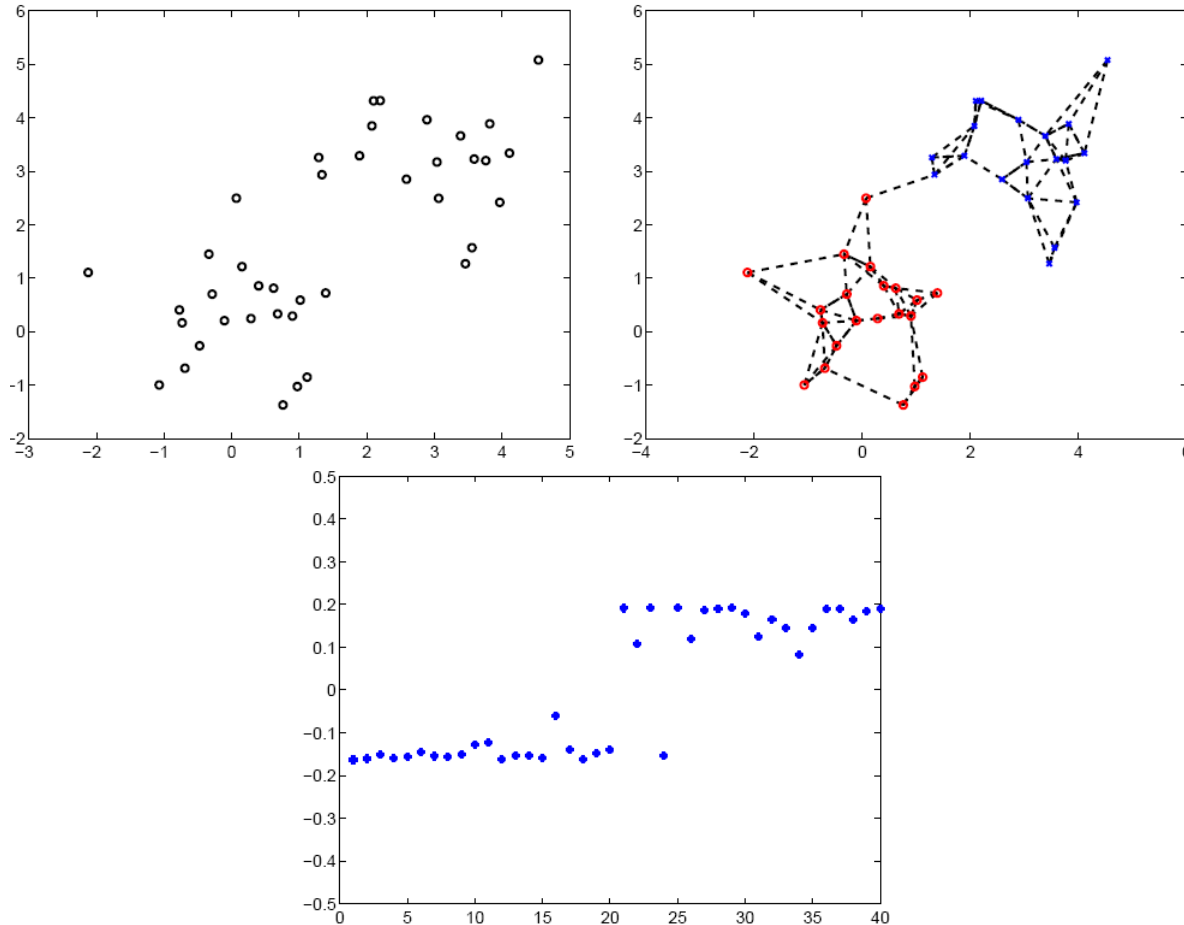
$$P^t \approx P^\infty + D^{-\frac{1}{2}} \left(\lambda_2^t \mathbf{z}_2 \mathbf{z}_2^T \right) D^{\frac{1}{2}}$$

Note: $[\mathbf{z}_2 \mathbf{z}_2^T]_{ij} = z_{2i} z_{2j}$ and thus the largest correction term increases the probability of transitions between points that share the same sign of z_{2i} and decreases transitions across points with different signs

