

HW2 Part 2

① $\{x_i\}_{i=1}^N$

Data Likelihood
of data

$$L = \prod_{i=1}^N p(x_i | \theta)$$

①.1 App.1: MLE : finds the parameters of
the data likelihood

$$\arg \max_{\theta} L$$

①.2 App.2: MAP : finds the parameters of
data likelihood assuming
a prior distribution

$$L_2 = \prod_{i=1}^N p(x_i | \theta) \cdot p(\theta | \alpha, \beta)$$

data likelihood

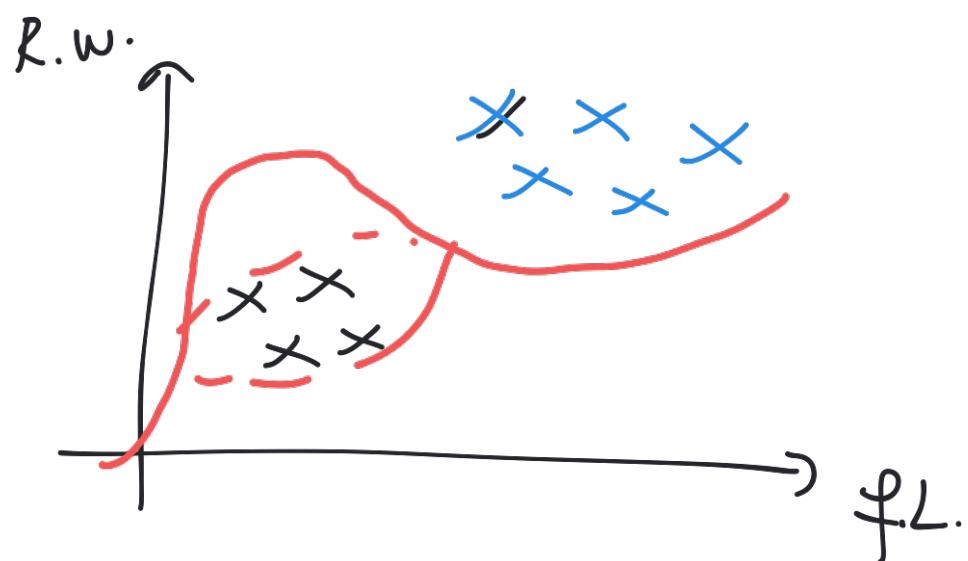
prior is
Gamma
distribution

$$\arg \max_{\theta} L_2$$

(2)	SPECIES	Frontal zip	rear width	Length	width	Depth	Mole	Fenest.
1	0							
2	1							
:	:							
140	0							
141								
:								
200								

FEATURE Matrix X

$\mathcal{L}_0 = \text{multivariate-gaussian}(\text{training}, M_0, \Sigma_0).$



- ① Find Mean and covariance for samples in species 0
- ② find Mean and covariance for samples in species 1

$$G(x | C_0) \times p(C_0) > G(x | C_1) \times p(C_1)$$

multivariate-gaussian C_1 training data

If we use 7 features,
we will obtain an error

"singular matrix"

$$\Rightarrow \det = 0$$

\Rightarrow inverse does not
exist

Multivariate
Gaussian

$$p(x|\mu, \Sigma) = \frac{1}{(2\pi)^{n/2} \cdot |\Sigma|^{1/2}} \cdot \exp\left(-(x-\mu)^T \Sigma^{-1} (x-\mu)\right)$$

If a matrix Σ is singular,

Σ^{-1} does not exist

$$(\Sigma + \alpha \cdot I)^{-1}$$

Lectures
6, 7

K-Means Obj. Fct:

$$J(\theta, \nu) = \sum_{i=1}^N \sum_{k=1}^K u_{ik} \cdot d(x_i, \theta_k)$$

Metric to define similarity between points
↓
DISTANCE METRIC

Crisp assignment: $u_{ik} = \{0, 1\}$, $\sum_{k=1}^K u_{ik} = 1$

MEMBERSHIP matrix U :

