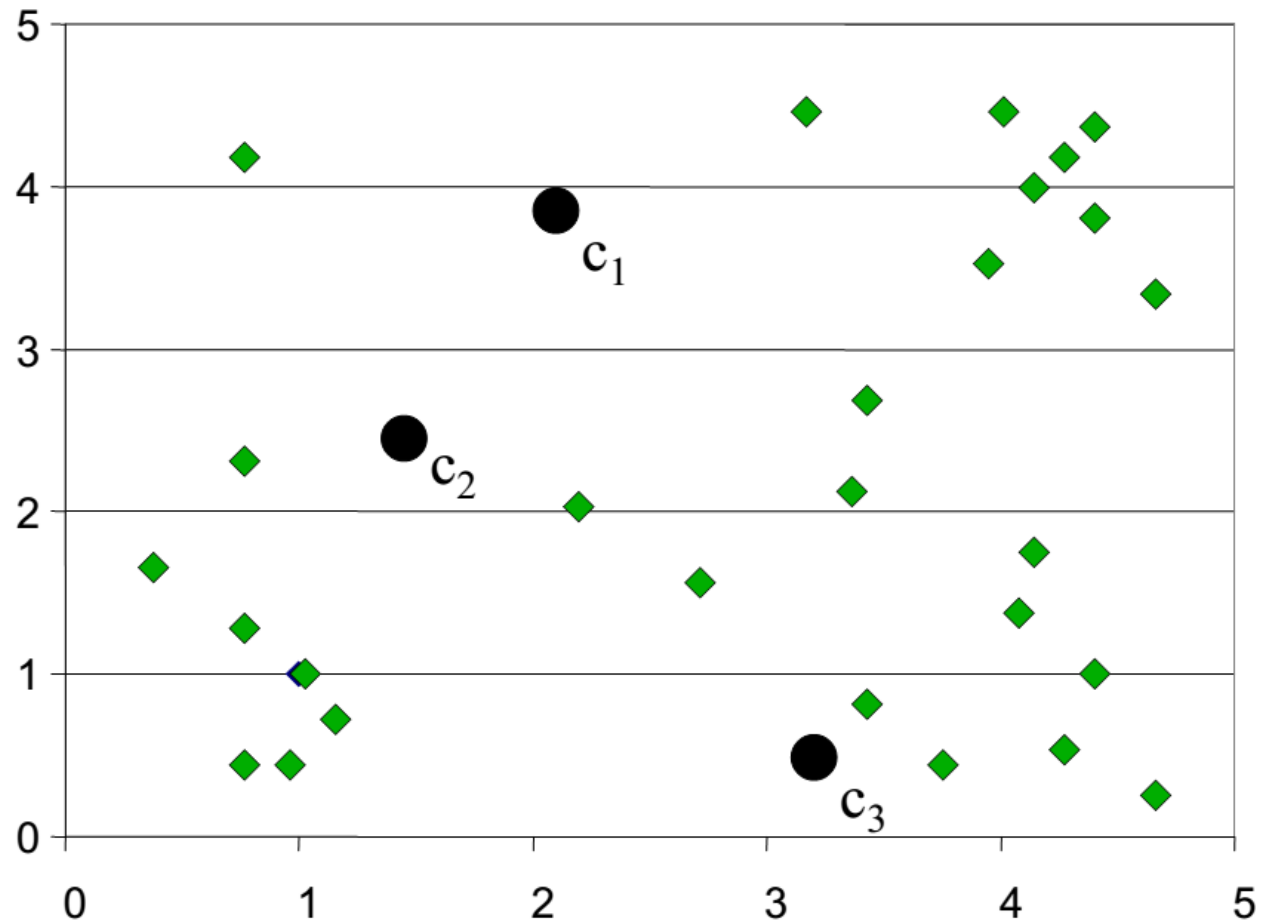


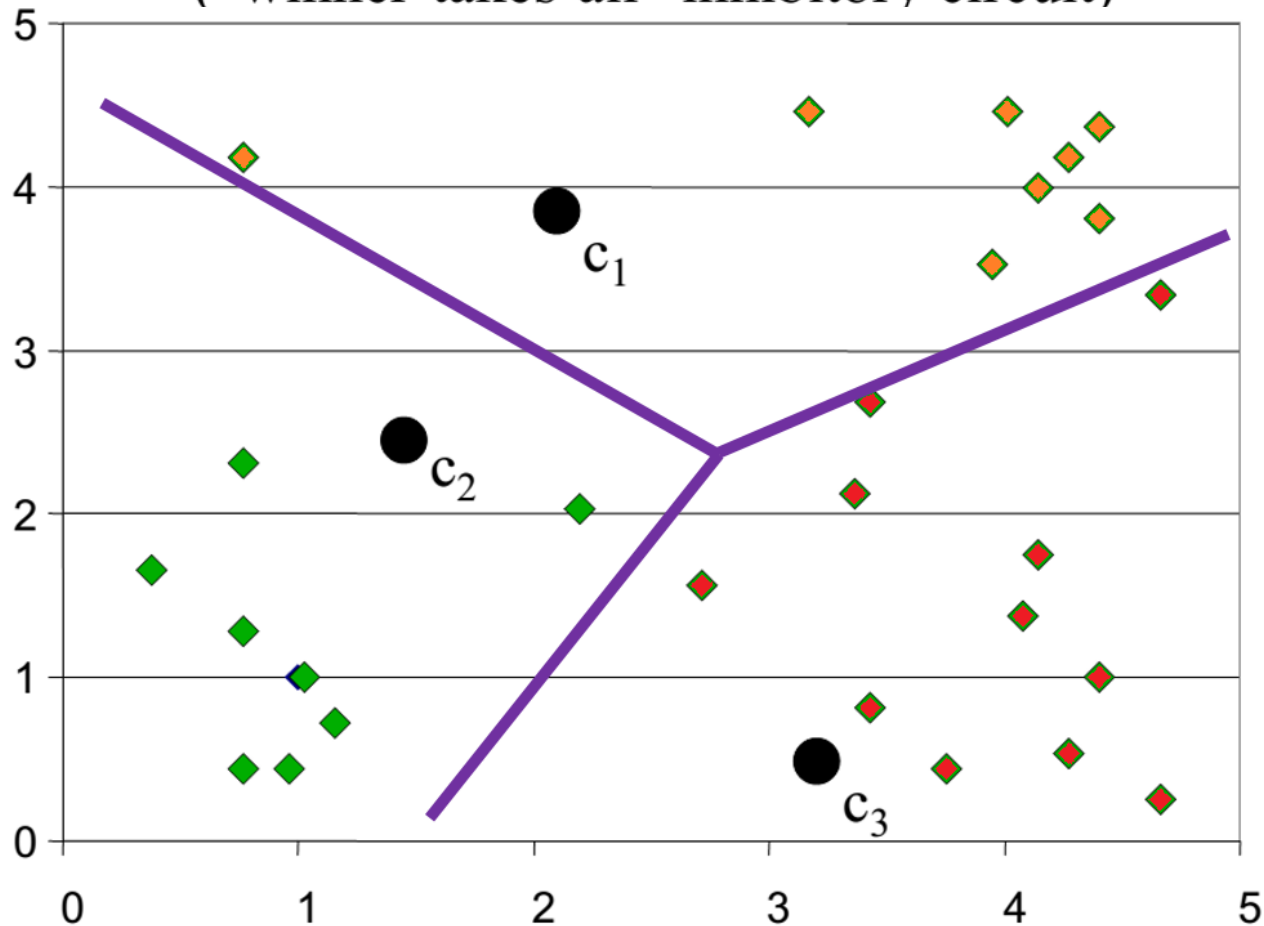
K-means clustering example: step 1

Randomly initialize the cluster centers (synaptic weights)



