

# ***MACHINE LEARNING***

## **Kernel for Clustering**

### ***kernel K-Means***



# Outline of Video

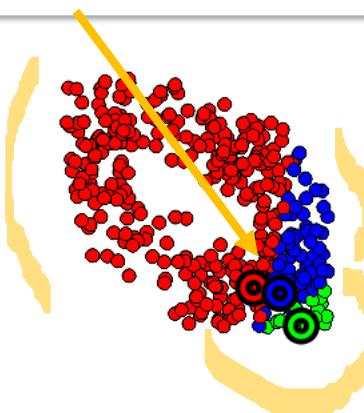
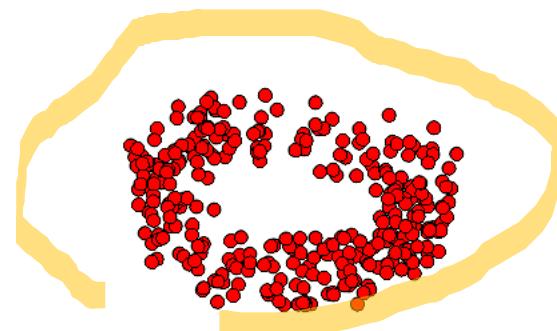
1. Review principle and steps of K-Means algorithm
2. Derive kernel version of K-means



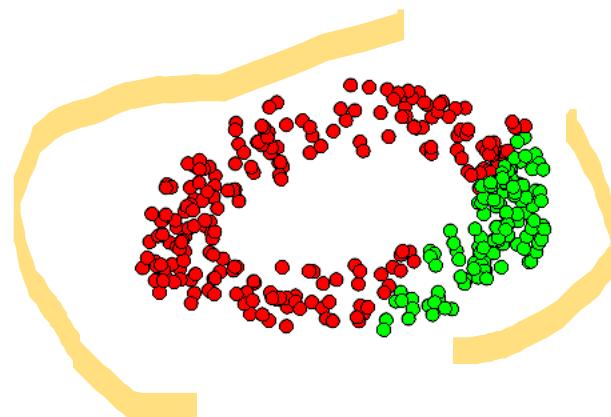
# K-means Clustering: Algorithm

Step 1: Make a guess on what is the correct number of clusters K

Step 2: Initialize randomly the center of the clusters & assign points to closest clusters.

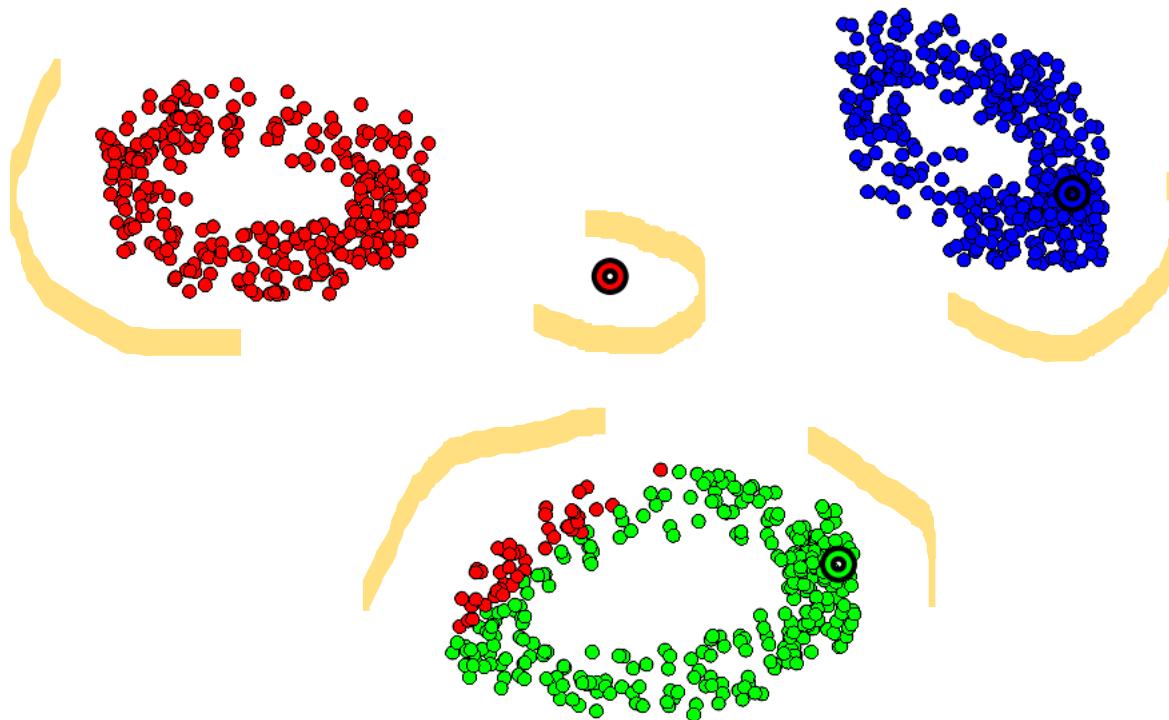


$K=3$

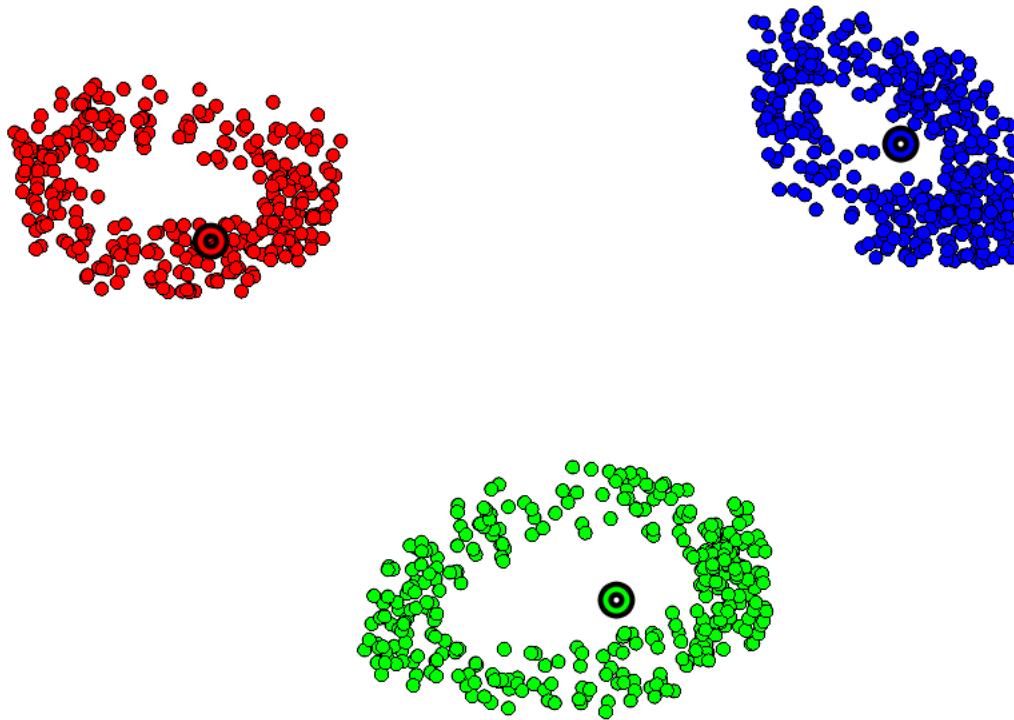


# K-means Clustering: Algorithm

Step 3: recompute the location of the centers of the clusters by taking the centroid of the datapoints assigned to each center.