If $x_1 \in C_2$

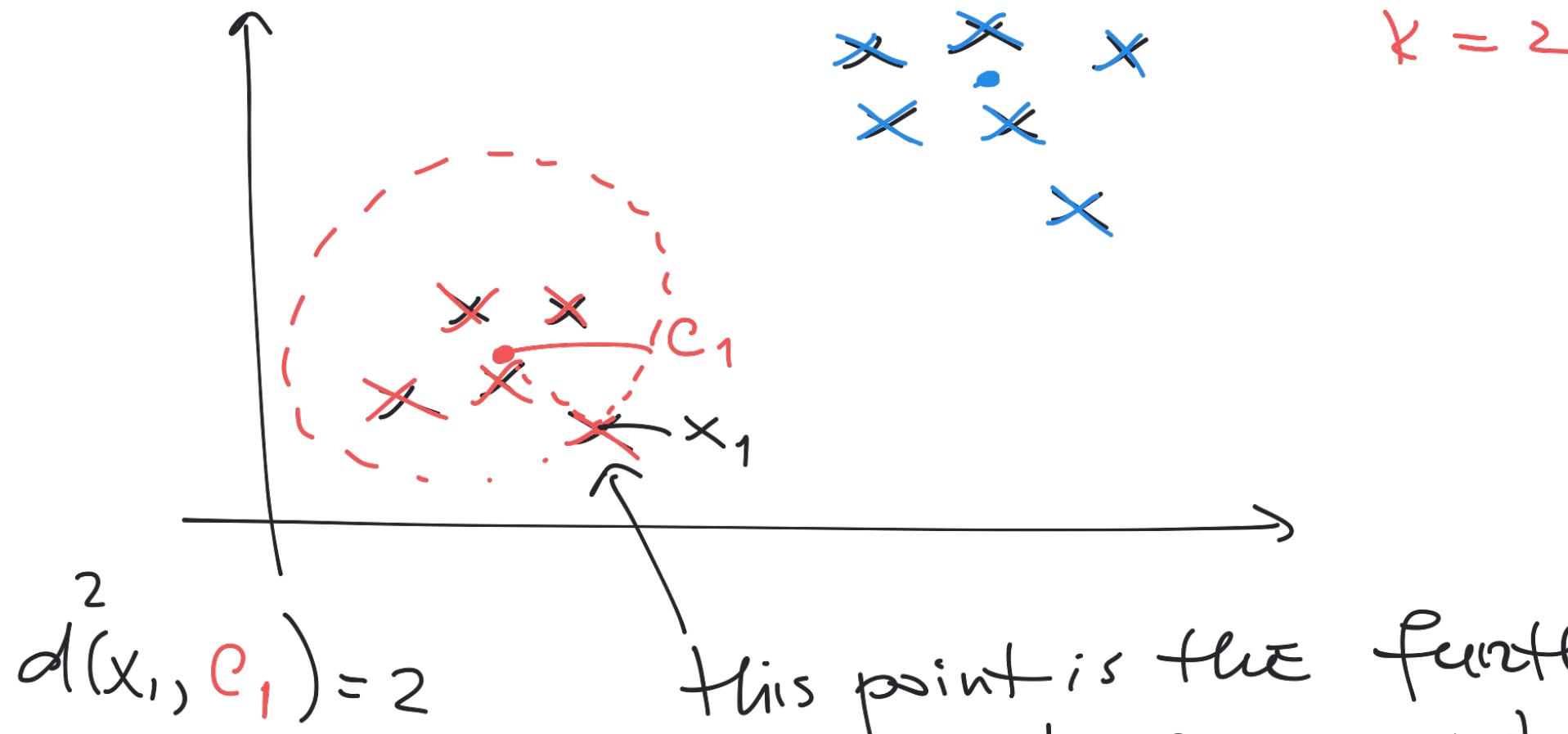
If $x_2 \in C_K$

	x_1	x_2	\vdots	x_n	1	2	$3..K$
$N \times K$	0	1			0	0	0