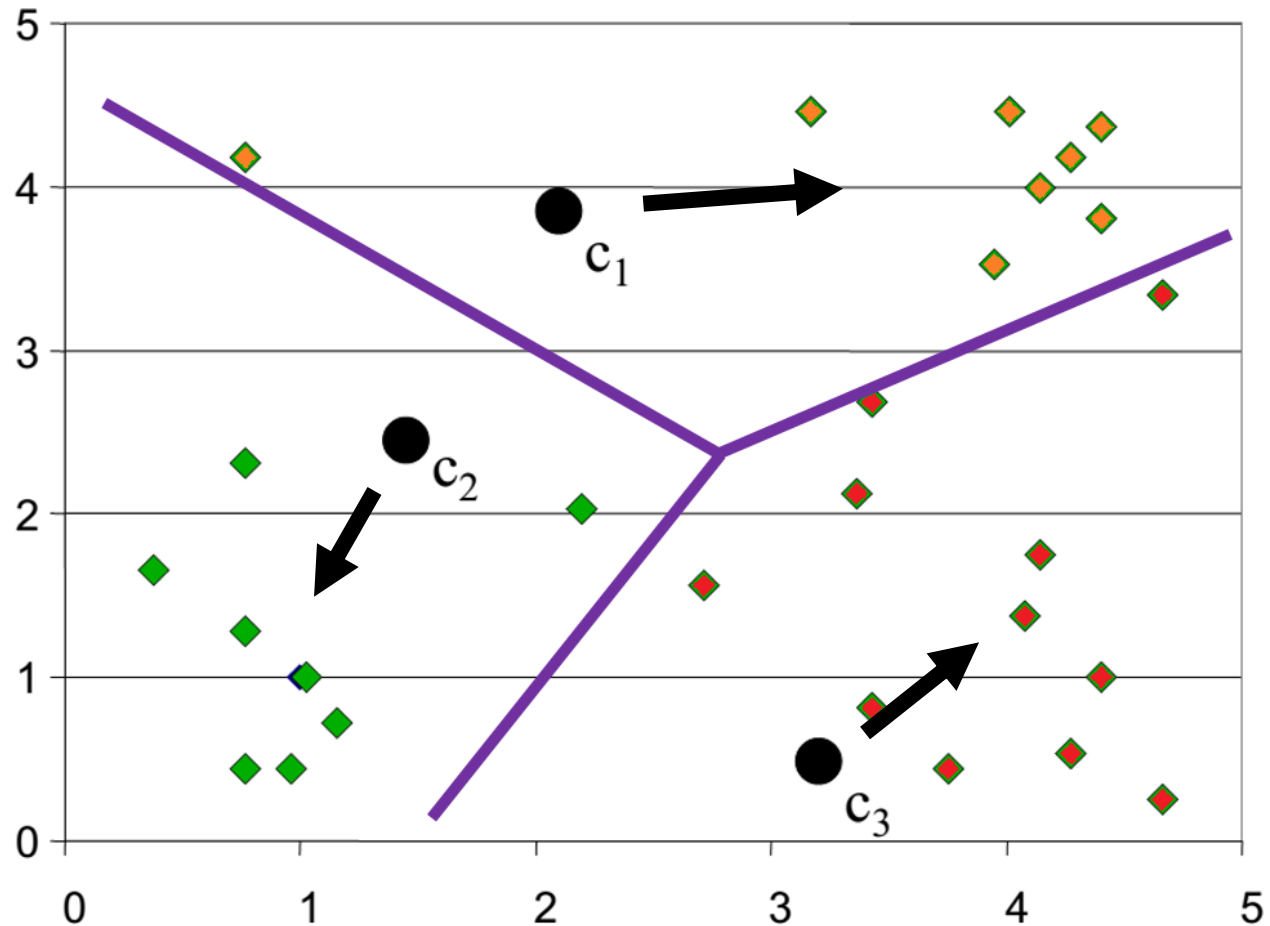
K-means clustering example – step 2

Determine cluster membership for each input
("winner-takes-all" inhibitory circuit)



