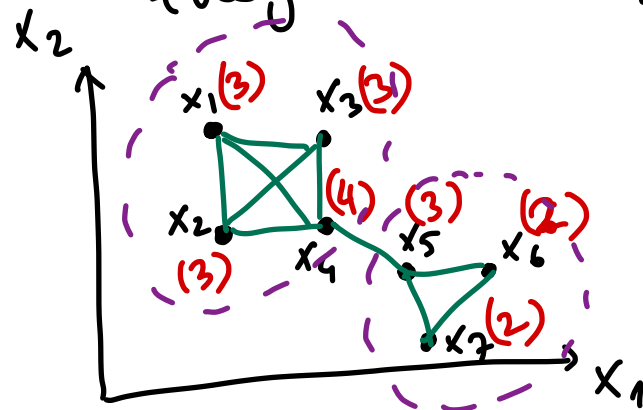


SPECTRAL CLUSTERING

- define local neighborhoods

- if the distance between x_i & x_j is smaller than a threshold they are neighbors.



$$b_{ij} = \begin{cases} 1 & \text{if } \|x_i - x_j\|_2 < \underline{\delta} \\ 0 & \text{otherwise} \end{cases}$$

$$b_{ij} = \begin{cases} \exp\left[-\frac{\|x_i - x_j\|_2^2}{2\sigma^2}\right] & \text{if } \|x_i - x_j\|_2 < \underline{\delta} \\ 0 & \text{otherwise} \end{cases}$$

$$b_{ii} = 0 \quad \forall i$$

$$d_{ii} = \sum_{j \neq i} b_{ij} \quad \forall i$$

$B =$

	x_1	x_2	x_3	x_4	x_5	x_6	x_7
x_1	0	1	1	1	0	0	0
x_2	1	0	1	1	0	0	0
x_3	1	1	0	1	0	0	0
x_4	1	1	1	0	0	0	0
x_5	0	0	0	0	1	1	1
x_6	0	0	0	0	1	0	1
x_7	0	0	0	0	1	1	0

connectivity or adjacency matrix

$D =$

	x_1	x_2	x_3	x_4	x_5	x_6	x_7
x_1	3						
x_2		3					
x_3			3				
x_4				4			
x_5					3		
x_6						2	
x_7							2

of neighbors of data point i

$$d_{ij} = 0 \quad \forall (i, j \neq i)$$

mass \rightarrow outgoing \rightarrow

$\rightarrow [3 \ -1 \ -1 \ -1 \ 0 \ 0 \ 0]$

Laplacian Matrix:

$$L_{N \times N} = D_{N \times N} - B_{N \times N}$$

↳ each row (column) sums up to 0.

$$L_{\text{RANDOM-WALK}} = \bar{D}^{-1} \cdot L = \bar{D}^{-1} \cdot (D - B) = \boxed{I - \bar{D}^{-1} \cdot B}$$

$$L_{\text{SYMMETRIC}} = \bar{D}^{-1/2} \cdot L \cdot \bar{D}^{-1/2} = \bar{D}^{-1/2} \cdot (D - B) \cdot \bar{D}^{-1/2} = \boxed{I - \bar{D}^{-1/2} \cdot B \cdot \bar{D}^{-1/2}}$$

SPECTRAL CLUSTERING Find the eigenvectors of normalized $L_{N \times N}$ matrix.

STEP #1: Find the eigenvectors of normalized $L_{N \times N}$ matrix.

STEP #2: Pick R smallest eigenvectors.

STEP #3: Construct Z matrix as follows:

$$Z = \begin{bmatrix} v_1 & v_2 & \dots & v_R \end{bmatrix}_{N \times R}$$

↳ 1st smallest eigenvector

↳ R^{th} smallest eigenvector

STEP #4: Run k-means clustering algorithm on Z matrix to find K clusters.

PARAMETERS:

δ : threshold.

R : # of eigenvectors to be included

K : # of clusters to be found.

HIERARCHICAL CLUSTERING

- finding groups such that instances (data points) in a group are more similar to each other than instances in different groups.

