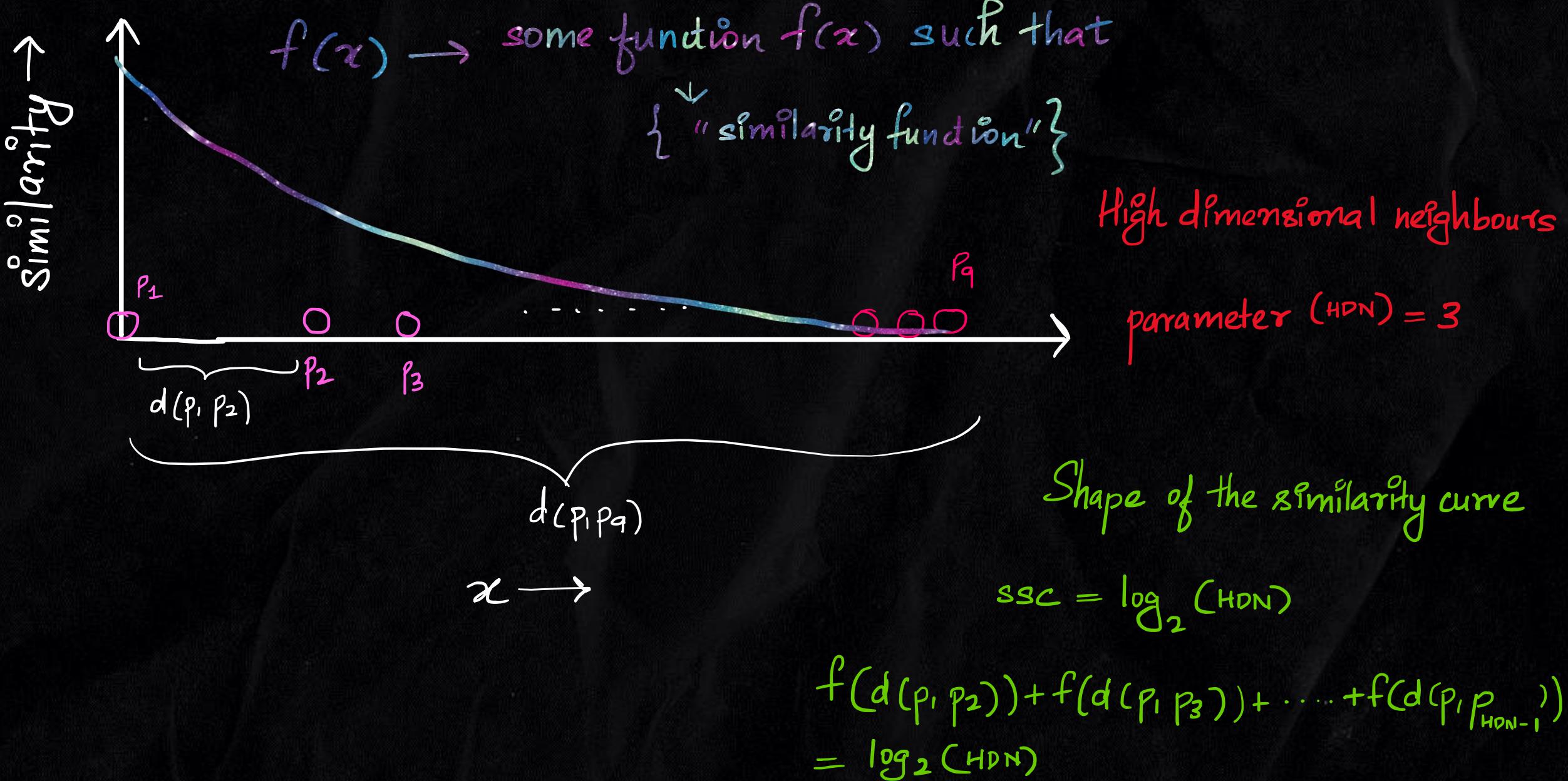


2) Calculating high dimensional similarity scores for each  $p_i \rightarrow$



$$f(x) = e^{-\frac{(\text{raw dist} - \text{dist to nearest neighbour})}{\sigma}}$$

$$f(p_i^o) = e^{-\frac{[d(o, p_i^o) - d(o, \text{nearest neighbour})]}{\sigma}}$$

We adjust  $\sigma$  such that

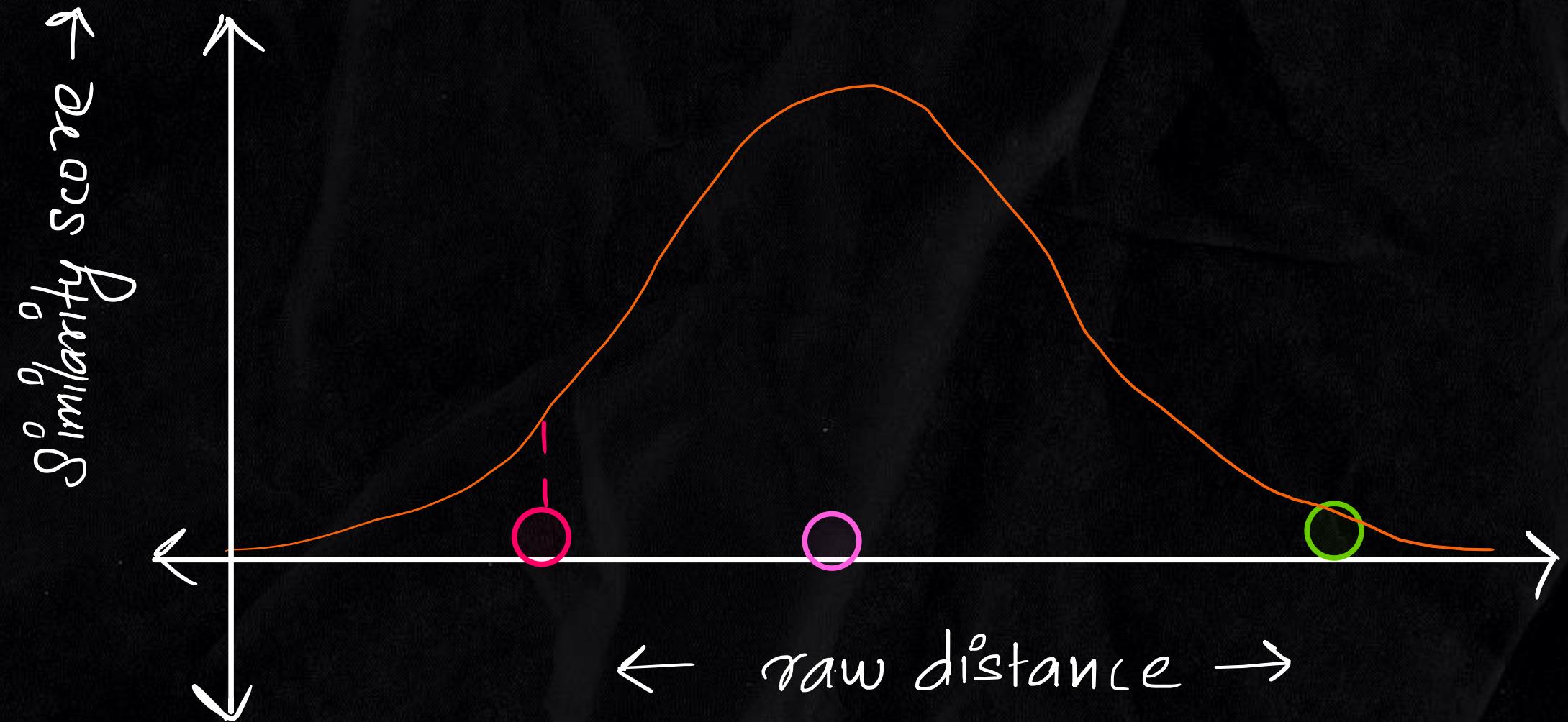
$$\sum_{i=1}^{\text{HDN}-1} f(p_i^o) = \log_2(\text{HDN})$$

