



# Cluster Analysis

**Dr. Rathachai Chawuthai**

Department of Computer Engineering

Faculty of Engineering

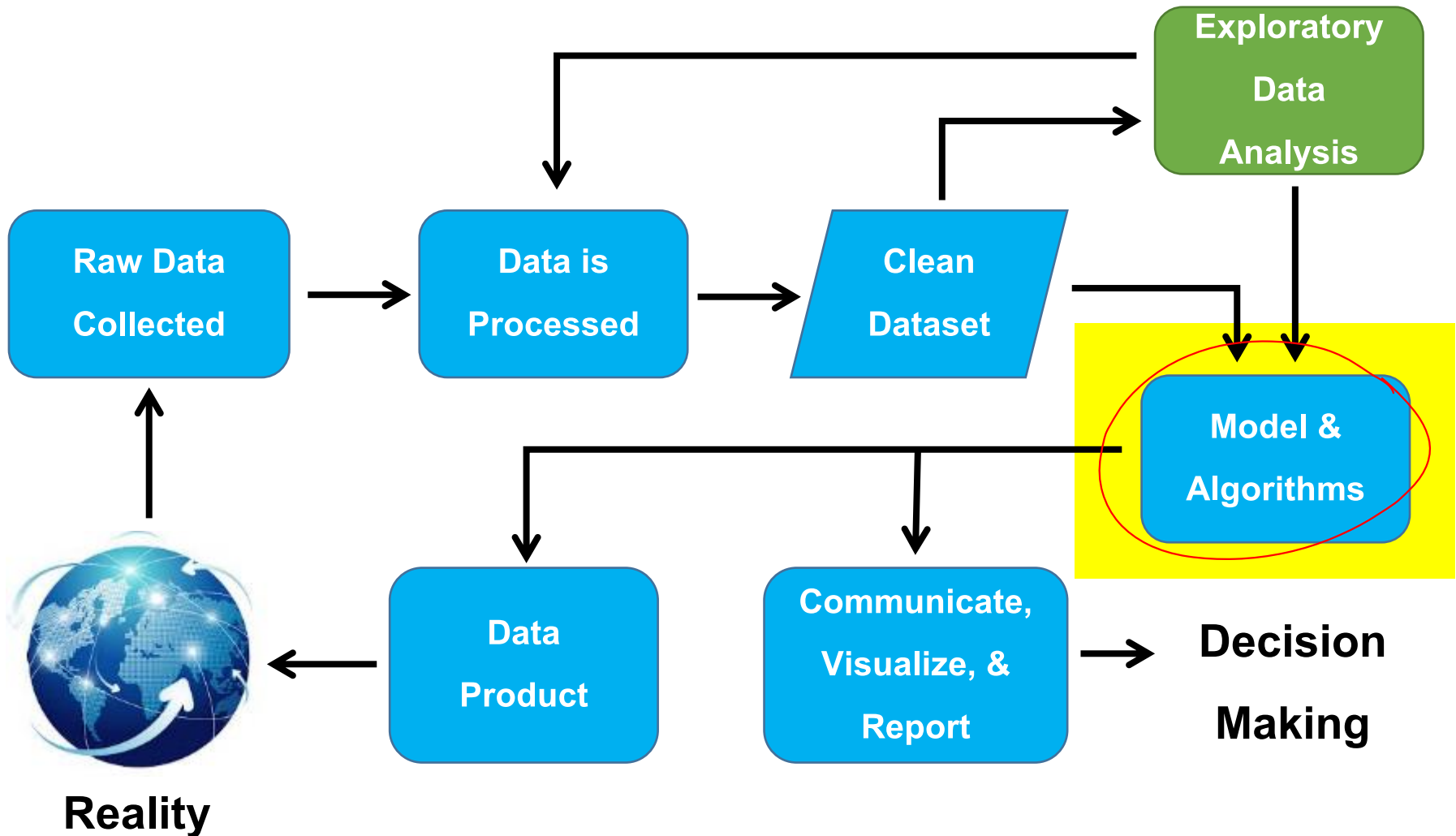
King Mongkut's Institute of Technology Ladkrabang

# Agenda

- Clustering
- K-Means
- DBSCAN
- Cluster Validation
- Dimensionality Reduction

# Data Science Process

---



# Machine Learning

## Supervised Learning

Develop predictive model based on both input and output data

### Regression

- Linear Regression
- Polynomial Regression

### Classification

- Decision Tree
- Logistic Regression
- Neural Network
- etc.

## Unsupervised Learning

Develop predictive model based on both input and output data

### Clustering

- K-Means
- DB-SCAN
- etc.