Component #1: The Distance Function Between Data Points

distance \Rightarrow dissimilarity

distance \uparrow similarity \downarrow
distance \downarrow similarity \uparrow

$$k(x_i, x_j) = \exp\left[-\frac{\|x_i - x_j\|_2^2}{2\sigma^2}\right]$$

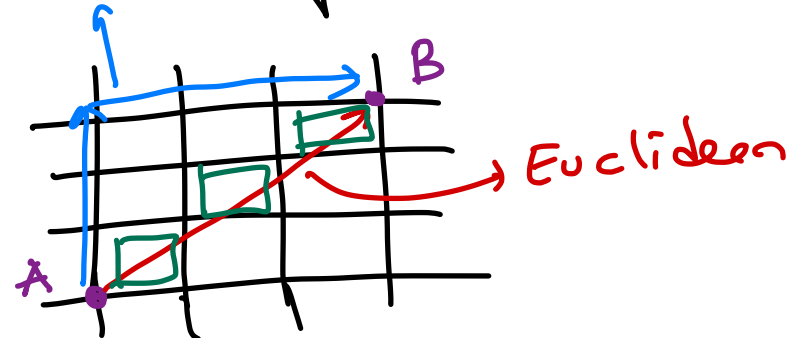
Diagram illustrating the Gaussian kernel function $k(x_i, x_j)$. Red arrows point from the components of the formula to their interpretations: 0 is labeled "dissimilar", 1 is labeled "similar", 0 is labeled "similar", and $+\infty$ is labeled "dissimilar". A double-headed arrow connects the 0 and 1 labels, with "dissimilar" and "similar" written below it. Purple arrows point from the 0 and $+\infty$ labels to the "dissimilar" label.

Manhattan Distance (city-Block Distance)

$$d(x_i, x_j) = \sum_{d=1}^D |x_{id} - x_{jd}|$$

Euclidean Distance

$$d(x_i, x_j) = \|x_i - x_j\|_2$$
$$= \sqrt{\sum_{d=1}^D (x_{id} - x_{jd})^2}$$
$$= \sqrt{x_i^T x_i - 2x_i^T x_j + x_j^T x_j}$$



Component #2: The Direction to Proceed

Agglomerative (bottom-to-top) Divisive (top-to-bottom)

⇒ combines small clusters into bigger ones
⇒ starts with "N" clusters.

⇒ divides big clusters into smaller ones
⇒ starts with "1" cluster

Component #3: The Distance Function Between Groups of Data Points.

Distance [{Paris, London}, {New York}]

Distance [{Paris, London}, {Rome}]

Distance [{London, Paris}, {Berlin, Rome}]



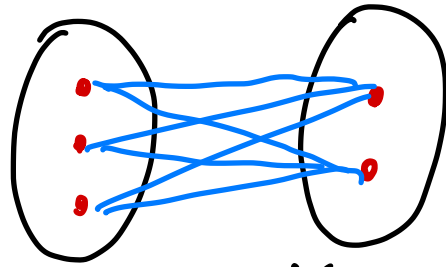
Centroid Clustering:

$$d(c_A, c_B) = \left\| \frac{\sum_{x_i \in C_A} x_i}{|C_A|} - \frac{\sum_{x_j \in C_B} x_j}{|C_B|} \right\|_2$$

Cardinality of C_A (# of members) \leftarrow centroid of C_A centroid of C_B

Single-Link Clustering:

$$d(c_A, c_B) = \min_{\substack{x_i \in C_A \\ x_j \in C_B}} d(x_i, x_j)$$



Complete-Link Clustering:

$$d(c_A, c_B) =$$

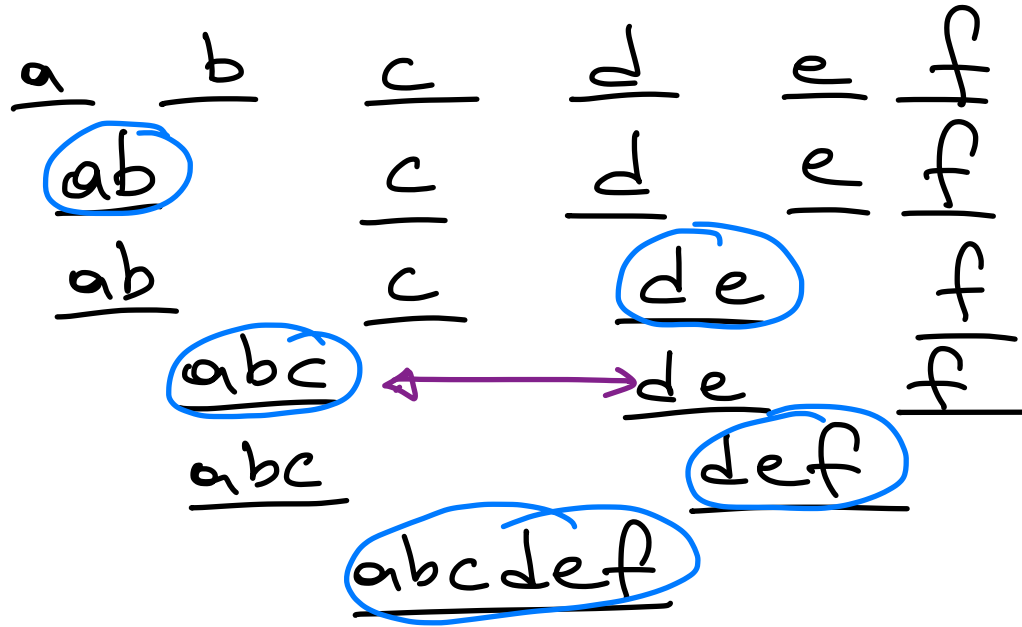
$$\max_{\substack{x_i \in C_A \\ x_j \in C_B}} d(x_i, x_j)$$

Average-Link Clustering:

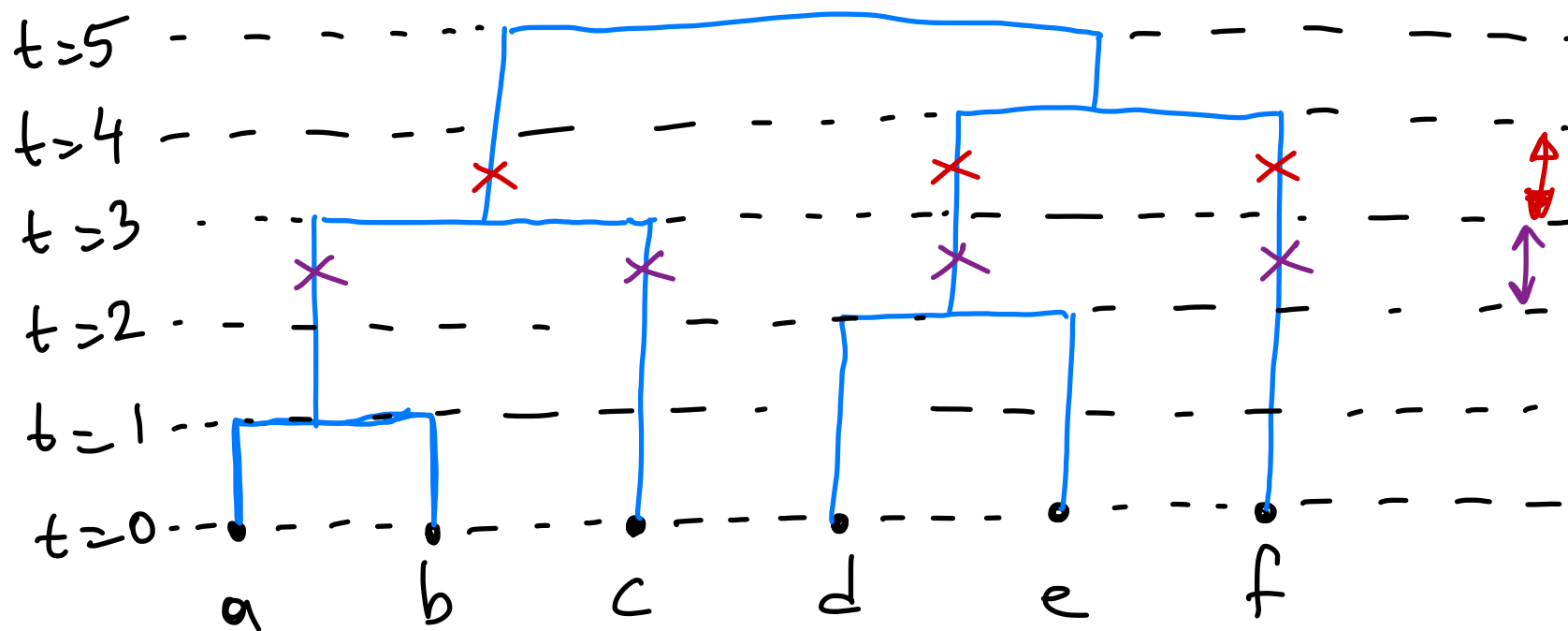
$$d(c_A, c_B) = \frac{\sum_{x_i \in C_A} \sum_{x_j \in C_B} d(x_i, x_j)}{|C_A| |C_B|}$$