Note :  $\rightarrow$  This is the reason why we get unique curves w.r.t each point.

\*

t-SNE  $\Rightarrow$  t-SNE differs from U-MAP

at this stage in that t-SNE uses a Gaussian distribution to calculate the similarity scores. The shape of the gaussian is controlled by the perplexity parameter { very similar to HDN in UMAP }.



← raw distance →

Shape of the curve determined by : $\rightarrow$  { HDN is the parameter that determines the shape of the curve }

$$\sum_{i=2}^{HDN-1} f(d(p_1 p_i)) = \log_2(HDN)$$

{ \* what is this function  
f or  $f^2$  ? }

$f(d(p_1 p_i))$  is the similarity score of point  $p_i$  w.r.t  $p_1$ .

Repeat step two for all the points

	$P_1$	$P_2$	$\dots$	$P_i^o$	$\dots \dots$	$P_n$
$P_1$						
$P_2$						
$P_j^o$	- - - - -		$f(d(p_i^o, p_j^o))$	$\Rightarrow$	similarity score of $p_j^o$ w.r.t $p_i^o$	
$\vdots$						
$P_n$						

HD

Similarity scores are not symmetrical  $\Rightarrow f(d(p_1, p_2)) \neq f(d(p_2, p_1))$

{you can also write this as  $f^*(p_1, p_2) \neq f^*(p_2, p_1)$ }

HD

UMAP makes the similarity score symmetrical by doing something like taking the average.

$$f(p_1 p_2) = \frac{f(p_1 p_2) + f(p_2 p_1)}{2} - f(p_1 p_2) \cdot f(p_2 p_1)$$

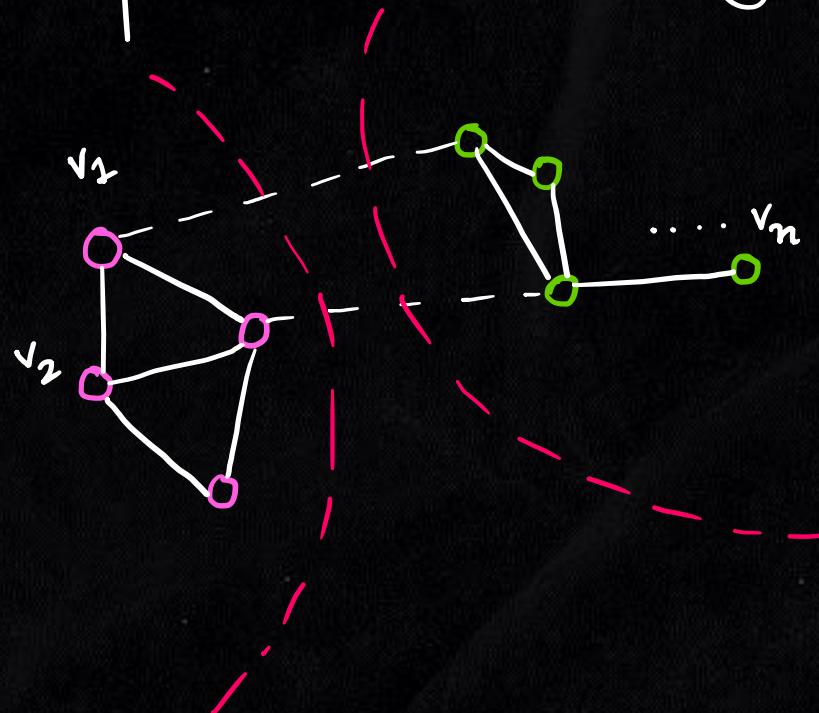
“Fuzzy Union Operation”

3)

Initialising a low dimensional graph {spectral embedding} : →



Spectral Embedding  $\rightarrow$



What is a good cut?

Matrix representation  
of an  
undirected  
graph

	$v_1$	$v_2$	$\dots$	$v_n$
$v_1$	0	1	$\dots$	0
$v_2$	1	0	$\dots$	0
$\vdots$	$\vdots$	$\vdots$	$\ddots$	0
$v_n$				0

$$\text{cut}(A) = \sum_{i \in A, j \notin A} w_{ij}$$

Criterion for good cut :  $\rightarrow$  Conductance / modularity

$$\phi(A) = \frac{|\{(i,j) \in E ; i \in A, j \notin A\}|}{\min(\text{vol}(A), 2m - \text{vol}(A))}$$

$m \rightarrow |E|$     ↴ this prevents you from choosing  
an unreasonably large  $A$

Adjacency matrix of a two component graph

		$v_1 \ v_2 \ \dots \ v_{n/2}$		$\dots \ \dots \ \dots \ v_n$	
$v_1$	$v_1$	0 1 1	0 0 0		
	$v_2$	1 0 1	0 0 0		
	$\vdots$	1 1 0	0 0 0		
	$v_{n/2}$	0 0 0	0 0 0		
$\vdots$	$v_n$	0 0 0	0 0 0		
		0 0 0	1 1 1		
		0 0 0	1 1 1		
		0 0 0	1 1 0		

Clusters

$$\begin{bmatrix} a_{11} & \cdots & \cdots & \cdots & a_{1n} \\ \vdots & & & & \vdots \\ a_{n1} & & & & a_{nn} \end{bmatrix} \begin{bmatrix} x_1 \\ \vdots \\ x_n \end{bmatrix} = \begin{bmatrix} y_1 \\ \vdots \\ y_n \end{bmatrix}$$

$y_i^o = \sum_{j=1}^n A_{ij}^o x_j^o = \sum_{(i,j) \in E} x_j^o$

for a vertex  $x_i^o$  this is  
 the sum of the vertex  
 values of all its  
 neighbours  $x_j^o$ .