	0	0			0	0	0
					0	0	1

$$\sum_{i=1}^N u_{ik} = N_k$$

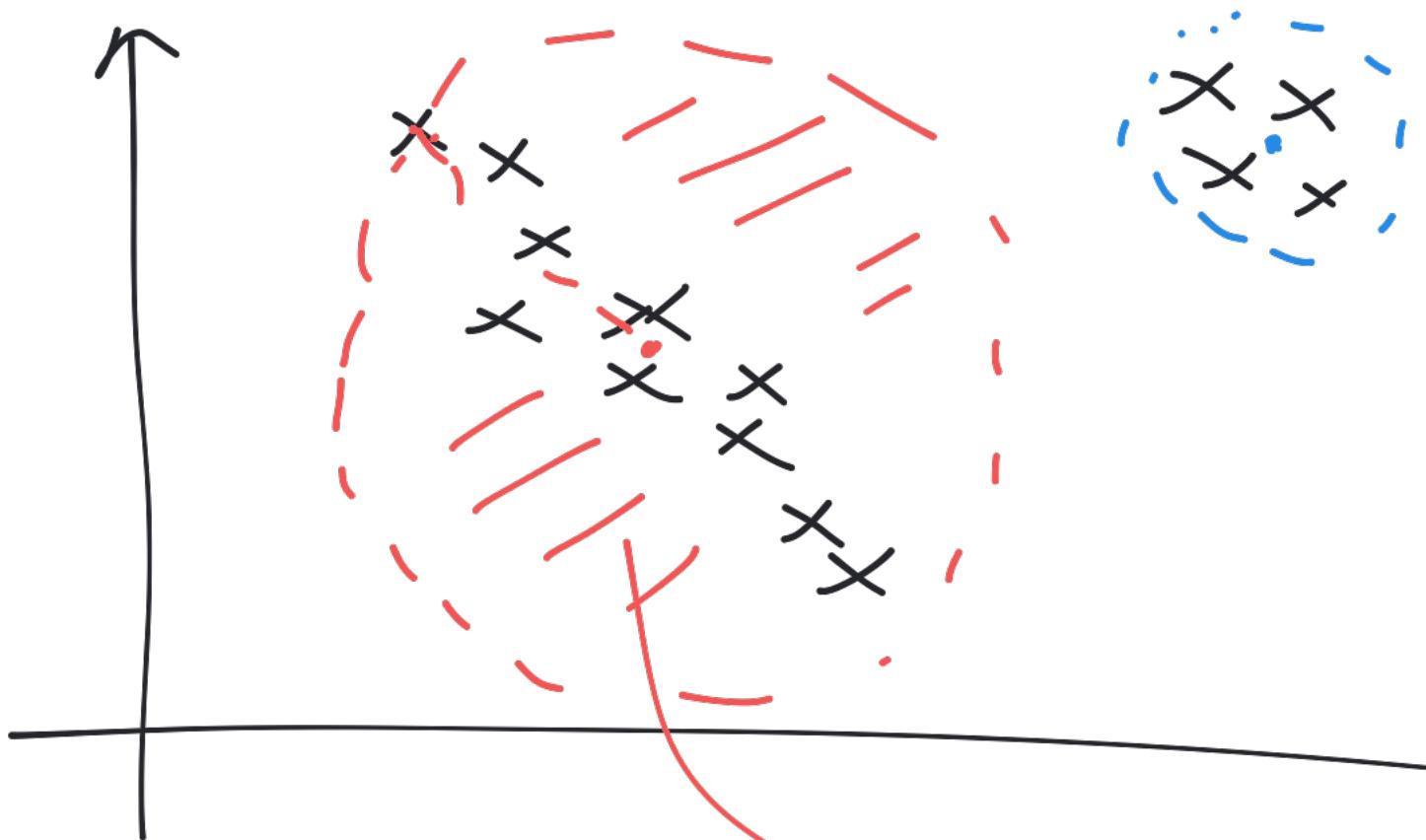
$N_k = \# \text{ of samples in cluster } k$