•
•
•

$t=0$ 6 clusters
 $t=1$ 5 clusters
 $t=2$ 4 clusters
 $t=3$ 3 clusters
 $t=4$ 2 clusters
 $t=5$ 1 cluster



Dendrogram



$K=3$ clusters

$C_1 = \{a, b, c\}$

$C_2 = \{d, e\}$

$C_3 = \{f\}$

$K=4$ clusters

$C_1 = \{a, b\}$

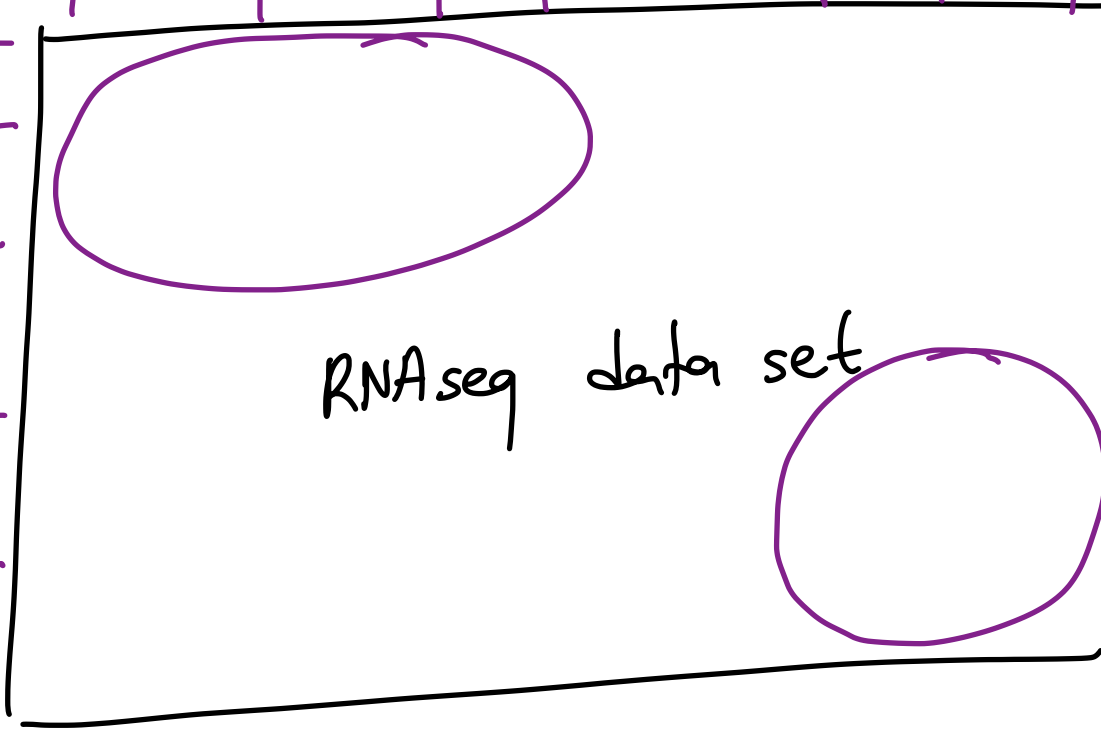
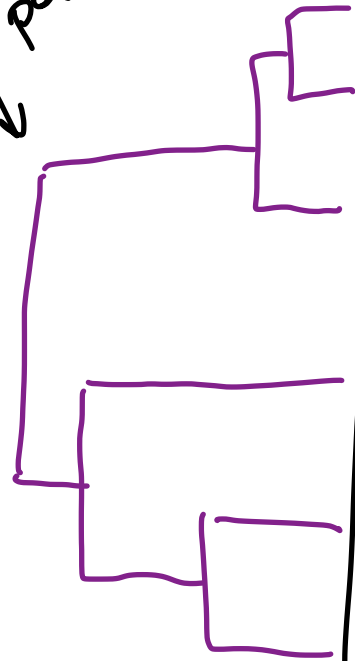
$C_2 = \{c\}$

$C_3 = \{d, e\}$

$C_4 = \{f\}$

clustering of
patients
↓

← clustering
of genes



RNAseq data set

patients

genes

"bi-clustering"