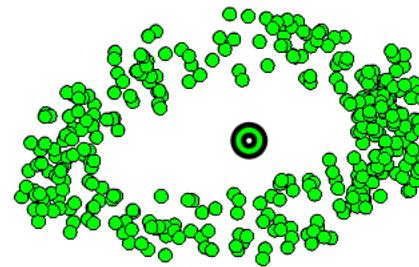
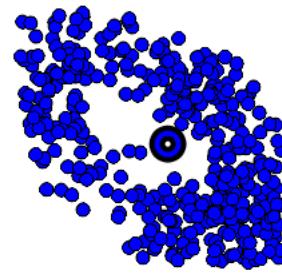
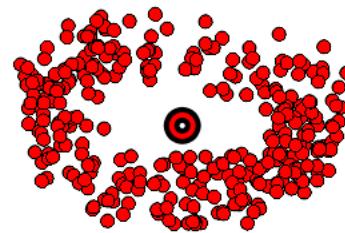
# K-means Clustering: Algorithm



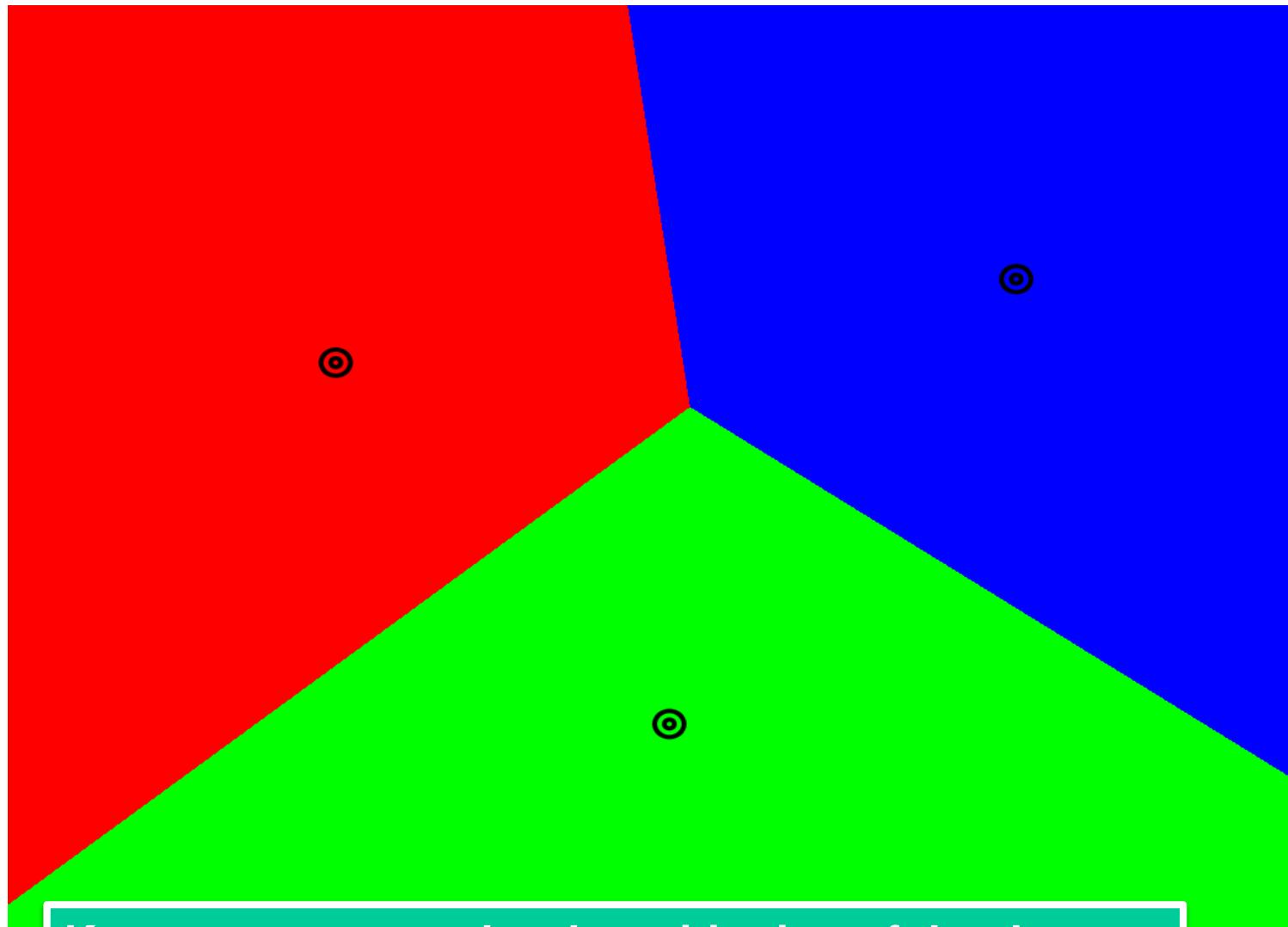
# K-means Clustering: Algorithm

Iterate until convergence

K-means is guaranteed to converge in a finite number of steps.



# K-means Clustering: Algorithm



**K-means creates a hard partitioning of the dataset**



# K-means Clustering: Objective function

K-Means clustering minimizes a quadratic cost function

$$J(\mu^1, \dots, \mu^K) = \sum_{k=1}^K \sum_{x^j \in C^k} \sqrt{(x - \mu^k)^T (x - \mu^k)}$$