$$\text{If } d^2(x_i, \theta_k) = \|x_i - \theta_k\|^2$$



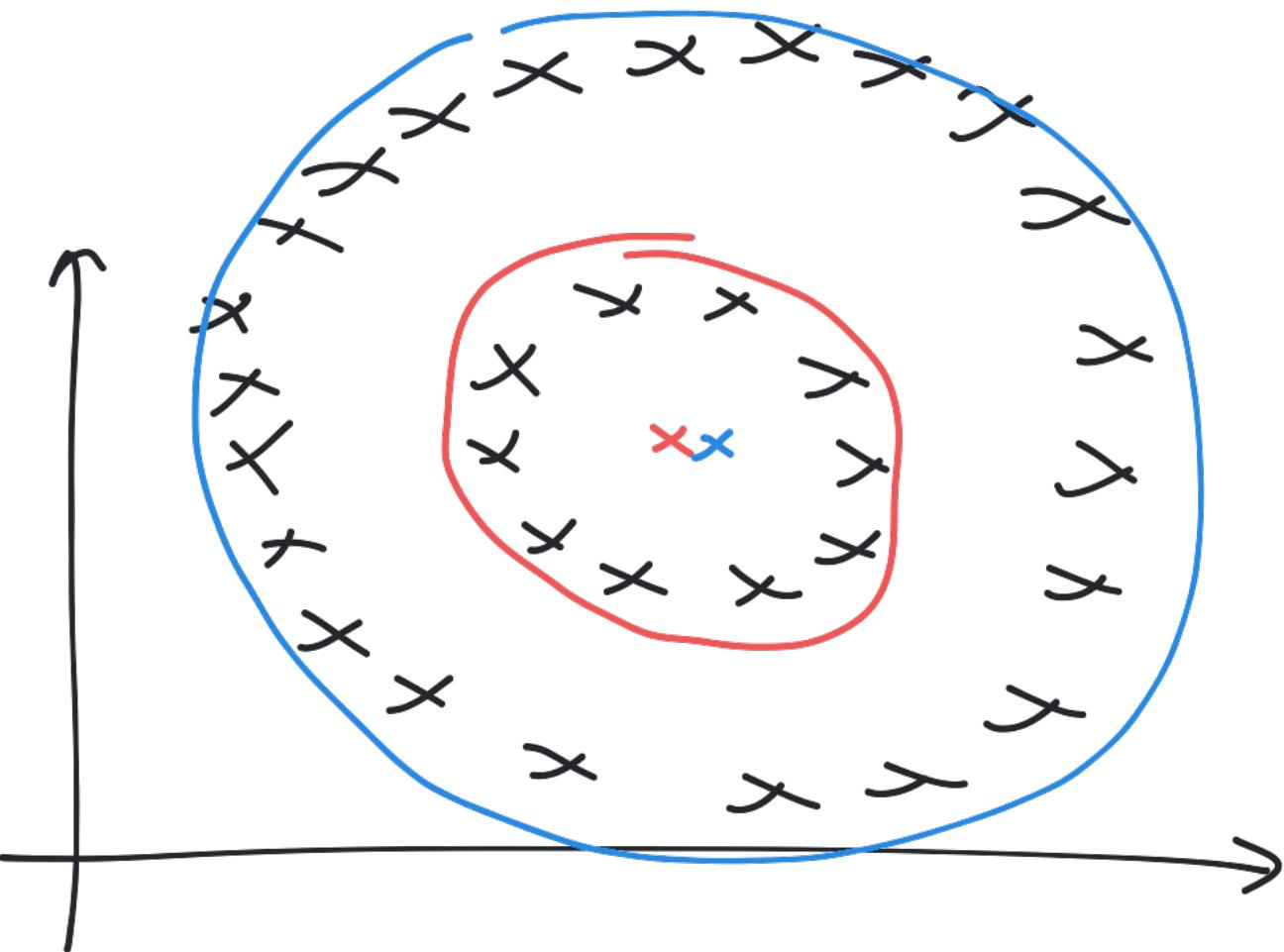
this point is the furthest point for centroid assign to this cluster

If distance is Euclidean then the clusters will be spherical.



WE do not have
samples.