Spectral Graph Theory :→

set of eigenvalues in ascending order  
and their corresponding eigenvectors.

Analyse the spectrum of matrix representing  
 $G_1$ .

Spectrum :→

$$\Lambda = \{ \lambda_1, \lambda_2, \dots, \lambda_n \} \quad \lambda_1 \leq \lambda_2 \dots \leq \lambda_n$$

$$A \times x_i = \lambda_i \cdot x_i$$

Consider a  $d$ -regular graph  $G_1 \rightarrow$

$$A \times x = \lambda x$$

solutions to this?  $\lambda?$   $x?$

For a  $d$ -regular graph,  $x = \underbrace{[1, 1, \dots, 1]}_n \left\{ [1]^n \right\}$  &  $\lambda = d$  is a solution

Now consider a graph with 2  $d$ -regular components.

Suppose you assign node values s.t

$$\text{Node value } (x) = \begin{cases} 1 & \text{if } x \in 1^{\text{st}} \text{ component (I)} \\ 0 & \text{if } x \in \text{II (2}^{\text{nd}} \text{ component)} \end{cases}$$

Then, the vector looks something like this:  $\rightarrow$

$$x^1 = [1, 1, \dots, 0, 1, 0, 0, \dots, 1]$$

$$A \times x' = [d, d, \dots 0, d, 0, 0, \dots d]$$

$\lambda_1 = \lambda_2 = d$ .  
some intuition :→

I haven't got this  
intuition fully, yet

If the two components are poorly connected rather  
than disconnected, then,

$$\lambda_1 \approx \lambda_2$$

Eigenvectors  $\Rightarrow$  They are real & orthogonal

$$\left\{ x_i^o \cdot x_j^o = 0 \right\}$$

	$v_1$	$v_2$	$\dots$	$\dots$	$v_n$
$v_1$	$d_{v_1}$	0	0	0	0
$v_2$	0	$d_{v_2}$			$\vdots$
$\vdots$	0				$\vdots$
$v_n$	0				$d_{v_n}$

$\rightarrow$  Degree matrix

Graph Laplacian matrix  $L \rightarrow$

$$L = D - A$$

	$v_1$	$v_2$	$\dots$	$\dots$	$v_n$
$v_1$	$d_{v_1}$	-1	0	-1	0
$v_2$	-1	$d_{v_2}$			
:	:	0	:	:	:
$v_n$	-1			0	$d_{v_n}$

Sum of each row  
and column is  
zero

What are the eigenvectors and eigenvalues of  
the L matrix?

Trivial solution exists,  $x = [1]^n$

$$Lx = [0]^n, \lambda = 0 = \lambda_1$$

{smallest eigenvalue}

Important fact that we will use without proving :→

$$\lambda_2 = \min_{\mathbf{x}} \frac{\mathbf{x}^T \mathbf{M} \mathbf{x}}{\mathbf{x}^T \mathbf{x}}$$

$\mathbf{M} \rightarrow$  matrix      What is  $\mathbf{x}^T \mathbf{L} \mathbf{x}$ ?

$$\mathbf{x}^T \mathbf{L} \mathbf{x} = \sum_{i,j=1}^n L_{ij} x_i x_j = \sum_{i,j=1}^n (D_{ij} - A_{ij}) x_i x_j$$

$$= \sum_{i^o} D_{ii^o} x_i^o - \sum_{(i^o, j^o) \in E} x_i^o x_j^o$$

$$= \sum_{(i^o, j^o) \in E} (x_i^o)^2 + (x_j^o)^2 - 2x_i^o x_j^o = \sum_{(i^o, j^o) \in E} (x_i^o - x_j^o)^2$$

$x^o$  is a unit vector since  $x$  is an eigenvector

$x^o$  is orthogonal to the first eigenvector  $[1]^n$

$$\Rightarrow \sum^l x_i^o = 0$$

$$\lambda_2 = \min_{\mathbf{x}} \frac{\mathbf{x}^T L \mathbf{x}}{\mathbf{x}^T \mathbf{x}}$$

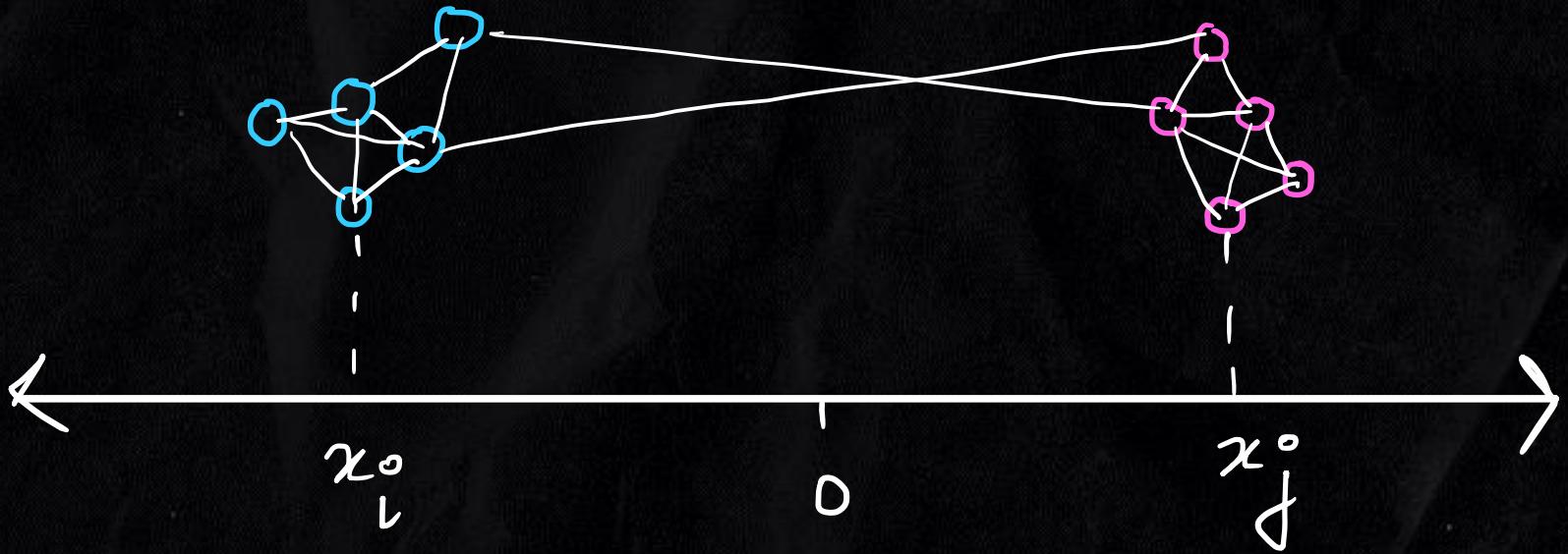
$$= \min_{\substack{\text{all labelling} \\ \text{of nodes}}} \frac{\sum_{(i,j) \in E} (x_i^o - x_j^o)^2}{\sum_i x_i^o} \rightsquigarrow 1 \quad \left\{ \begin{array}{l} \text{since } \mathbf{x} \\ \text{is a} \\ \text{unit vector} \end{array} \right\}$$