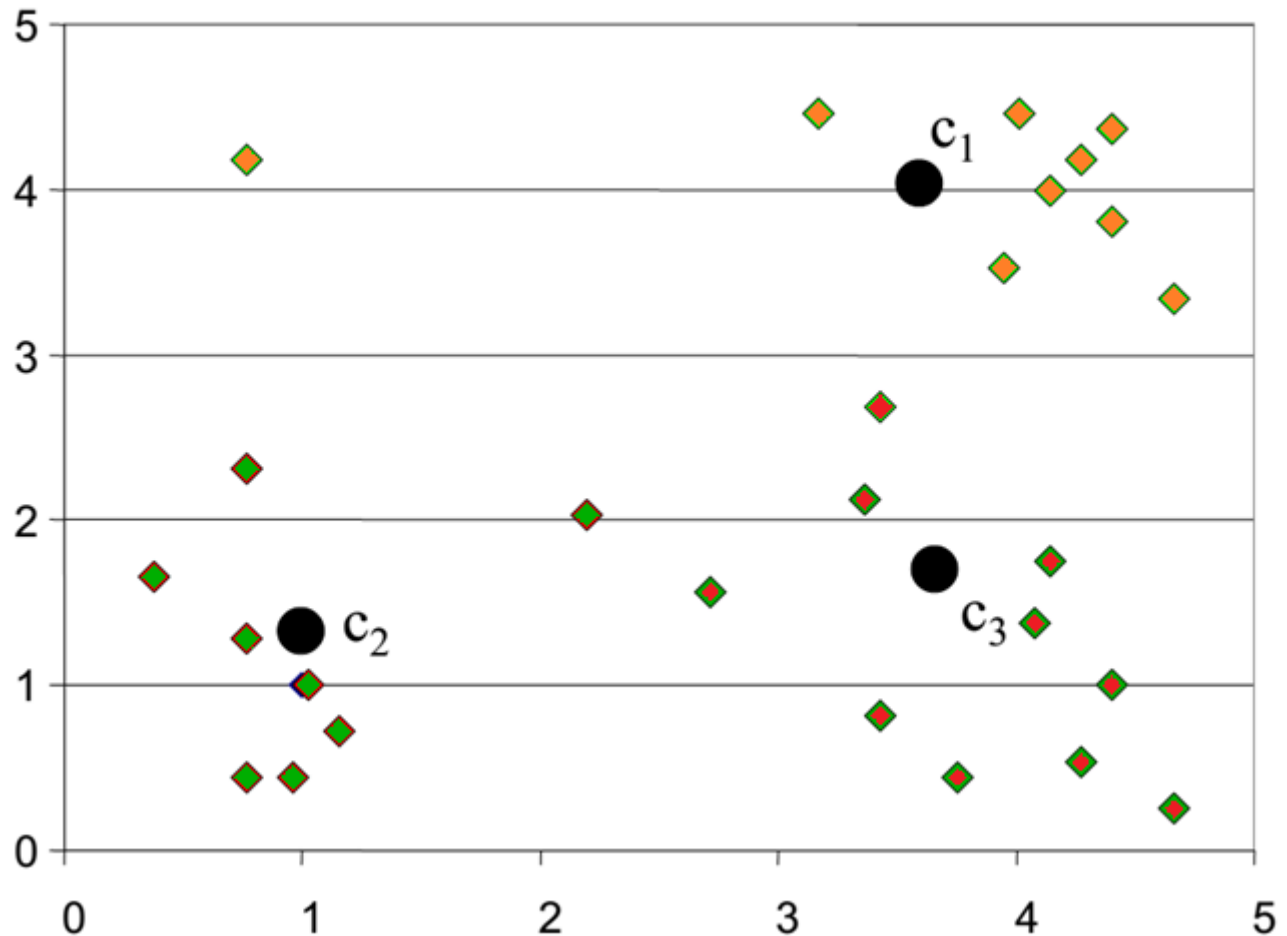
K-means clustering example – step 3

Re-estimate cluster centers (adapt synaptic weights)