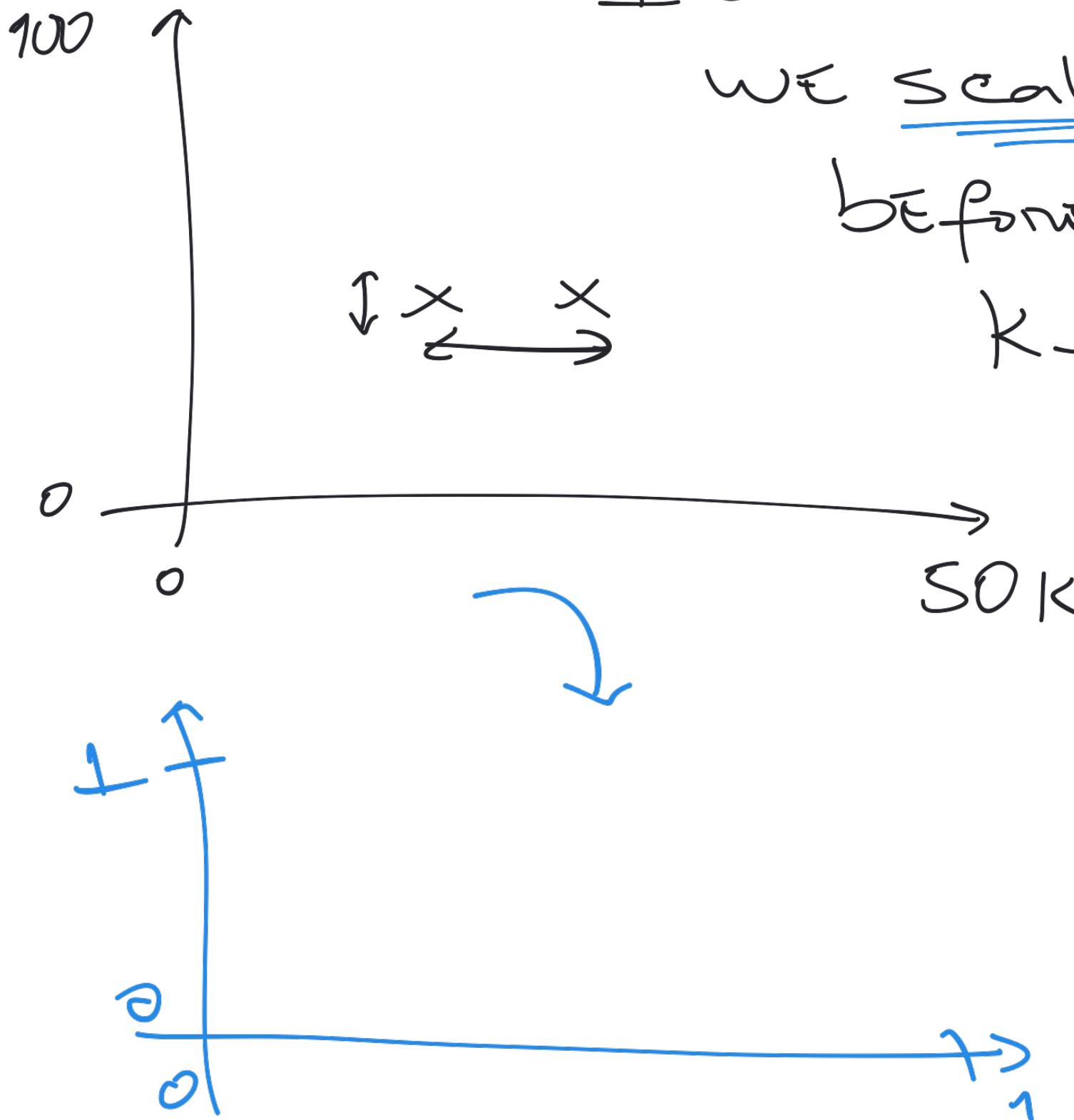
yet, k-means will cluster
any point in that region
as belonging to group
Red.



In practice we will have high-dimensional data
and so we will not be able to plot.
So we need to define quantitative
metrics to validate the clustering
results.

$$J = \sum_i \sum_k u_{ik} d^2(x_i, \theta_k)$$

It is crucial that
we scale the data
before applying
K-Means.