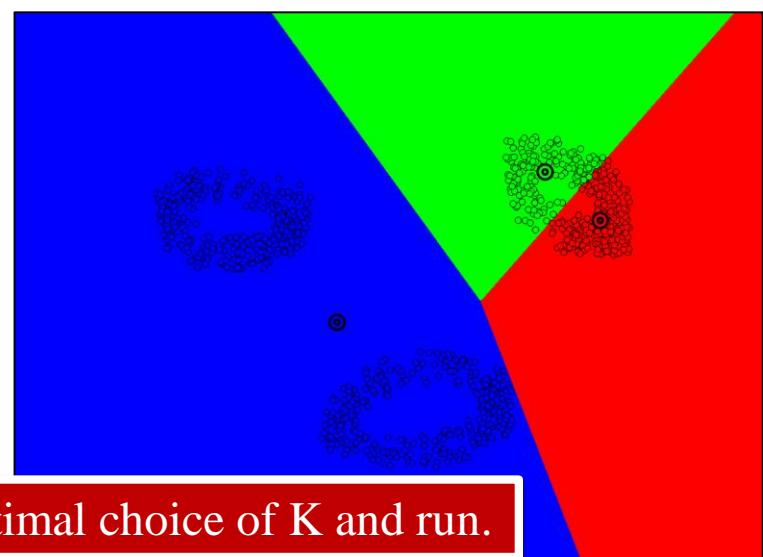
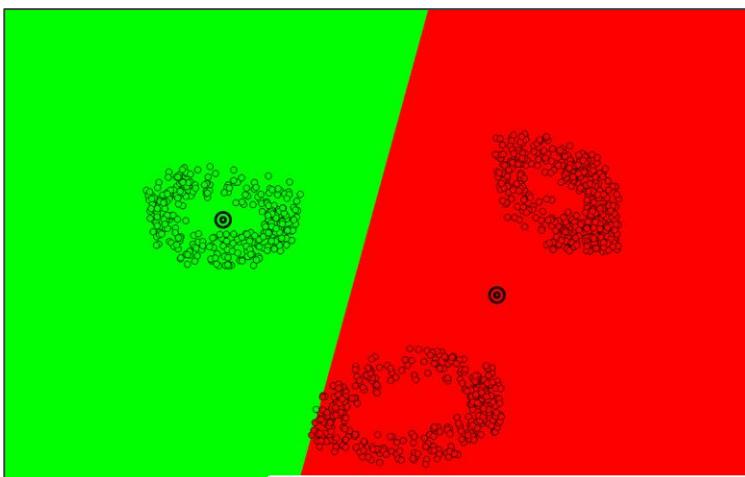
$\mu^1, \dots, \mu^K$  : centers of the  $K$  clusters



## K-means Clustering: Advantages

- The algorithm is guaranteed to converge in a finite number of iterations (but it converges to a local optimum!)
- It is computationally **cheap** and **faster** than other clustering techniques - update step is  $\sim O(N)$ .

## K-means Clustering: Disadvantages



Use RSS or BIC to determine optimal choice of K and run.

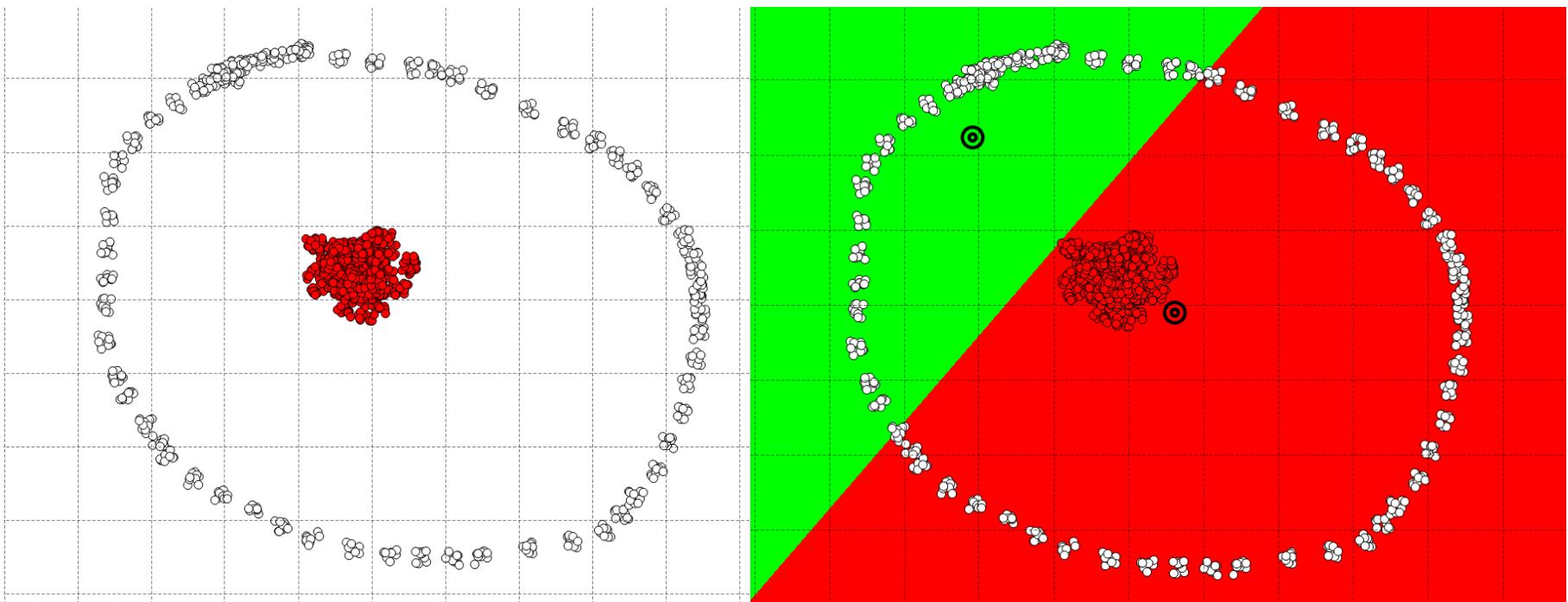
Good performance depends on properly choosing K.

Good performance depends on initialization.



# K-means Clustering: Limitations

K-means can separate clusters linearly, or quasi-linearly (with norm-p) only.



Kernel K-means can generate **non-linear separations** of groups of points.



# Kernel K-means: Principle

## Idea:

- Exploit the principle of the kernel to perform classical K-means clustering with norm-2 in feature space:
  - This yields non-linear boundaries.
  - This retains simplicity of computation of linear K-means.
- The objective function of K-means is composed of an inner product across datapoints.
  - One can replace the inner product with a kernel to perform inner product in feature space.



# Kernel K-means: Derivation

K – Means algorithm minimizes the objective function :

$$J(\mu^1, \dots, \mu^K) = \sum_{k=1}^K \sum_{x^j \in C^k} \|x^j - \mu^k\|^2 \quad \text{with } \mu^k = \frac{\sum_{x^j \in C^k} x^j}{m_k}$$

$m_k$  : number of datapoints in cluster  $C^k$

Project the points into feature space

$$\{x^i\}_{i=1}^M \xrightarrow{\text{red arrow}} \{\phi(x^i)\}_{i=1}^M$$

Express the same problem in feature space:

$$J(\mu^1, \dots, \mu^K) = \sum_{k=1}^K \sum_{x^j \in C^k} \|\phi(x^j) - \phi(\mu^k)\|^2$$

$$\frac{\sum_{x^j \in C^k} \phi(x^j)}{m_k}$$

We cannot observe the image of  $\mu^k$  in feature space.

→ Construct  $\phi(\mu^k)$  using images of points in same cluster.