so that  $\sum_i x_i^o = 0$

$$\lambda_2 = \min_{\substack{\text{all labelling} \\ \text{of node } i}} \sum_{(i,j) \in E} (x_i^o - x_j^o)^2$$

so that

$$\sum x_i^o = 0$$



So for each  $e \in E$ , we would like  
 $e_{ij}$ ,  $i, j \in$  same cluster i.e same side  
of  $O$ .

When we are punished the most? when we  
have an edge  $e_{ij}$  where

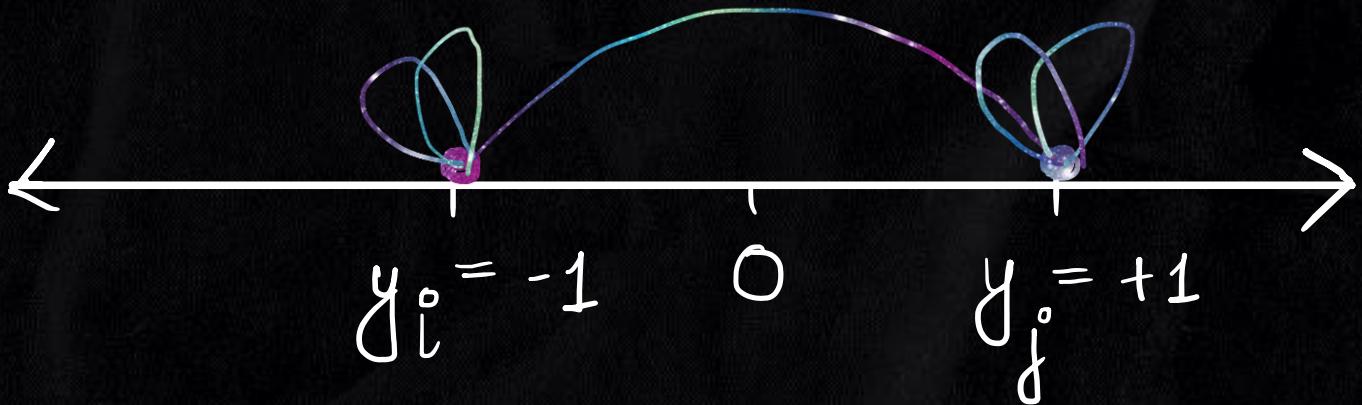
$$x_i^o < 0, x_j^o > 0$$

If given a graph & asked to find 2 clusters  
in it, then

express the two clusters as a vector

$$y_i^o = \begin{cases} 1 & \text{if } i \in A \\ -1 & \text{if } i \in B \end{cases}$$