# Clustering

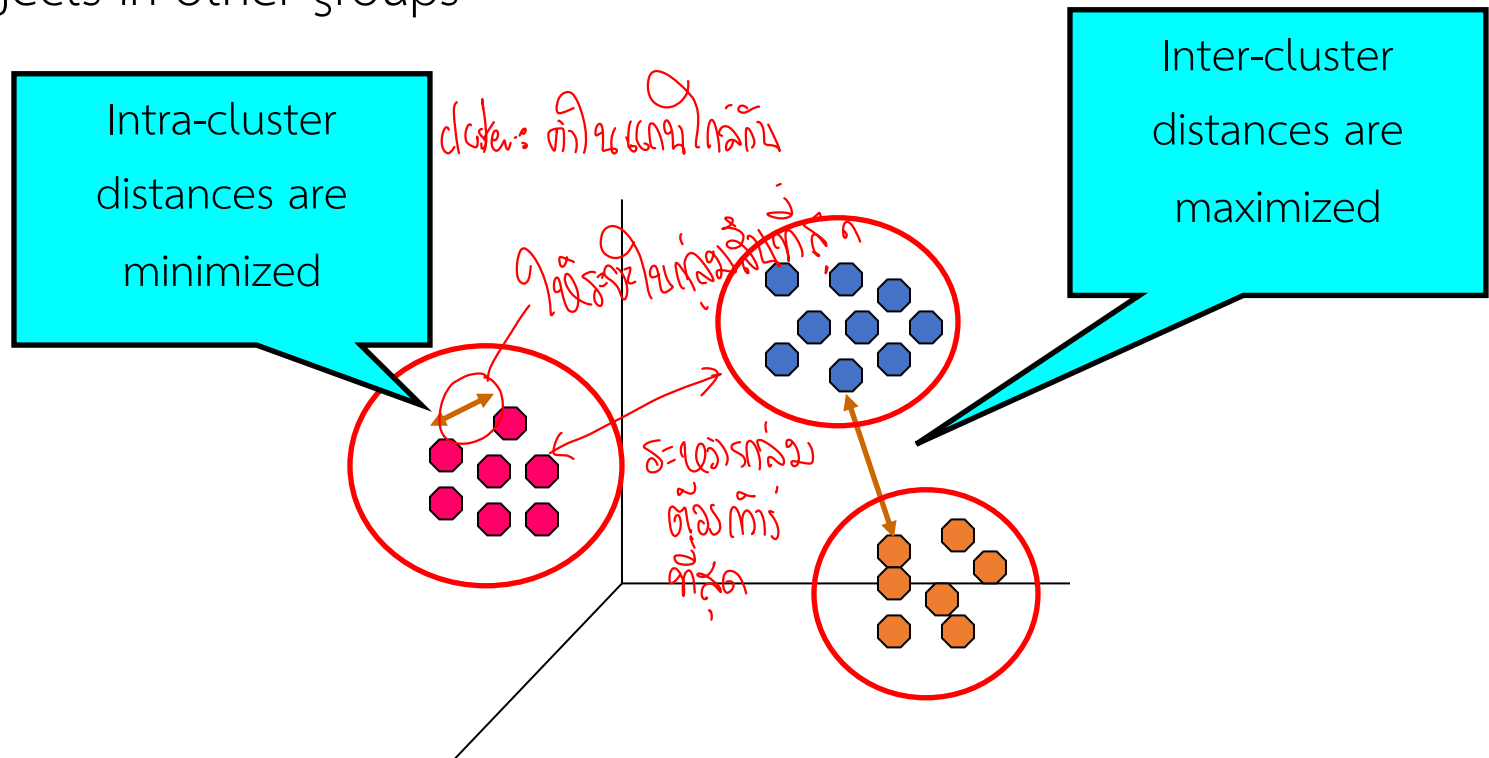
การจัดกลุ่มของข้อมูลที่ไม่สนใจไปข้างหน้า

↓  
อะไรที่เหมือนกันจับมาอยู่ด้วยกัน



# What is a Clustering?

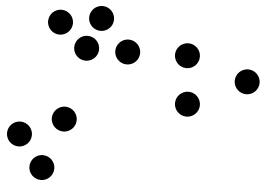
- In general a grouping of objects such that the objects in a group (cluster) are similar (or related) to one another and different from (or unrelated to) the objects in other groups



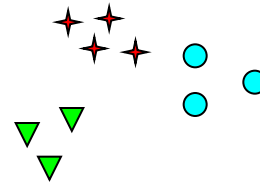
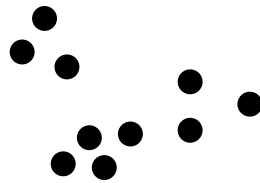
# Notion of a Cluster can be Ambiguous

แล้วจริงๆ กลุ่มของเหตุการณ์นี้กี่กลุ่มล่ะ?