# Kernel K-means: Solution

$$\begin{aligned}
 J(\mu^1, \dots, \mu^K) &= \sum_{k=1}^K \sum_{x^j \in C^k} \left\| \phi(x^j) - \frac{\sum_{x^l \in C^k} \phi(x^l)}{m_k} \right\|^2 \\
 &\quad k(x^l, x^j) \quad k(x^j, x^l) \\
 &= \sum_{k=1}^K \sum_{x^j \in C^k} \left( \phi(x^j)^T \phi(x^j) - \frac{2 \sum_{x^l \in C^k} \phi(x^l)^T \phi(x^j)}{m_k} + \frac{\sum_{x^t, x^l \in C^k} \phi(x^t)^T \phi(x^l)}{(m_k)^2} \right) \\
 &\quad k(x^j, x^j) \\
 &= \sum_{k=1}^K \sum_{x^j \in C^k} \left( k(x^j, x^j) - \frac{2 \sum_{x^t \in C^k} k(x^i, x^t)}{m_k} + \frac{\sum_{x^t, x^l \in C^k} k(x^t, x^l)}{(m_k)^2} \right)
 \end{aligned}$$

Objective function in feature space



# Kernel K-means: Algorithm

Kernel K-means algorithm is also an iterative procedure:

- 1. Initialization:** pick K clusters (random assignment of points to a cluster, or use K-means at initialization)
- 2. Assignment Step:** Assign each data point to its “closest” centroid (E-step).

$$\arg \min_k d(x, C^k) = \min_k \left( k(x, x) - \frac{2 \sum_{x^j \in C^k} k(x, x^j)}{m_k} + \frac{\sum_{x^j, x^l \in C^k} k(x^j, x^l)}{(m_k)^2} \right)$$

- 3. Update Step:** Update the list of points belonging to each centroid (M-step)
4. Go back to step 2 and repeat the process until the clusters are stable.



# Kernel K-means: interpreting the solution

Consider a RBF kernel. Let us interpret the different terms.

Cst of value 1

$$\arg \min_k d(x, C^k) = \min_k \left[ k(x, x) - \frac{2 \sum_{x^j \in C^k} k(x, x^j)}{\sum_{x^j, x^l \in C^k} k(x^j, x^l)} + \frac{(m_k)^2}{\sum_{x^j, x^l \in C^k} k(x^j, x^l)} \right]$$

If  $x$  is close to all points in cluster  $k$ , this sum is close to 2.

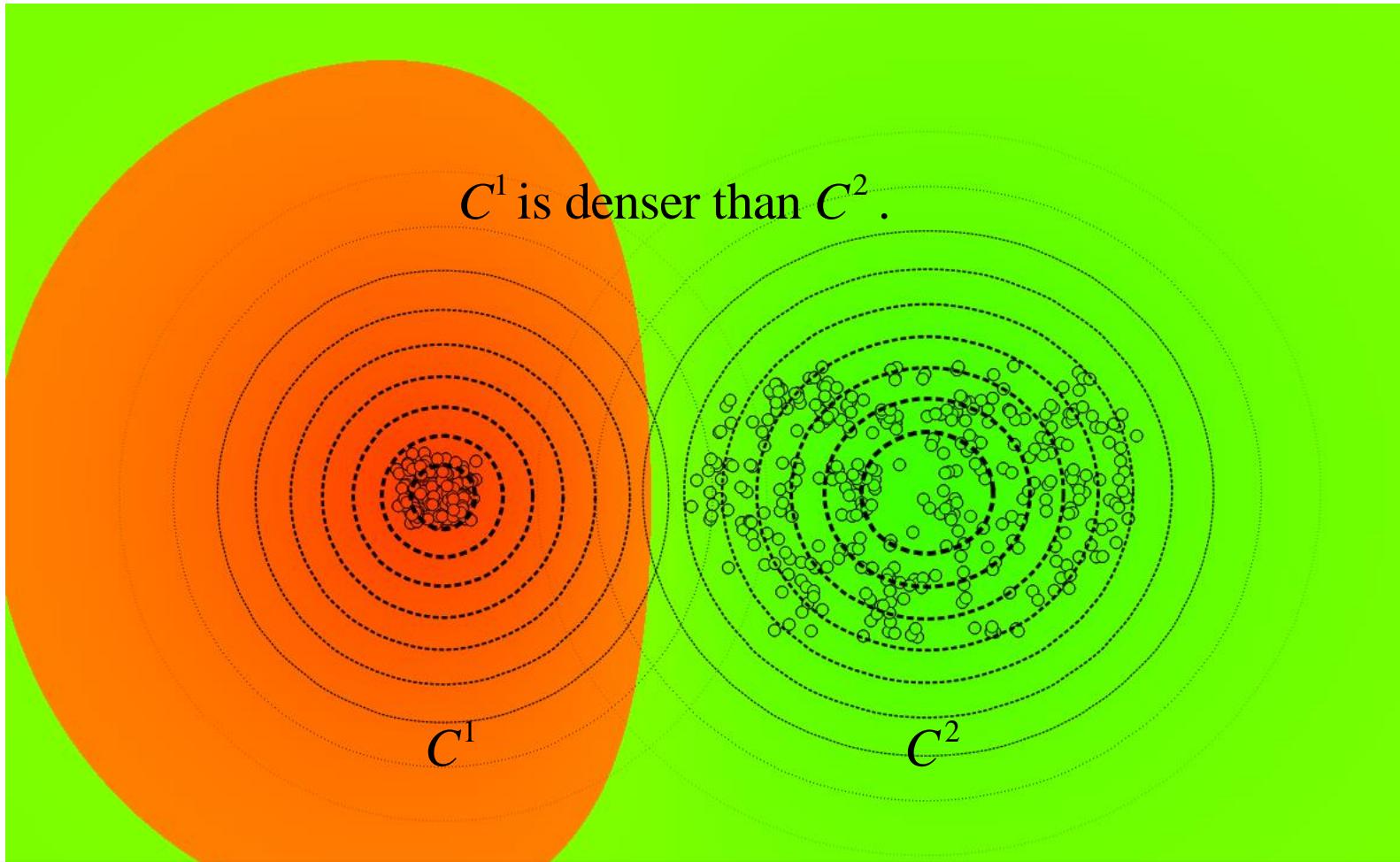
Normalization factor

If the points are tightly grouped in the cluster, this sum is close to 1.

The tighter the cluster, the closer the datapoint must be to the cluster to be assigned.