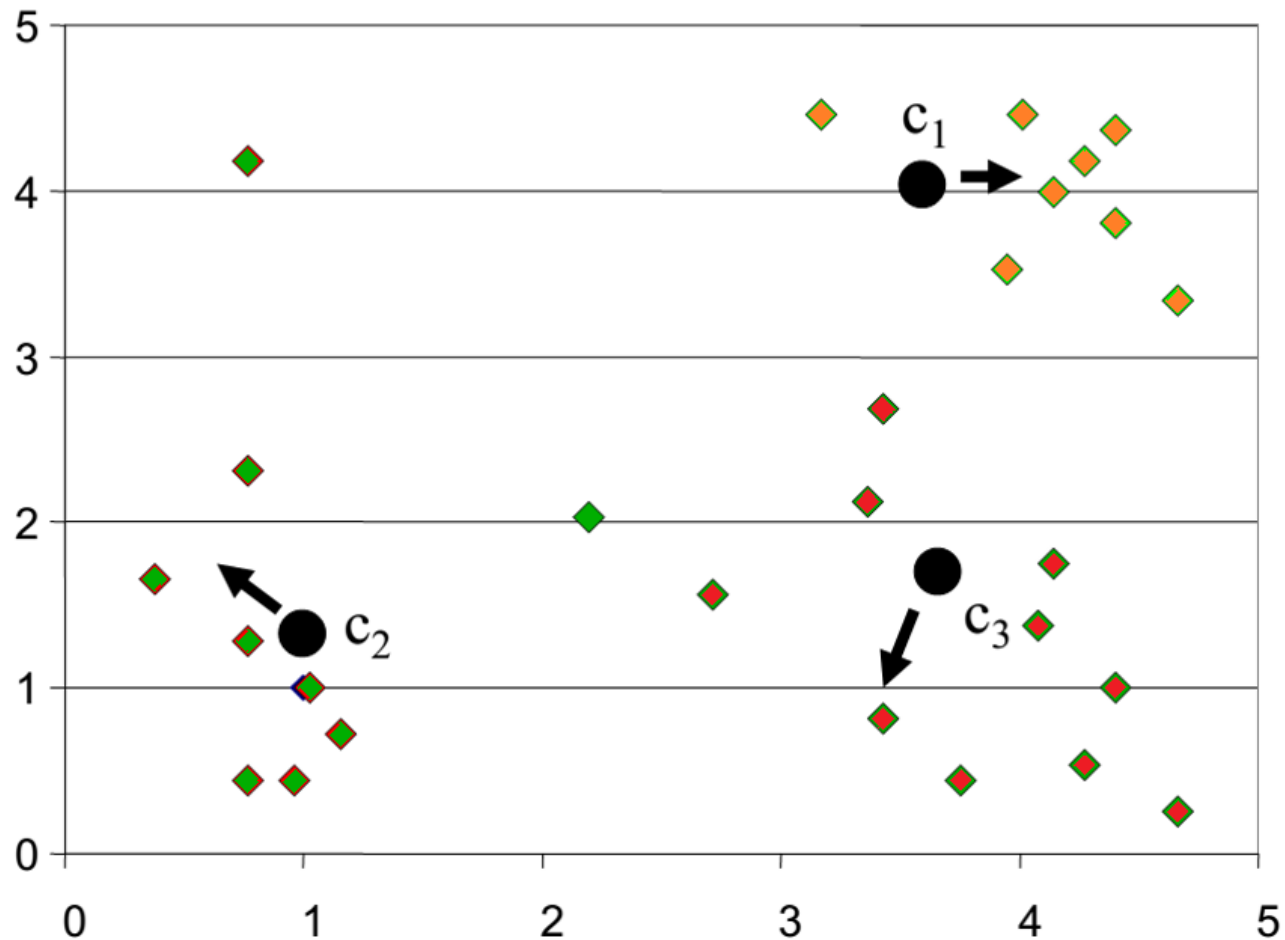
K-means clustering example

Result of first iteration



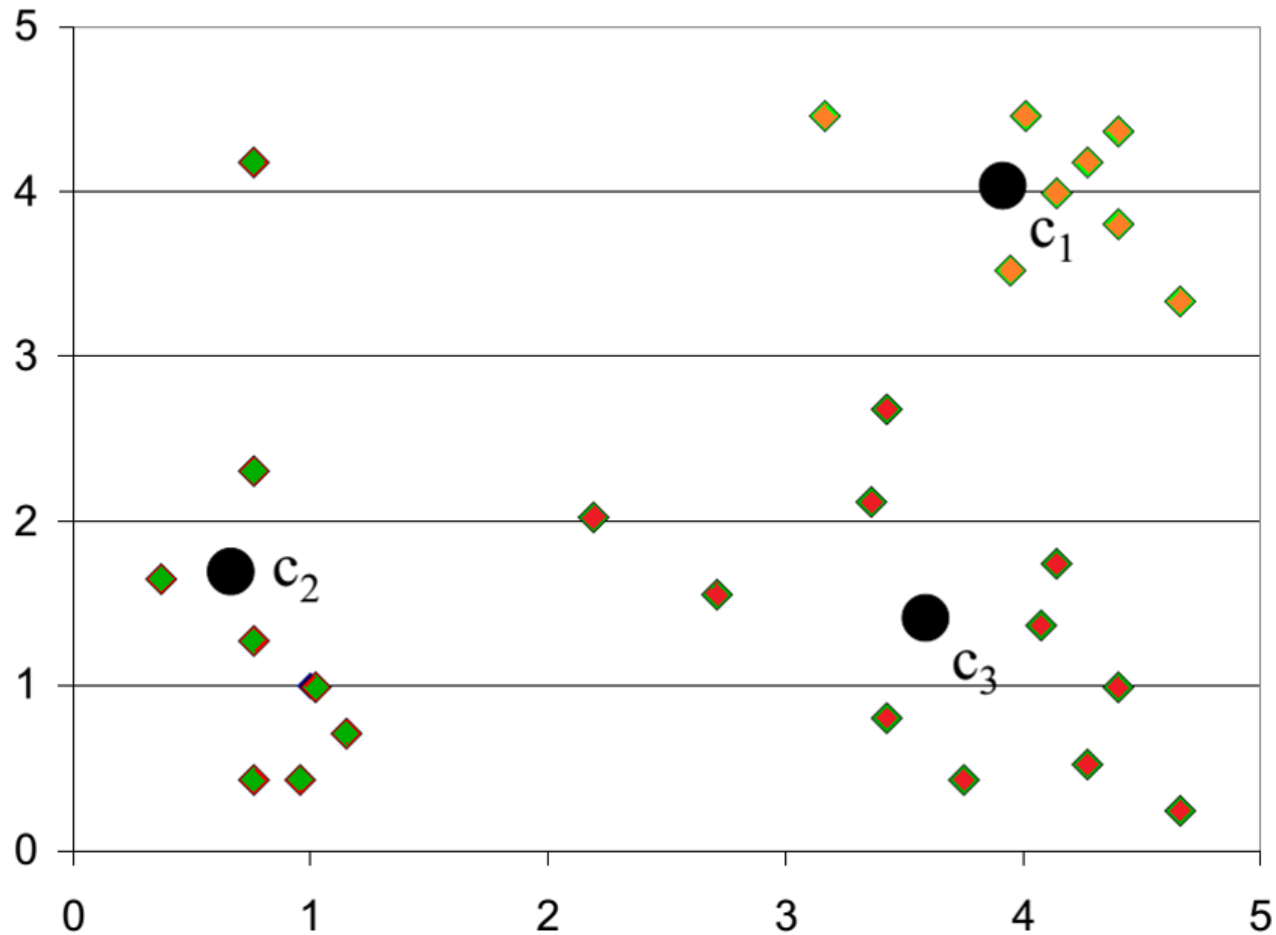
K-means clustering example

Second iteration



K-means clustering example

Result of second iteration



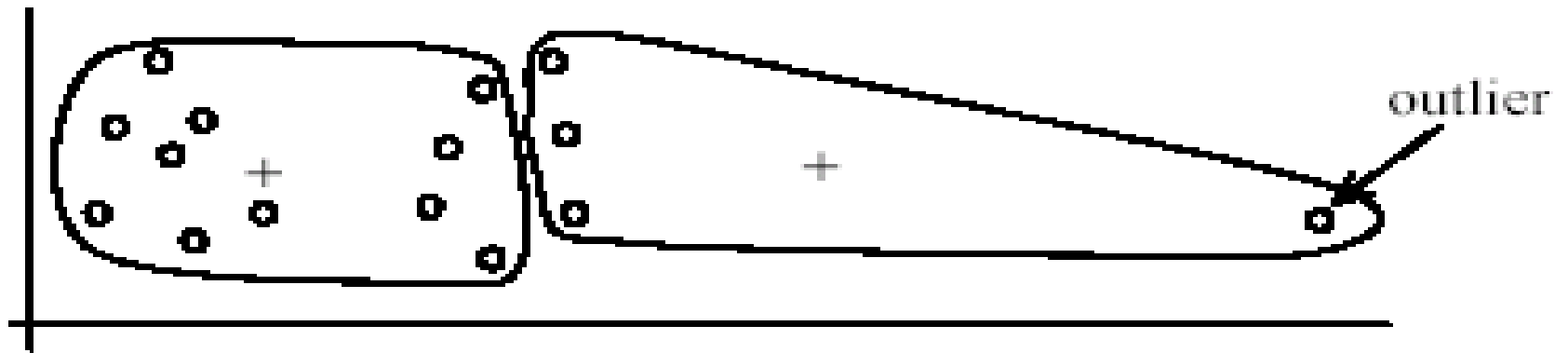
Why use K-means?

- Strengths:
 - Simple: easy to understand and to implement
 - Efficient: Time complexity: $O(tkn)$,
where n is the number of data points,