① Find Membership Matrix:

$$\frac{\partial J}{\partial u_{ik}} = 0 \Rightarrow$$

Assign point x_i
to the cluster
with closest
centroid θ_k

② $\frac{\partial J}{\partial \theta_k} = 0 \Leftrightarrow \sum_{i=1}^N u_{ik} \cdot 2 \cdot (-1) \cdot (x_i - \theta_k) = 0$

$$\Leftrightarrow \sum_{i=1}^N u_{ik} (x_i - \theta_k) = 0$$

$$\Leftrightarrow \sum_{i=1}^N u_{ik} \cdot x_i - \sum_{i=1}^N u_{ik} \cdot \theta_k = 0$$

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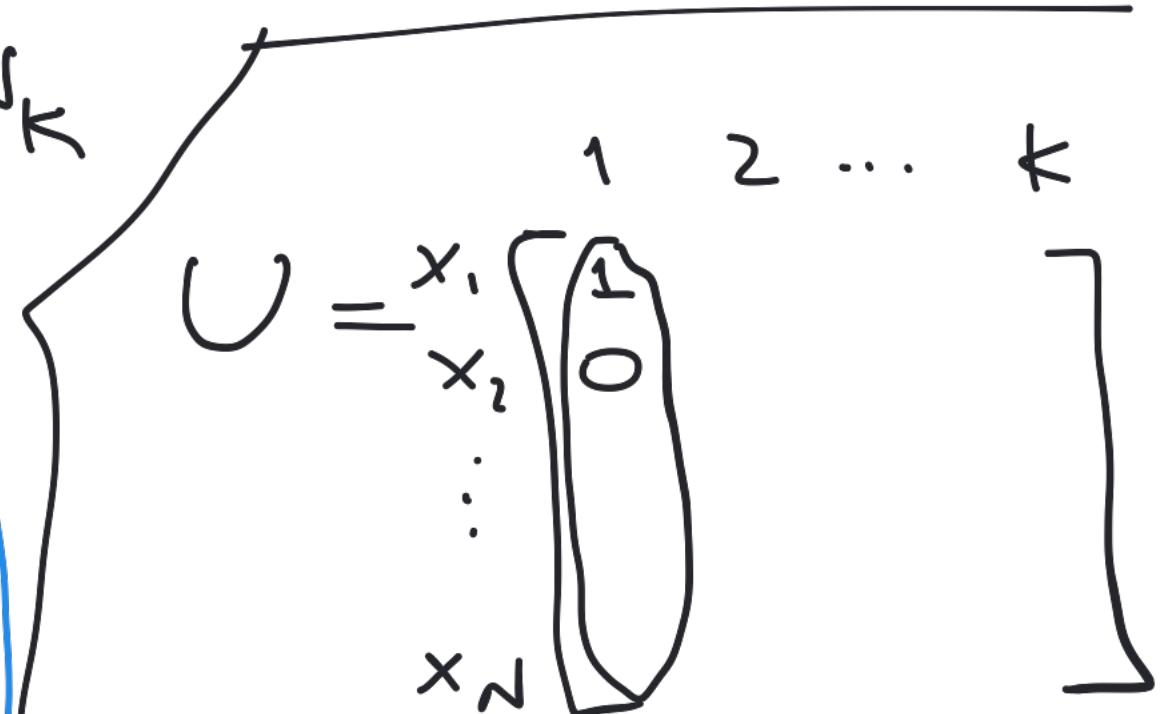
$$\left[\sum_{i=1}^n u_{ik} x_i \right] = \vartheta_k \left[\sum_{i=1}^n u_{ik} \right]$$

*

$$\Leftrightarrow \sum_{n \in C_k} x_i = \vartheta_k \cdot z_k$$

$$\Leftrightarrow \vartheta_k = \frac{1}{N_k} \cdot \sum_{n \in C_k} x_i$$

\hookrightarrow average for samples
belonging to cluster k .



$N_k \equiv$ # of points
assigned to
cluster k

$$\begin{aligned} \left[\sum_{i=1}^n u_{ik} x_i \right] &= 0 \cdot x_1 + 1 \cdot x_2 + 1 \cdot x_3 + \dots \\ &= \sum_{n \in C_k} x_n \end{aligned}$$