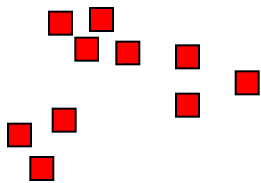
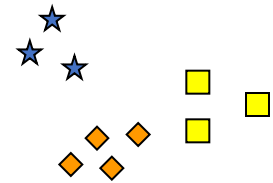
↓  
แล้วแต่เราสนใจ



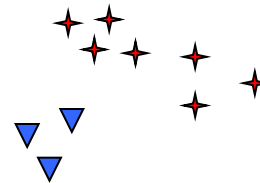
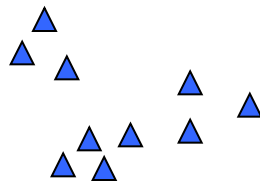
How many clusters?



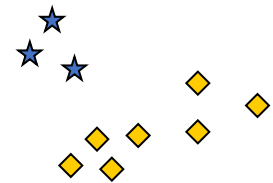
6 Clusters



2 Clusters



4 Clusters

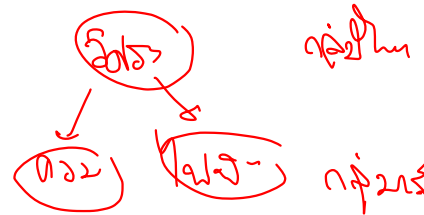


# Types of Clusterings

- A clustering is a set of clusters
- Important distinction between hierarchical and partitional sets of clusters

- ကွဲစုခြင်း  
• **Partitional Clustering** → ဘုံကလပ်ခွဲခြားနိုင်ဖို့ လိုအပ်တယ်  
  - A division data objects into subsets (clusters) such that each data object is in exactly one subset

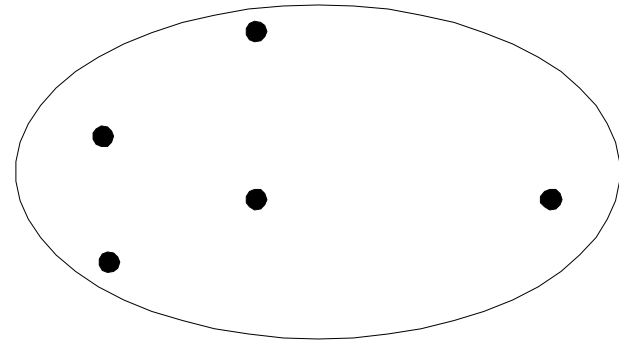
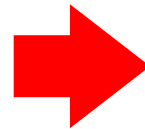
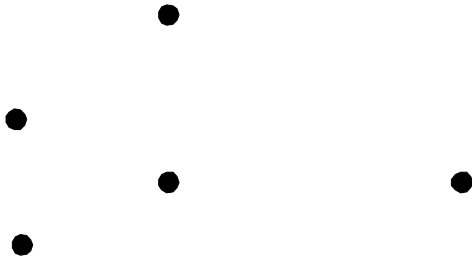
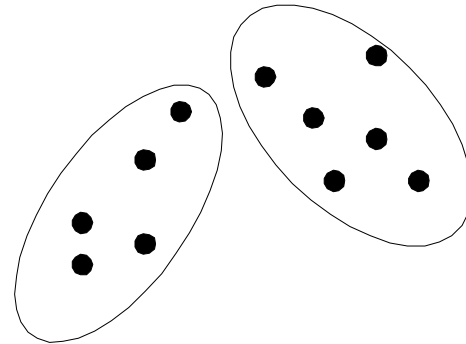
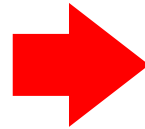
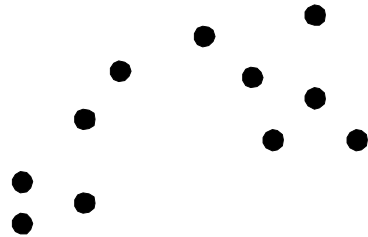
- **Hierarchical clustering** → ချဲ့ချဲ့ဝင်နေခြင်း  
  - A set of nested clusters organized as a hierarchical tree