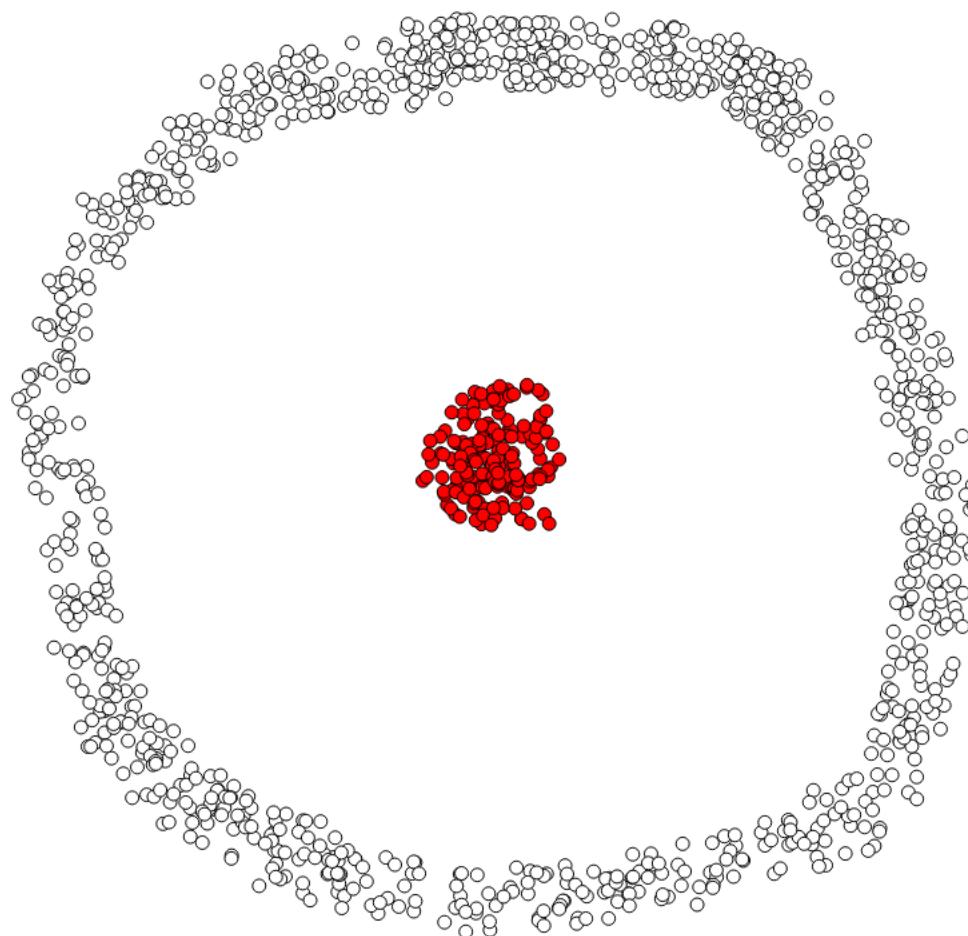
# Kernel K-means: interpreting the solution



The tighter the cluster, the closer the datapoint must be to the cluster to be assigned.



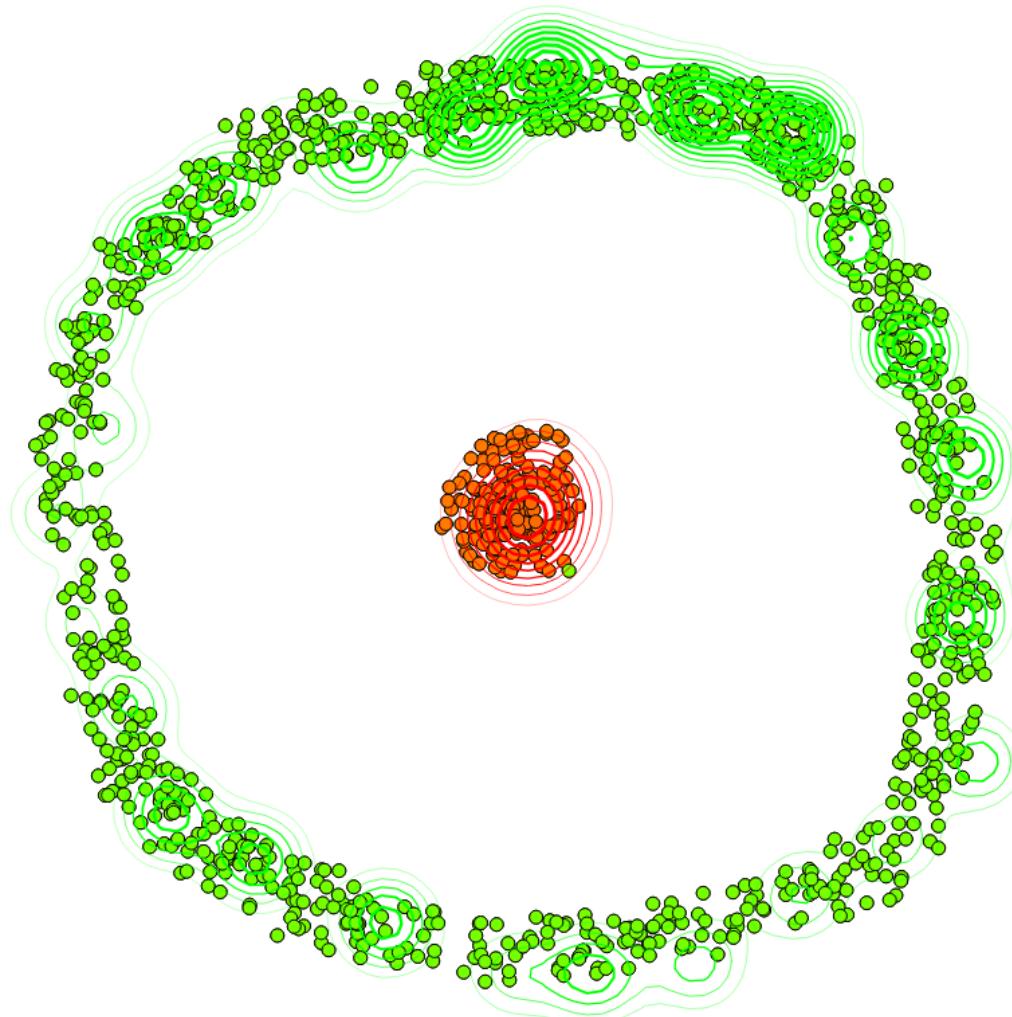
# Nonlinear boundaries with kernel K-means