 k is the number of clusters, and
 t is the number of iterations.
 - Since both k and t are small. k -means is considered a linear algorithm.
- K-means is the most popular clustering algorithm.
- Note that: it terminates at a **local optimum** if SSE is used. The **global optimum** is hard to find due to complexity.

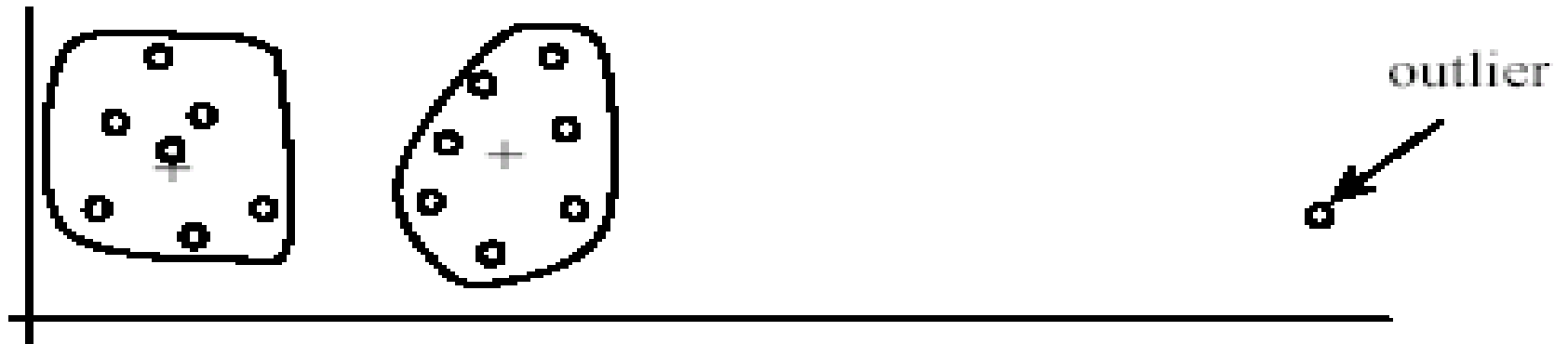
Weaknesses of K-means

- The algorithm is only applicable if the **mean** is defined.
 - For categorical data, *k*-mode - the centroid is represented by most frequent values.
- The user needs to specify ***k***.
- The algorithm is sensitive to **outliers**
 - Outliers are data points that are very far away from other data points.
 - Outliers could be errors in the data recording or some special data points with very different values.

Outliers



(A): Undesirable clusters



(B): Ideal clusters