# Partitional Clustering

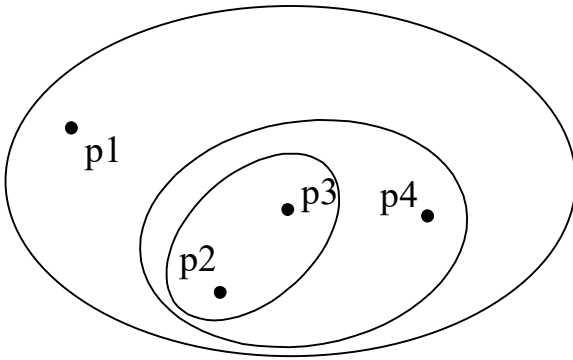
---



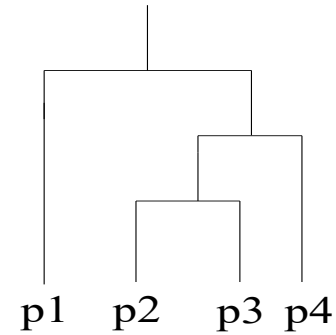
Original Points

A Partitional Clustering

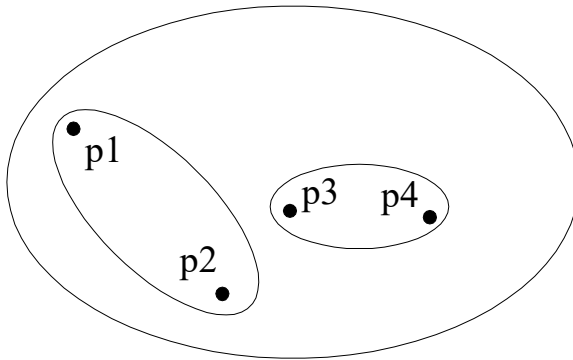
# Hierarchical Clustering



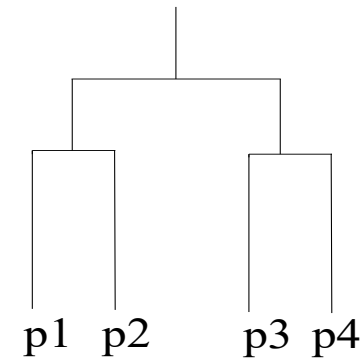
Traditional Hierarchical Clustering



Traditional Dendrogram



Non-traditional Hierarchical Clustering



Non-traditional Dendrogram

# Other types of clustering

---

- **Exclusive** (or non-overlapping) versus **non-exclusive** (or overlapping)
  - In non-exclusive clusterings, points may belong to multiple clusters.
    - Points that belong to multiple classes, or ‘border’ points
- **Fuzzy** (or soft) versus **non-fuzzy** (or hard)
  - In fuzzy clustering, a point belongs to every cluster with some weight between 0 and 1
    - Weights usually must sum to 1 (often interpreted as probabilities)
- **Partial** versus **complete**
  - In some cases, we only want to cluster some of the data

# Types of Clusters: Well-Separated

---

- Well-Separated Clusters: ပြားပြားဝေးဝေး
  - A cluster is a set of points such that any point in a cluster is closer (or more similar) to every other point in the cluster than to any point not in the cluster.



# Types of Clusters: Center-Based

---

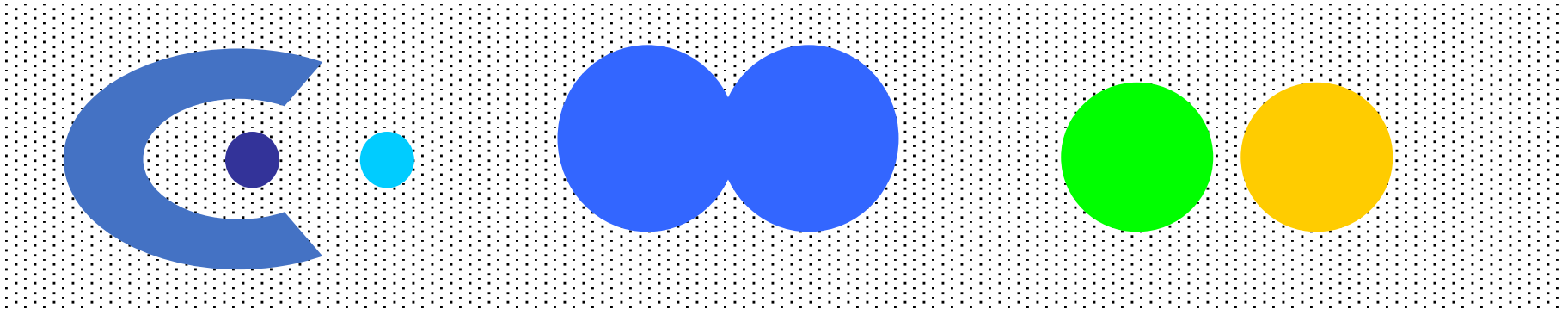
- Center-based
  - A cluster is a set of objects such that an object in a cluster is closer (more similar) to the “center” of a cluster, than to the center of any other cluster
  - The center of a cluster is often a centroid, the minimizer of distances from all the points in the cluster, or a medoid, the most “representative” point of a cluster



4 center-based clusters

# Types of Clusters: Density-Based

- Density-based *จ=รอย/กลุ่มที่แน่นติดกัน*
  - A cluster is a dense region of points, which is separated by low-density regions, from other regions of high density.
  - Used when the clusters are irregular or intertwined, and when noise and outliers are present.