Best clustering =  $\min_y \sum_{(i,j) \in E} (y_i^o - y_j^o)^2$



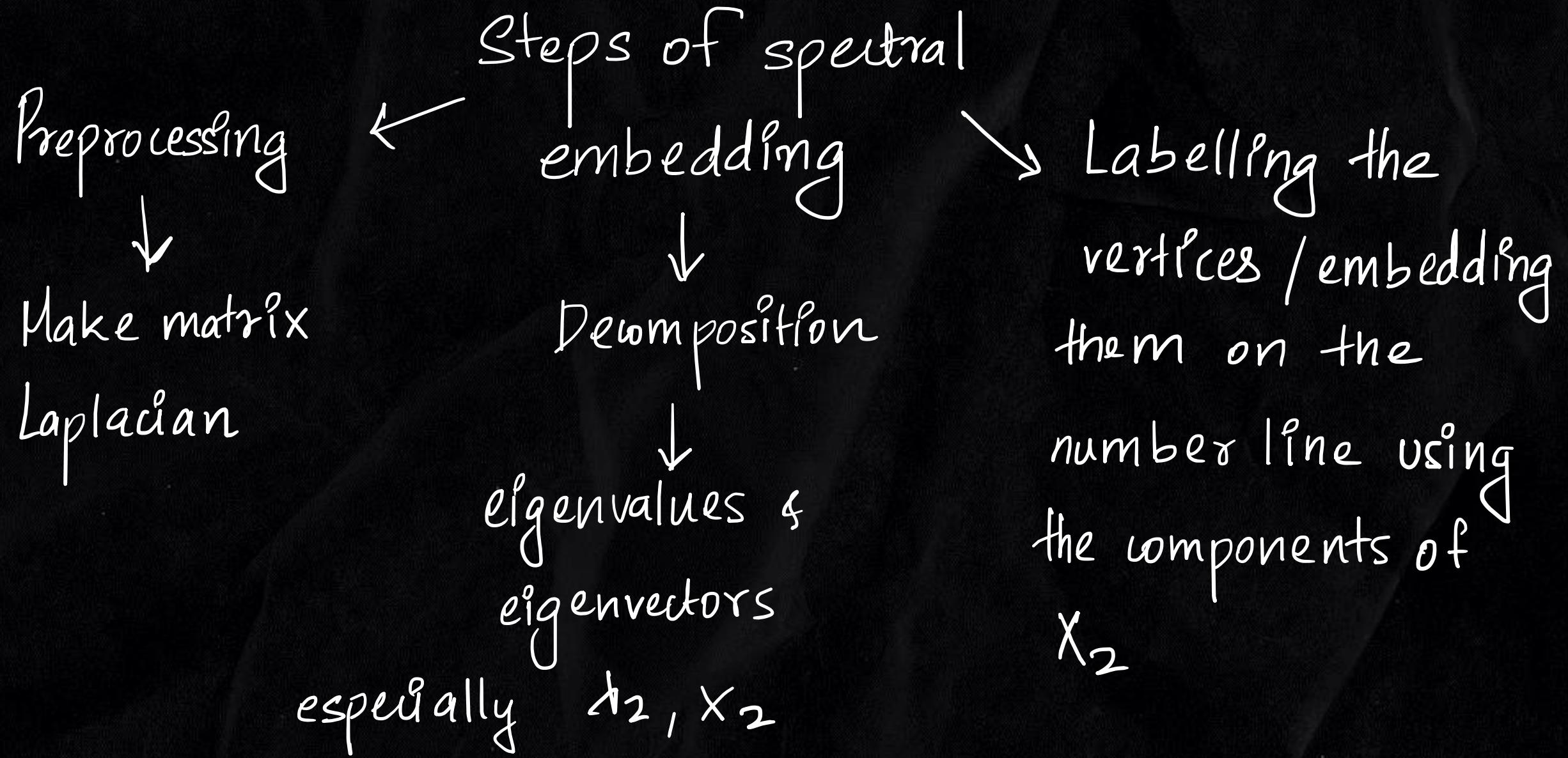
Relaxation  $\Rightarrow$

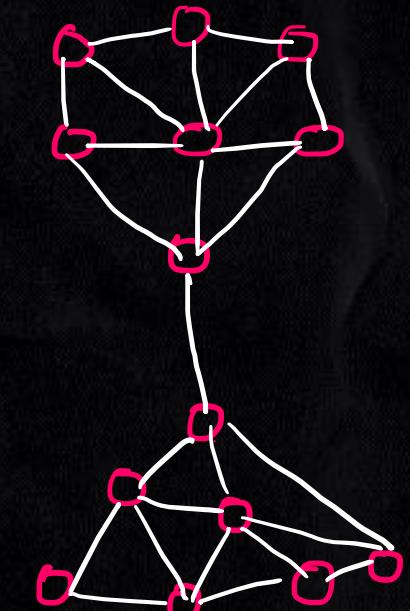
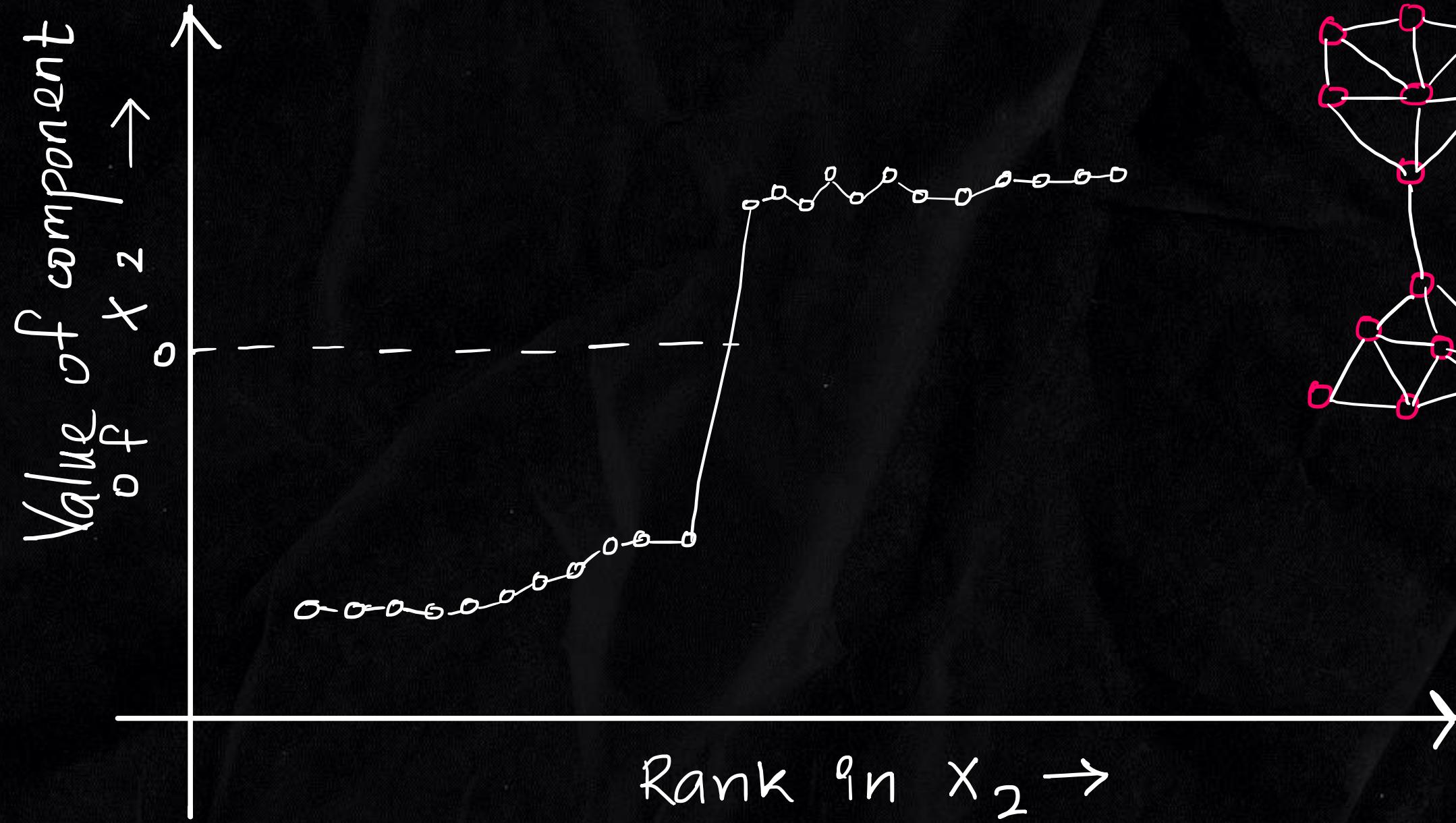
instead of  $y_i^o \in \{-1, 1\}$