- Binary spectral clustering: we divide the points into clusters based on the sign of the elements of \mathbf{z}_2

$$z_{2j} > 0 \Rightarrow \text{cluster 1, otherwise cluster 0}$$

Spectral clustering: example



Components of the eigenvector corresponding to the second largest eigenvalue



Reference papers of SC

- A. Y. Ng, M. I. Jordan, and Y. Weiss, *On spectral clustering: Analysis and an algorithm*, NIPS, (2001)
- Y. Weiss, *Segmentation using eigenvectors: a unifying view*. ICCV, (1999)
- J. Shi and J. Malik, *Normalized cuts and image segmentation*, IEEE TPAMI, 22 (2000)
- And more about image segmentations ...
 - Graph cut
 - Mean-shift



Classical methods on cluster distance



- A *linkage* method: we have to be able to measure distances between clusters of examples C_k and C_l

a) Single linkage:

Nearest neighbor

$$d_{kl} = \min_{i \in C_k, j \in C_l} d(\mathbf{x}_i, \mathbf{x}_j)$$

b) Average linkage:

$$d_{kl} = \frac{1}{|C_l| |C_k|} \sum_{i \in C_k, j \in C_l} d(\mathbf{x}_i, \mathbf{x}_j)$$

c) Centroid linkage:

$$d_{kl} = d(\bar{\mathbf{x}}_k, \bar{\mathbf{x}}_l), \quad \bar{\mathbf{x}}_l = \frac{1}{|C_l|} \sum_{i \in C_l} \mathbf{x}_i$$

Hierarchical (bottom-up) clustering

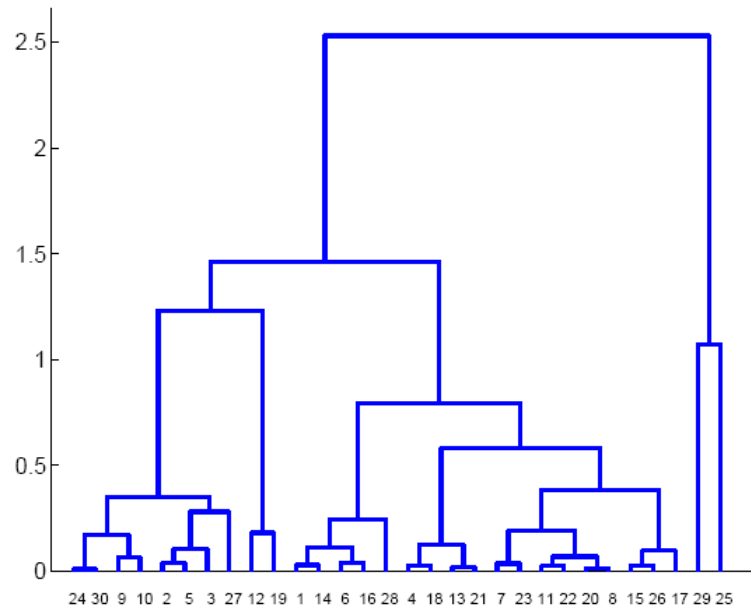


- Hierarchical agglomerative clustering: we sequentially merge the pair of “closest” points/clusters
- The procedure
 1. Find two closest points (clusters) and merge them
 2. Proceed until we have a single cluster (all the points)
- Two prerequisites:
 1. distance measure $d(x_i, x_j)$ between two points
 2. distance measure between clusters (cluster linkage)

Hierarchical (bottom-up) clustering



- A dendrogram representation of hierarchical clustering



The height of each pair represents the distance between the merged clusters; the specific linear ordering of points is chosen for clarity



Clustering vs. Classification

- **Clustering**

- **Instance:** $\{\mathbf{x}_i\}_{i=1}^N$
- **Learn:** $\langle \mathbf{x}_i, t_i \rangle$ and/or mapping from \mathbf{x} to $t(\mathbf{x})$

- **Classification/Regression**

- **Instance:** $\langle \mathbf{x}_i, t_i \rangle$
- **Learn:** mapping from \mathbf{x} to $t(\mathbf{x})$

The End

新浪微博： @浙大张宏鑫

微信公众号：