6 density-based clusters

# Clustering Algorithms

---

- K-Means

အားပြုလုပ်ရန်

- DBSCAN

- Hierarchical clustering
- PAM, CLARANS: Solutions for the [k-medoids](#) problem
- BIRCH: Constructs a [hierarchical tree](#) that acts a summary of the data, and then clusters the leaves.
- MST: Clustering using the [Minimum Spanning Tree](#).
- ROCK: clustering [categorical data](#) by neighbor and link analysis
- LIMBO, COOLCAT: Clustering [categorical data](#) using [information theoretic](#) tools.
- CURE: [Hierarchical](#) algorithm uses different representation of the cluster
- CHAMELEON: [Hierarchical](#) algorithm uses [closeness and interconnectivity](#) for merging

# K-Means





# K-means Clustering → បែងចែកទិន្នន័យ (ឬ ទិន្នន័យ) ទៅជា ក្រុម ដែលមាន ចំណុច កណ្តាល ក្នុង ក្រុម ទាំងនោះ

↓ រក ចំណុច កណ្តាល ក្នុង ក្រុម ទាំងនោះ → ចំណុច កណ្តាល ក្នុង ក្រុម ទាំងនោះ

---

- Partitional clustering approach
- Each cluster is associated with a **centroid** (center point)
- Each point is assigned to the cluster with the **closest** centroid
- Number of clusters, **K**, must be specified
- The objective is to **minimize the sum of distances** of the points to their respective **centroid**

# K-means Clustering

---

- **Problem:** Given a set  $X$  of  $n$  points in a  $d$ -dimensional space and an integer  $K$  group the points into  $K$  clusters  $C = \{C_1, C_2, \dots, C_k\}$  such that

$$\text{Cost}(C) = \sum_{i=1}^k \sum_{x \in C_i} \text{dist}(x, c_i)$$

*cost = sum of distances to centroid*

is minimized, where  $c_i$  is the centroid of the points in cluster  $C_i$

# K-means Algorithm

---

- Also known as **Lloyd's algorithm**.
- K-means is sometimes synonymous with this algorithm

1: Select  $K$  points as the initial centroids.

ตัวแรกมาเลือกเป็นจุดไปเรื่อยๆ

2: **repeat**

3: Form  $K$  clusters by assigning all points to the closest centroid. **ขบวนการ**

4: Recompute the centroid of each cluster.