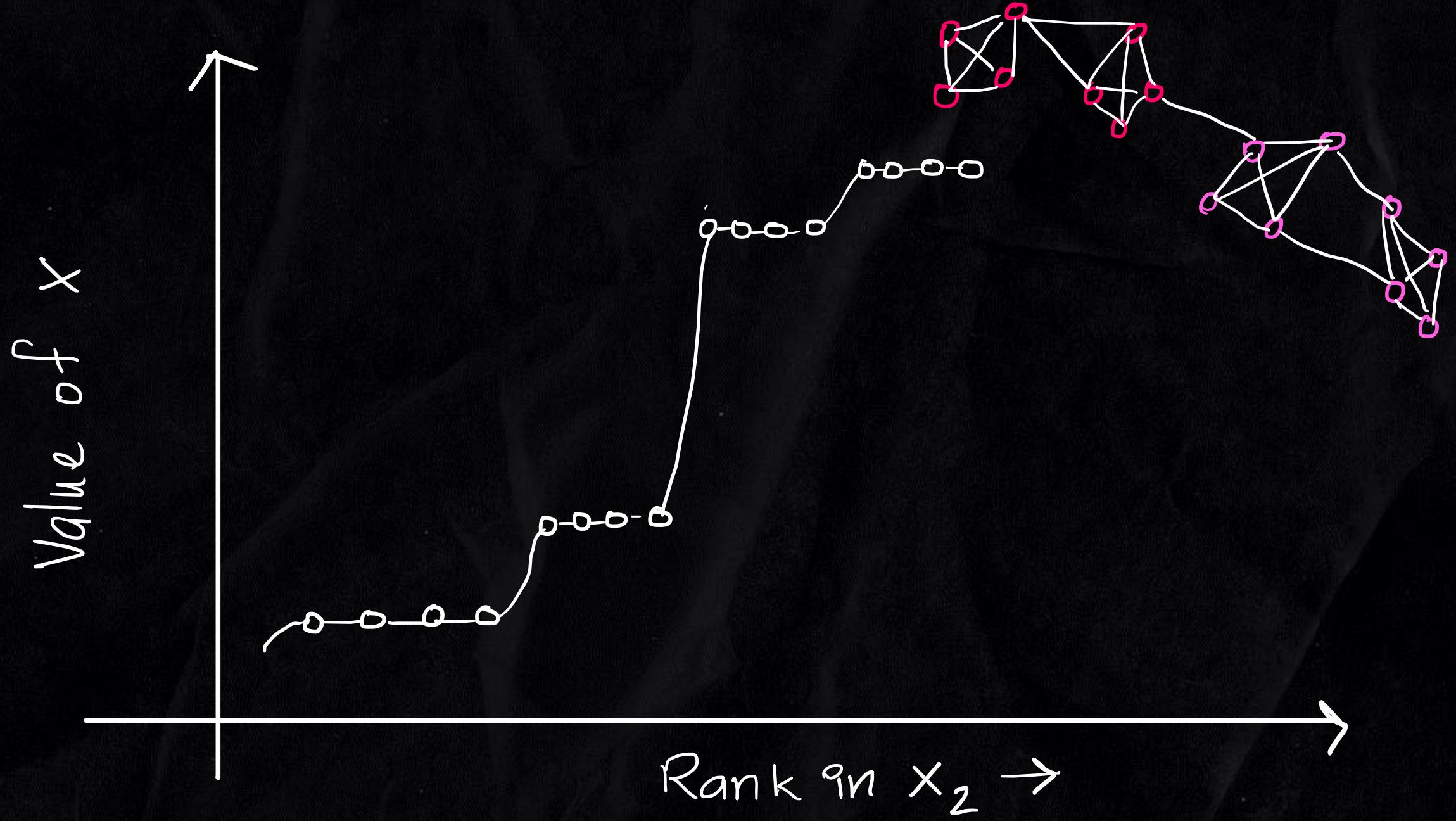
$$y_i^o \in \mathbb{R}$$

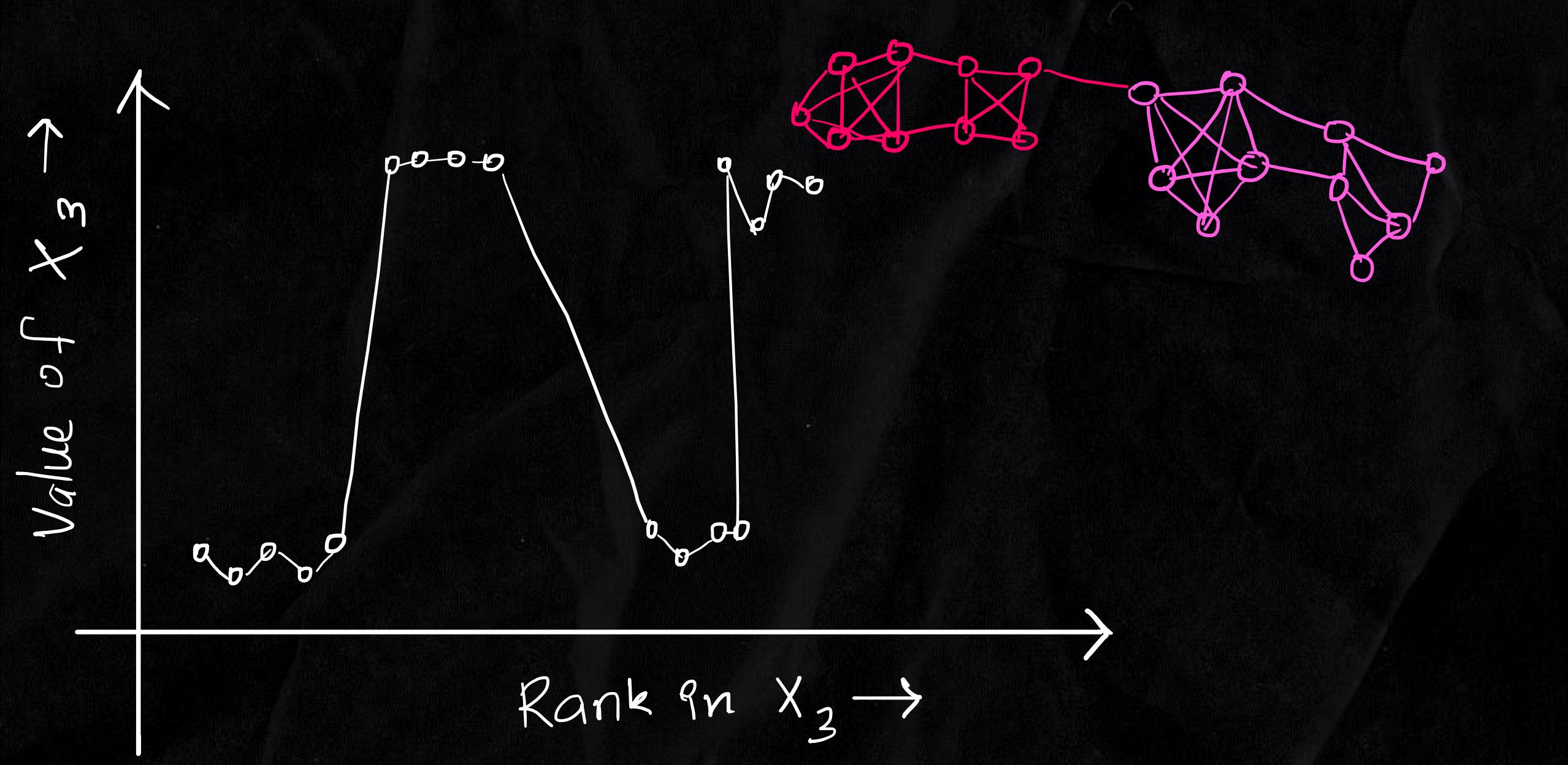
The vector  $\gamma$  that solves this minimization is the second smallest eigenvector, also known as the Fiedler vector.