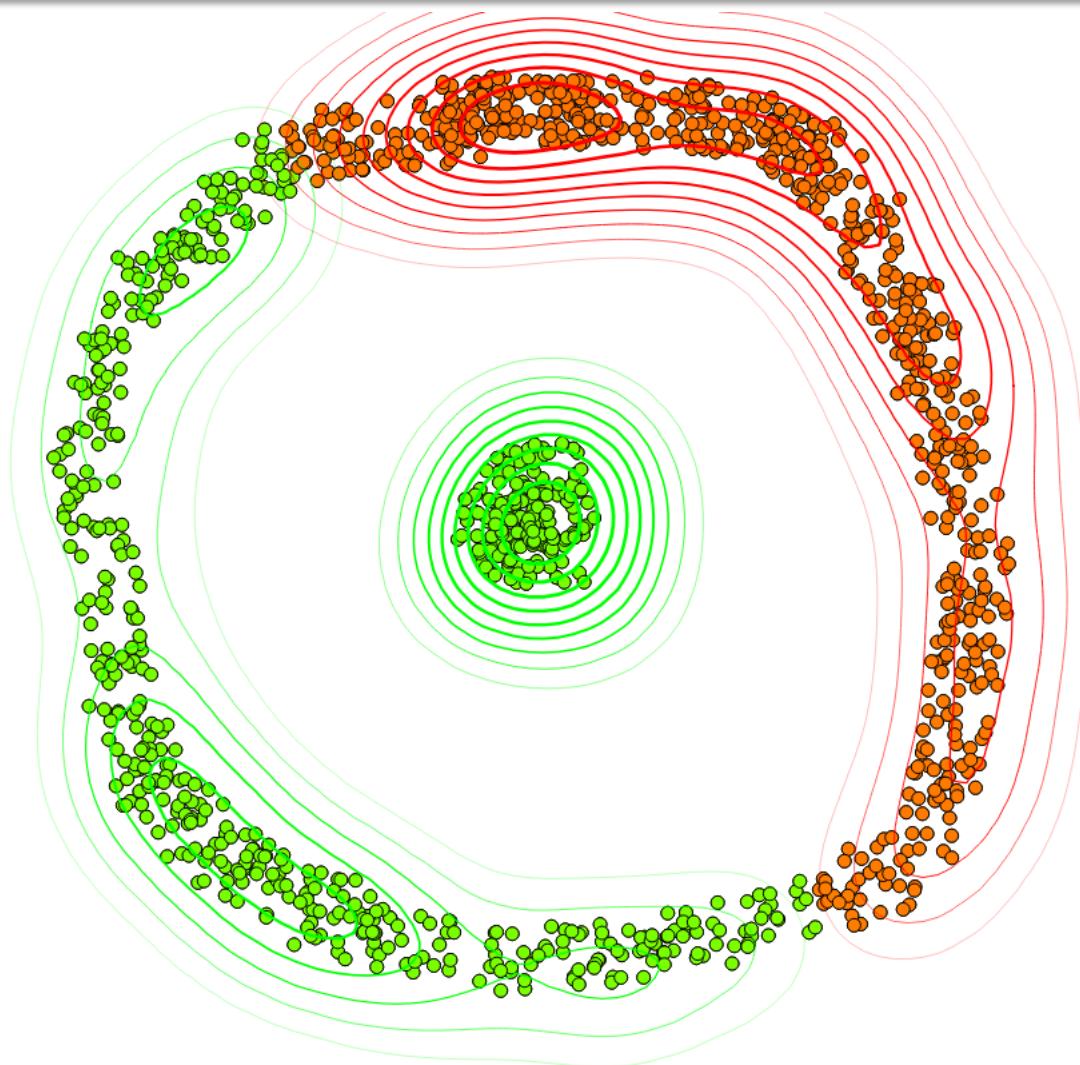
2 Clusters



# Nonlinear boundaries with kernel K-means



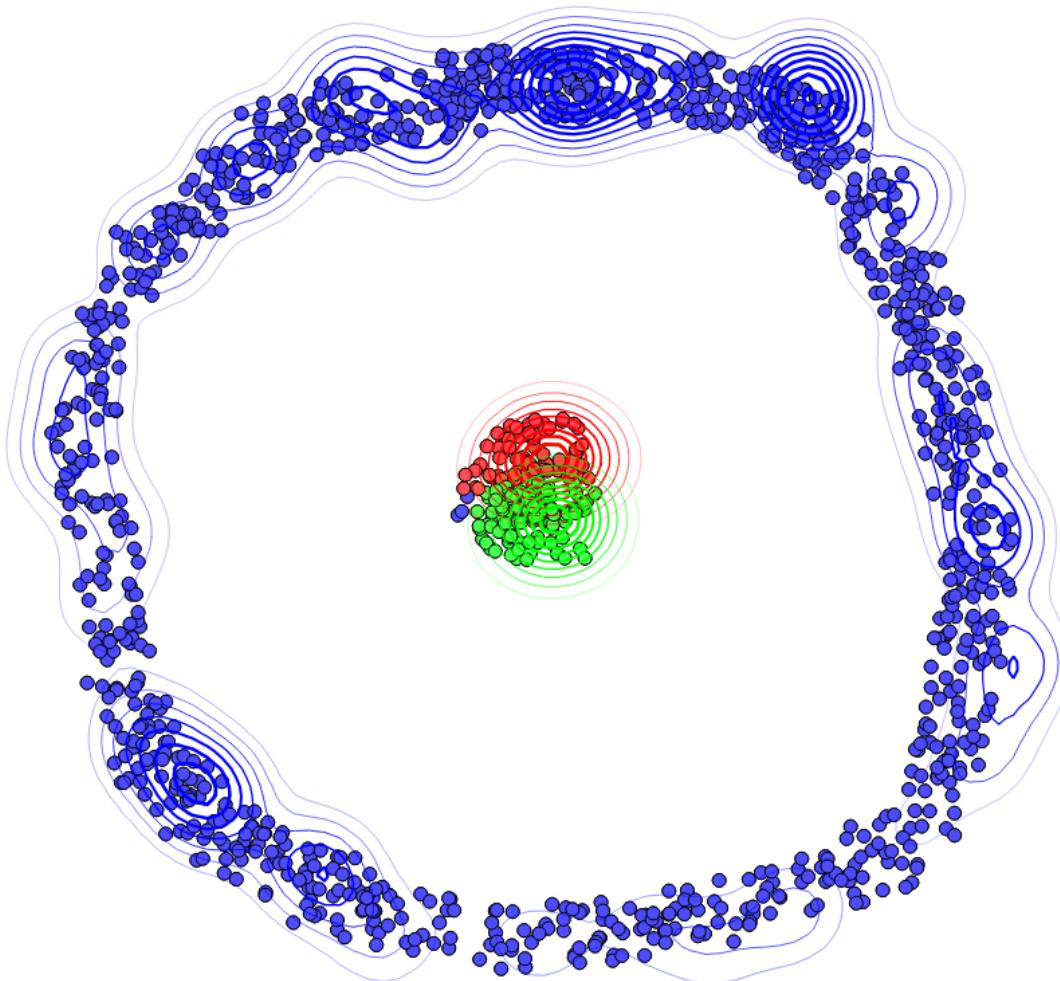
# Sensitivity to kernel width



Kernel width too large



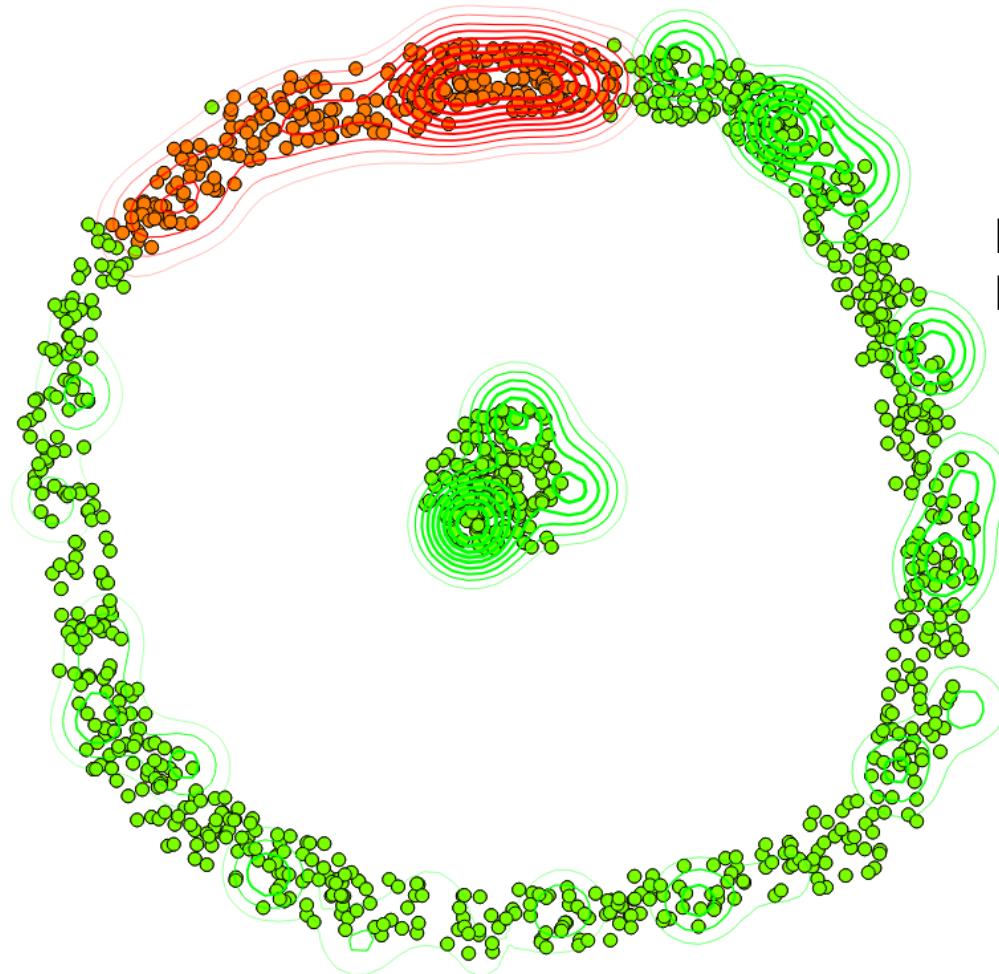
# Sensitivity to choice of K



Still sensitive to choice of K. Here K=3.



# Sensitivity to initialization

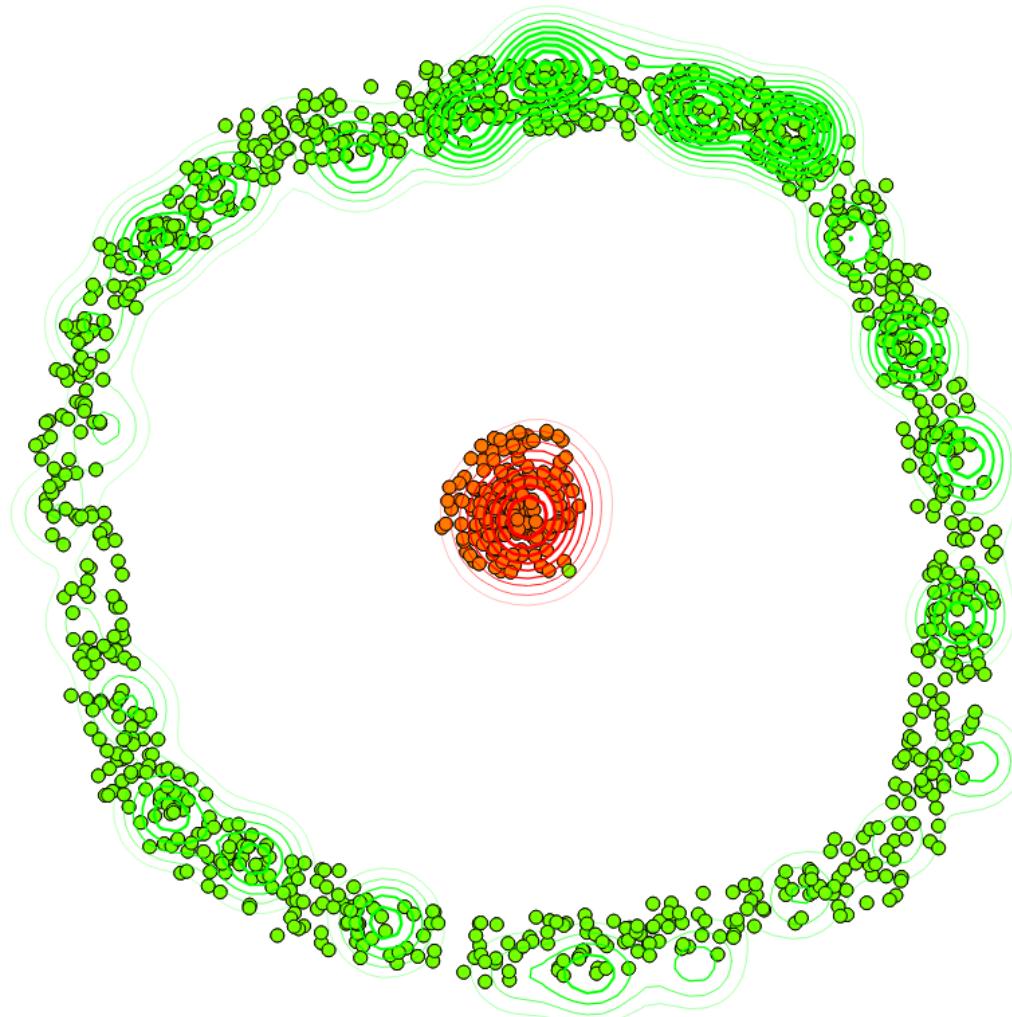


RSS measure: 2773  
BIC: -5500

But still sensitive to initialization



# Finding the right kernel width



**RSS measure: 2625  
BIC: -5522**

Better values on  
both RSS and BIC

With a correct Kernel width



# Kernel K-means: interpreting the solution

With a polynomial kernel

A: Affected by the position of the points from the origin (norm).

B: Affected by the relative angle across the points.