แล้วจบ

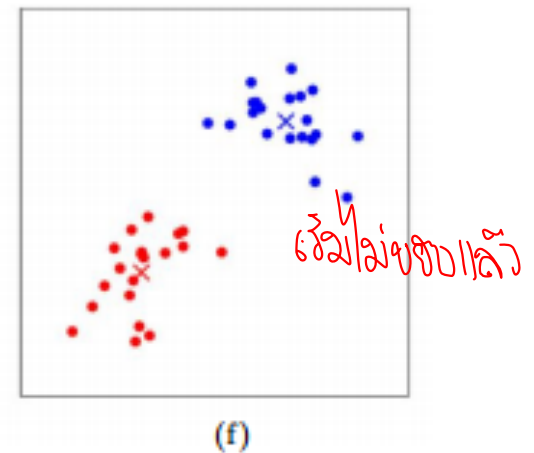
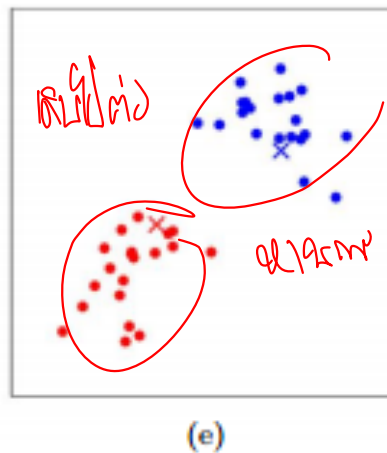
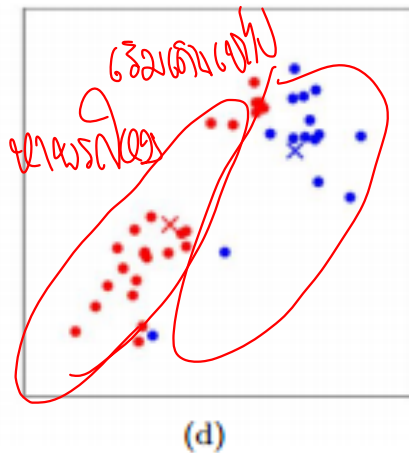
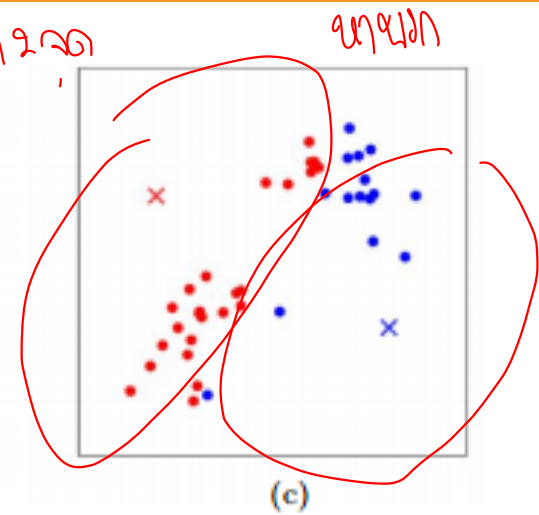
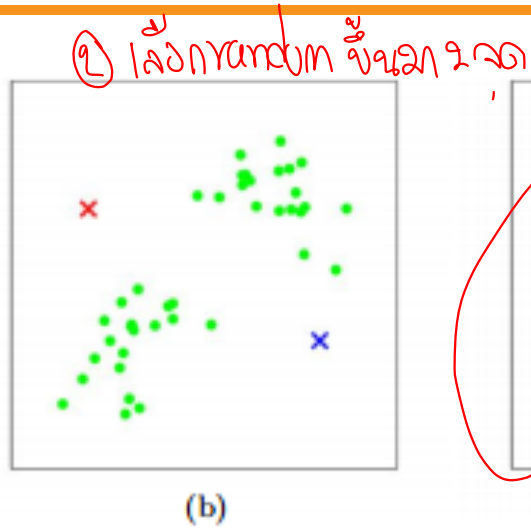
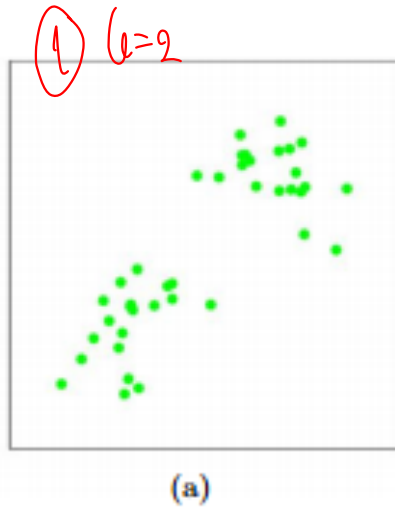
5: **until** The centroids don't change

---

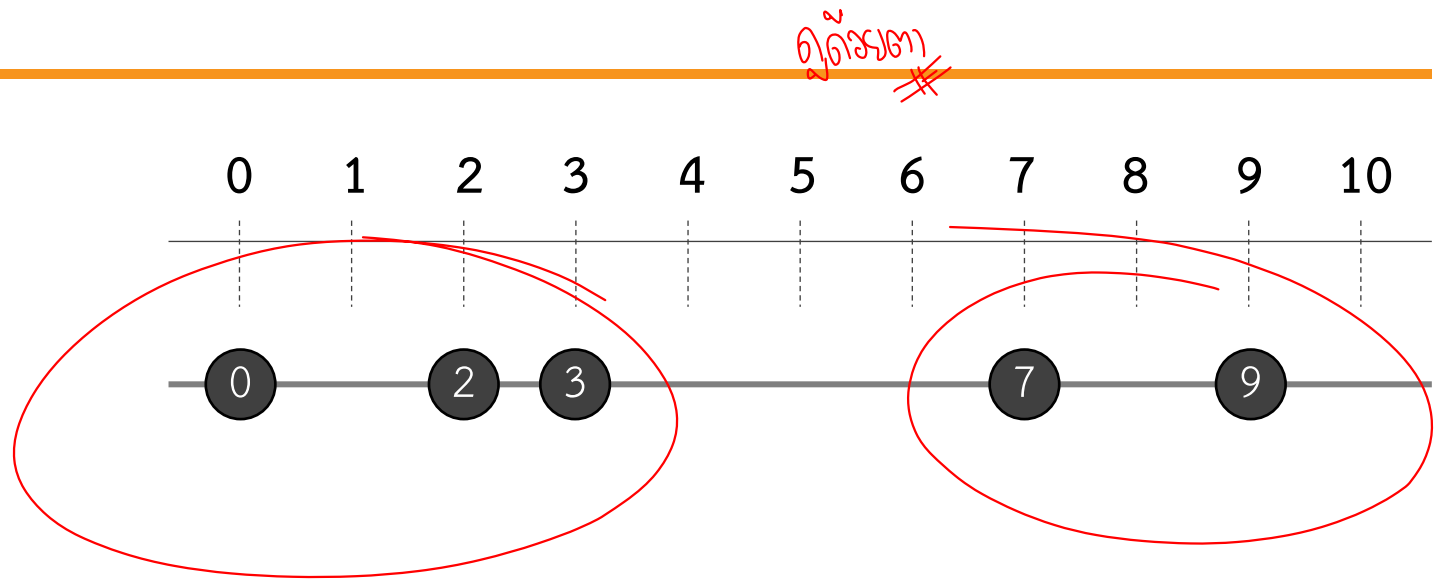
- Initial centroids are often chosen **randomly**.
  - Clusters produced vary from one run to another.