$$P \cdot \pi^* = 0$$









Partitioning a graph into k-clusters : →

Express each node of the graph as a vector

for node  $i \rightarrow [x_{2i}, x_{3i}, \dots, x_{ni}]$

$x_{ji} \rightarrow$   $\downarrow$   
 $i^{th}$  component of the  $j^{th}$  smallest  
eigenvector

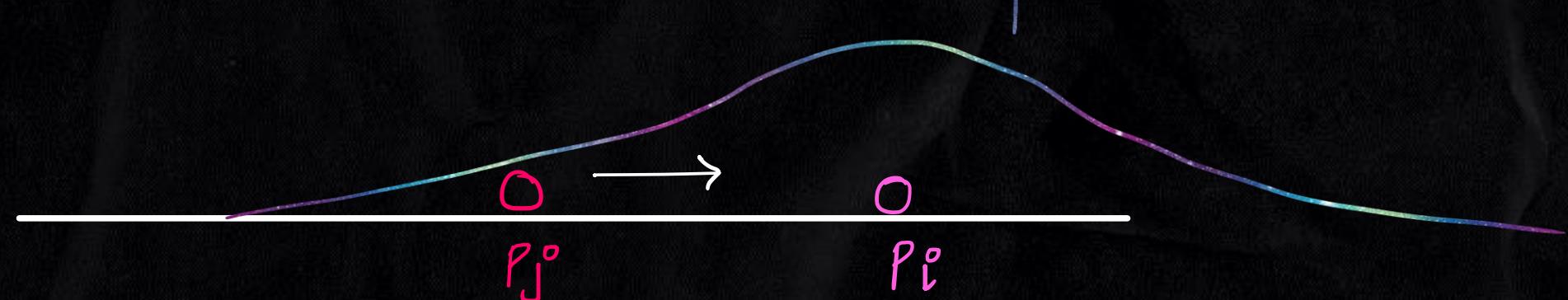
Now that each node is expressed by a vector  
we can use k-means to identify the clusters  
in the n-dimensional space.

## 5) Adjustments

- Randomly choose 2 points within a cluster  $\{ p_i \text{ and } p_j \}$ 
  - { how does UMAP know whether they are in the same cluster or not? }
  - non-zero similarity score
- Randomly choose to move one of the two points  $\{ p_j \}$
- Choose a point from another cluster that the first point should move away from.  $\{ p_k \}$ 
  - ↓ HD
  - { using similarity scores? }
- Calculate low dimensional similarity scores for  $p_j, p_k$  &  $p_j, p_i$   
(LD)

5.1) Calculating LD similarity score :→

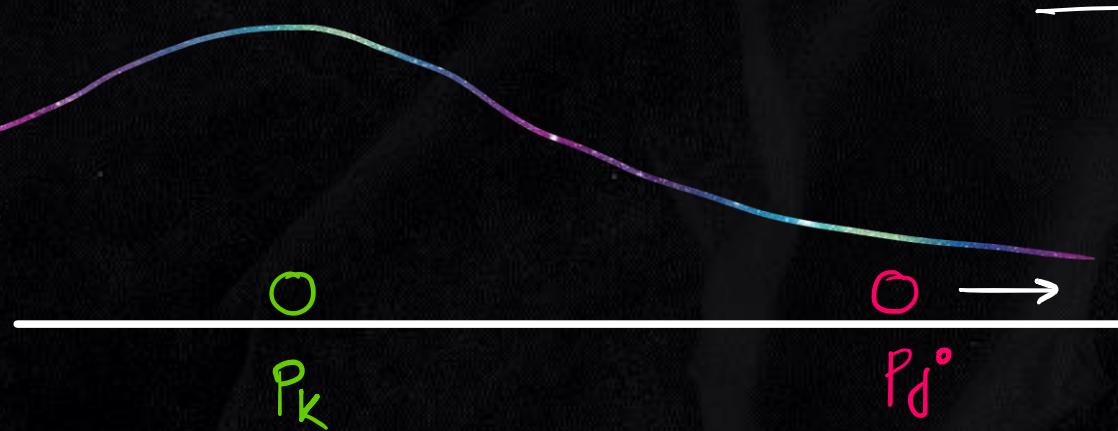
→ t-distribution



Increases the LD similarity

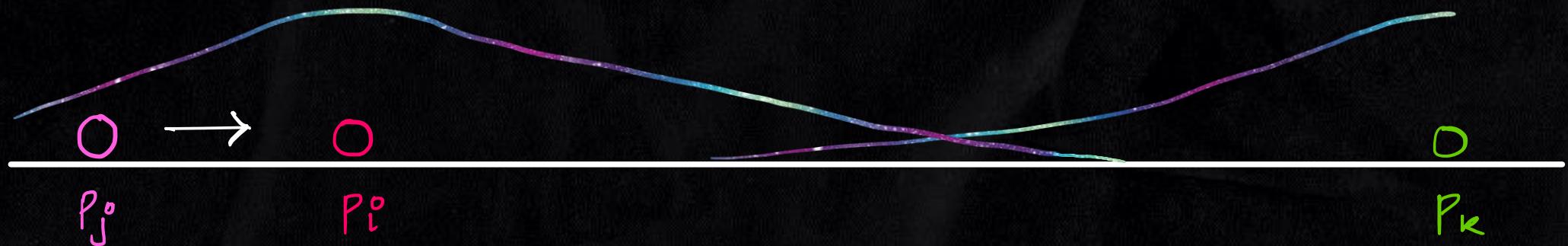
score of  $p_j^o, p_i^o$  &

decreases LDSS  $p_j^o, p_k$ .



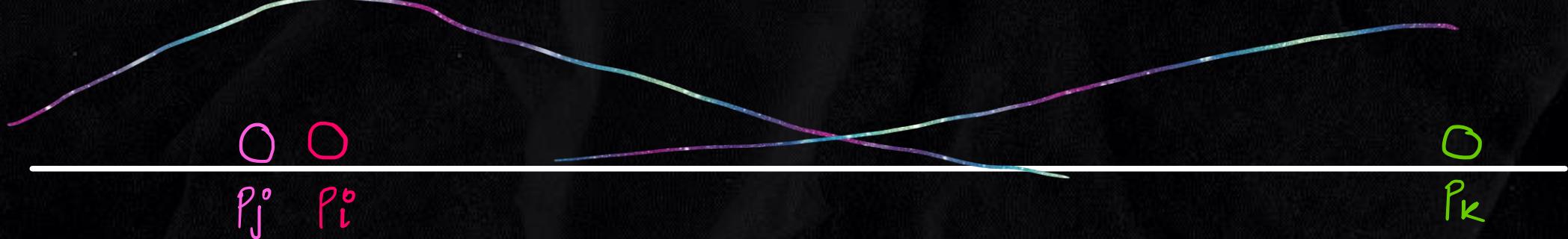
{ how much to move? A lil bit, how little? }