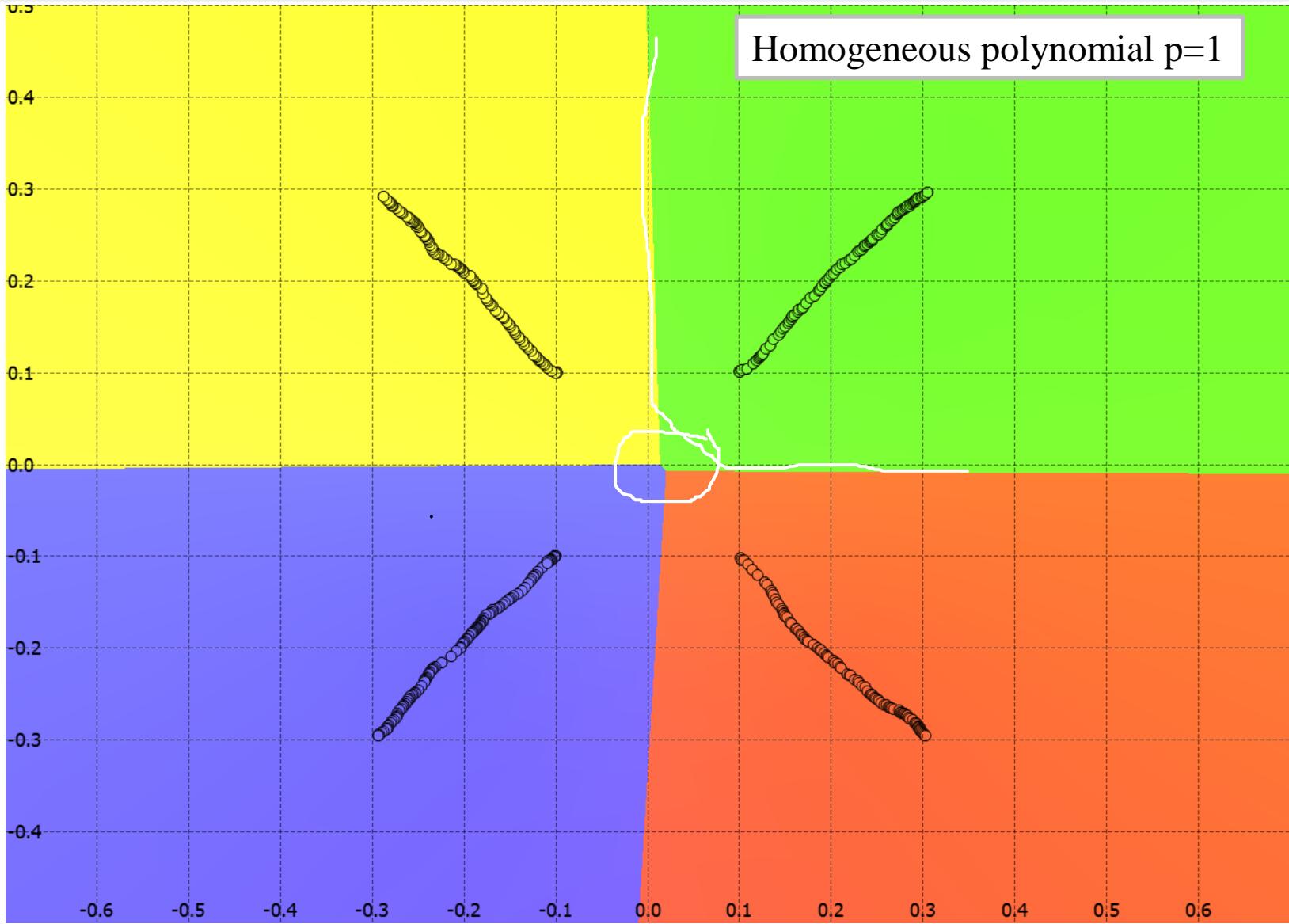
Norm - Positive value

$$\arg \min_k d(x, C^k) = \min_k \left( k(x, x) - \frac{2 \sum_{x^j \in C^k} k(x, x^j)}{m_k} + \frac{\sum_{x^j, x^l \in C^k} k(x^j, x^l)}{(m_k)^2} \right)$$

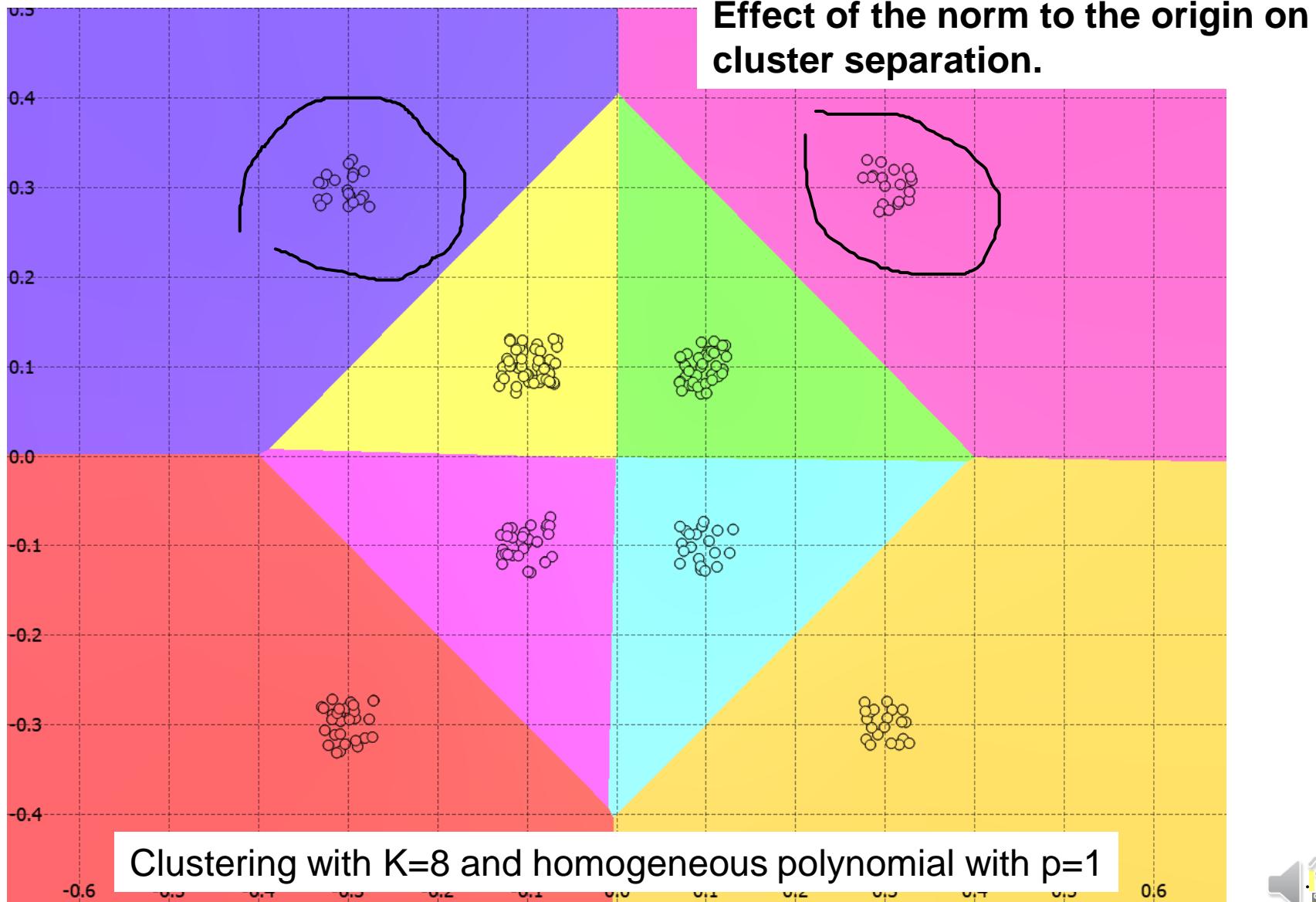
A datapoint will be assigned to the closest cluster in the closest partition.



# Quadran partitioning



# Quadran partitioning



# Summary

- ❑ Kernel K-means follows the same principle as K-means. It is an iterative procedure, akin to Expectation-Maximization.
- ❑ It allows to build highly non-linear boundaries when using the RBF kernel.
- ❑ When using the polynomial kernel, it allows to separate groups of datapoints exploiting the geometrical distribution of the points.
- ❑ As K-means, it depends on initialization of the centers of the clusters which is random.
- ❑ As K-means, the solution depends on choosing well the number of clusters  $K$ . To choose  $K$ , one can use the AIC, BIC or RSS criteria.
- ❑ It is computationally more expensive: K-means is  $O(N)$ . Kernel K-means requires to store a Gram matrix which is  $O(M^2)$ .