# K-means: Steps

ตัวอย่างการคำนวณ

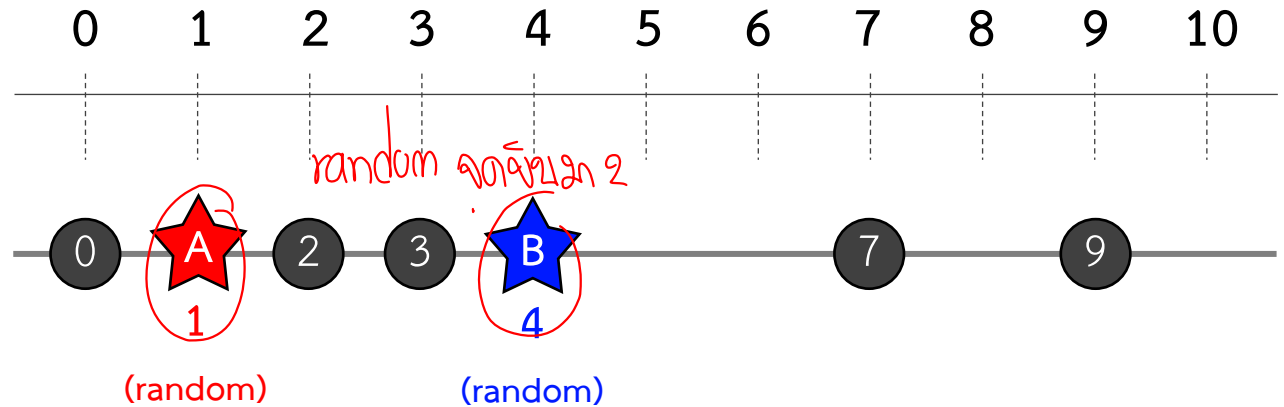


# K-means: Example Steps ( 1 feature )



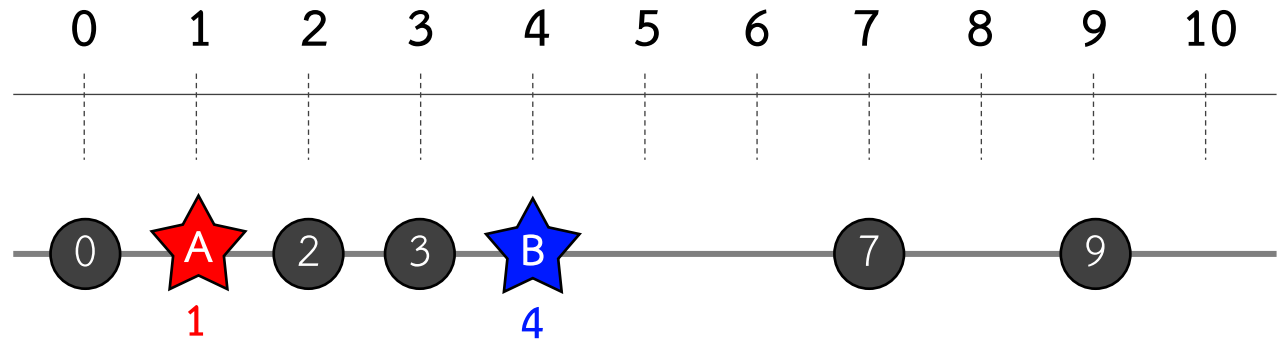
# K-means: Example Steps ( 1 feature )