Alternate case  $\rightarrow$



$$f_{LD}(P_j^o P_i^o) = \chi$$

$$\downarrow \quad f_{LD}(P_j^o P_k^o) \approx 0$$



$$f_{LD}(P_j^o P_i^o) > \chi$$

$$f_{LD}(P_j^o P_k^o) \approx 0$$

LDSS between  $p_i^o$  &  $p_j^o$        $\alpha, \beta \rightarrow$  user defined parameters

$$\text{LDSS}(p_i^o, p_j^o) = \frac{1}{1 + \alpha d^*(p_i^o, p_j^o)^{2\beta}}$$

$d^*(p_i^o, p_j^o)$   $\rightarrow$  LD distance b/w  $p_i^o$  &  $p_j^o$

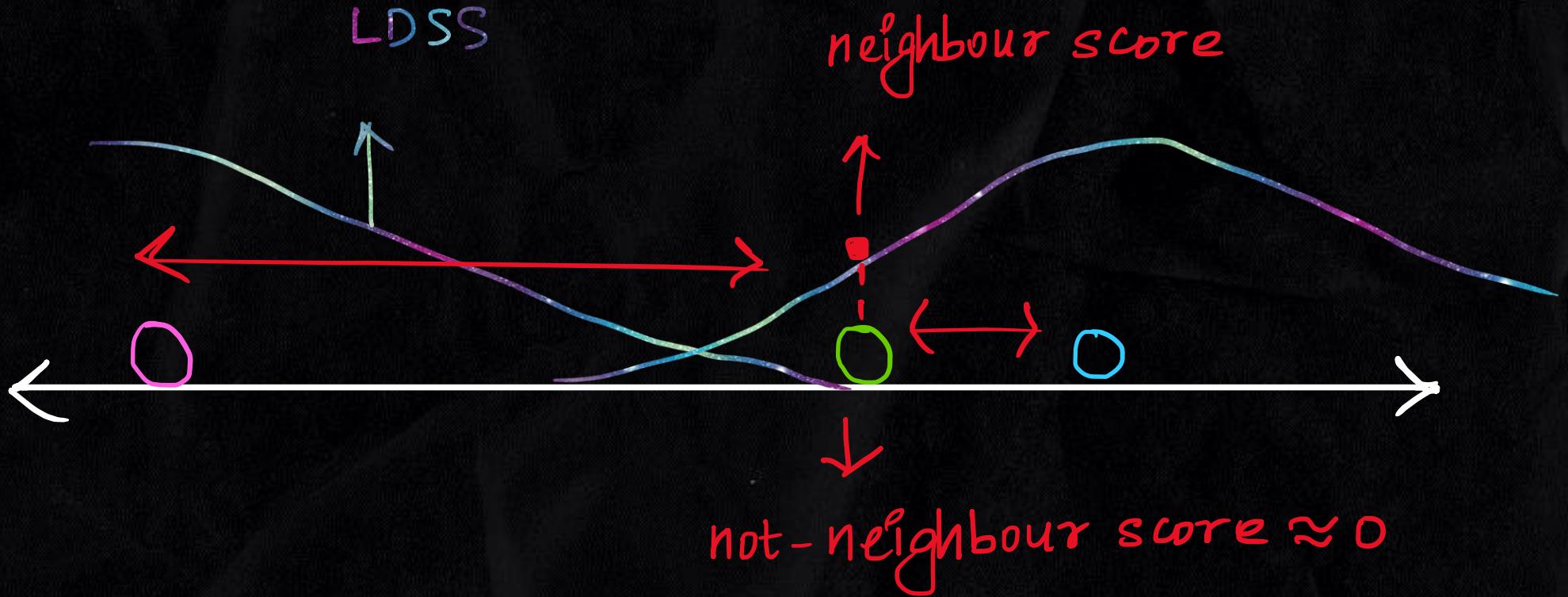
by default

$$\alpha = 1.577$$

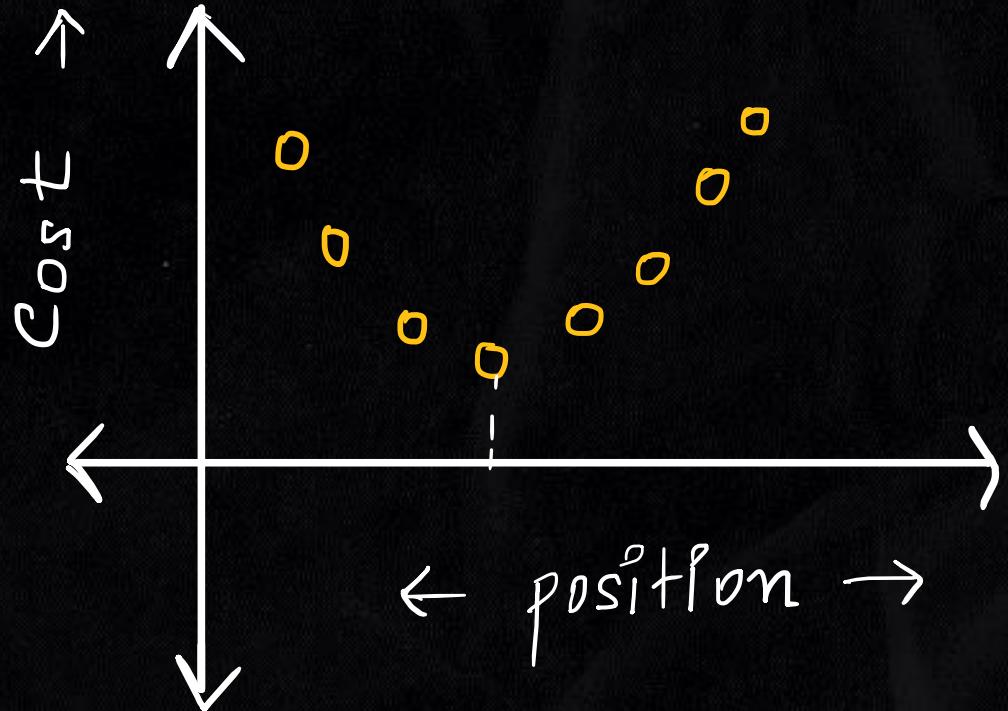
$$\beta = 0.8951$$

t-SNE uses  $\alpha = \beta = 1$

that decide the minimum distance b/w points on the LD graph & their spread.



$$\text{Cost} = \log\left(\frac{1}{\text{neighbour}}\right) + \log\left(\frac{1}{1-\text{not neighbour}}\right)$$



Cost vs Position for a  
particular point

UMAP uses stochastic gradient  
descent to optimise the cost.

- Move the points based on calculated LDSS
- ⑤ repeat till desired same number of clusters as that in the HD data is achieved.