Round 0



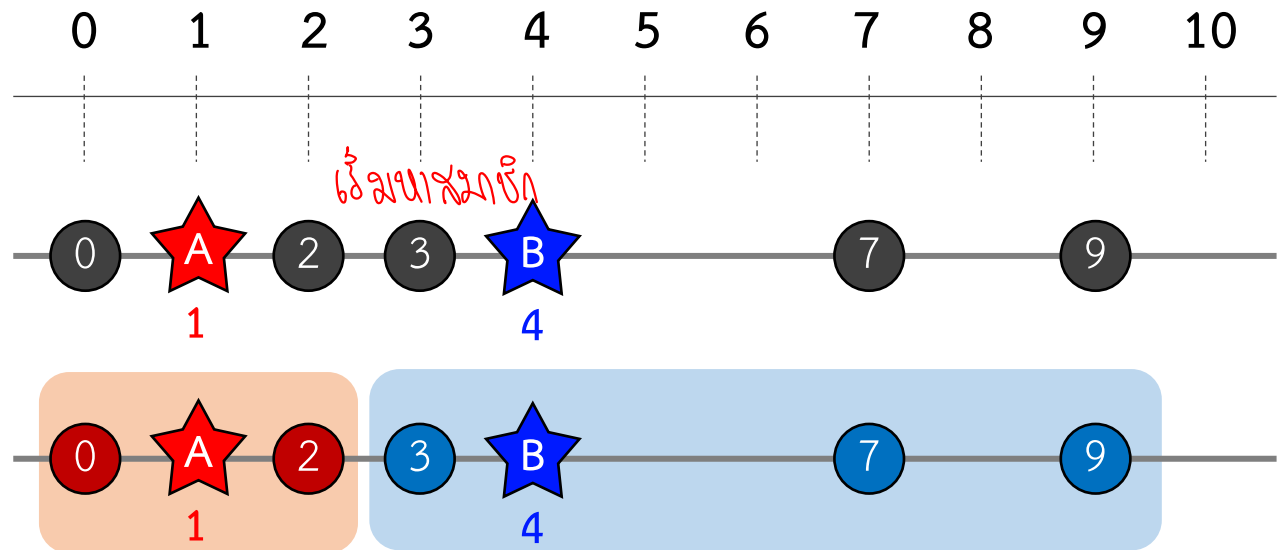
# K-means: Example Steps ( 1 feature )

Round 1



# K-means: Example Steps ( 1 feature )

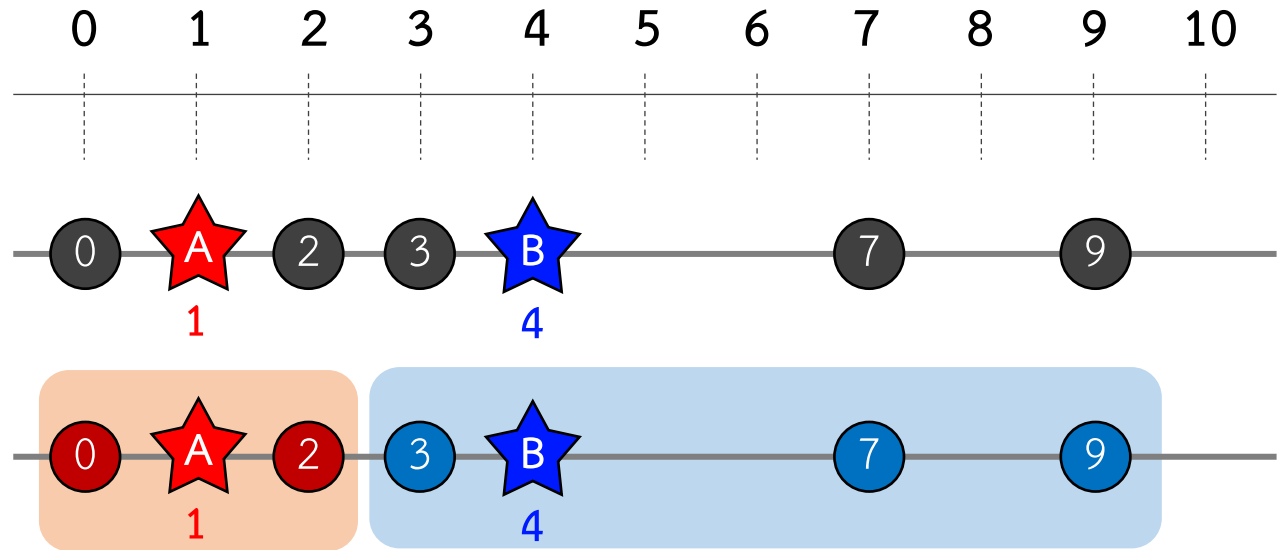
Round 1





# K-means: Example Steps ( 1 feature )

Round 1



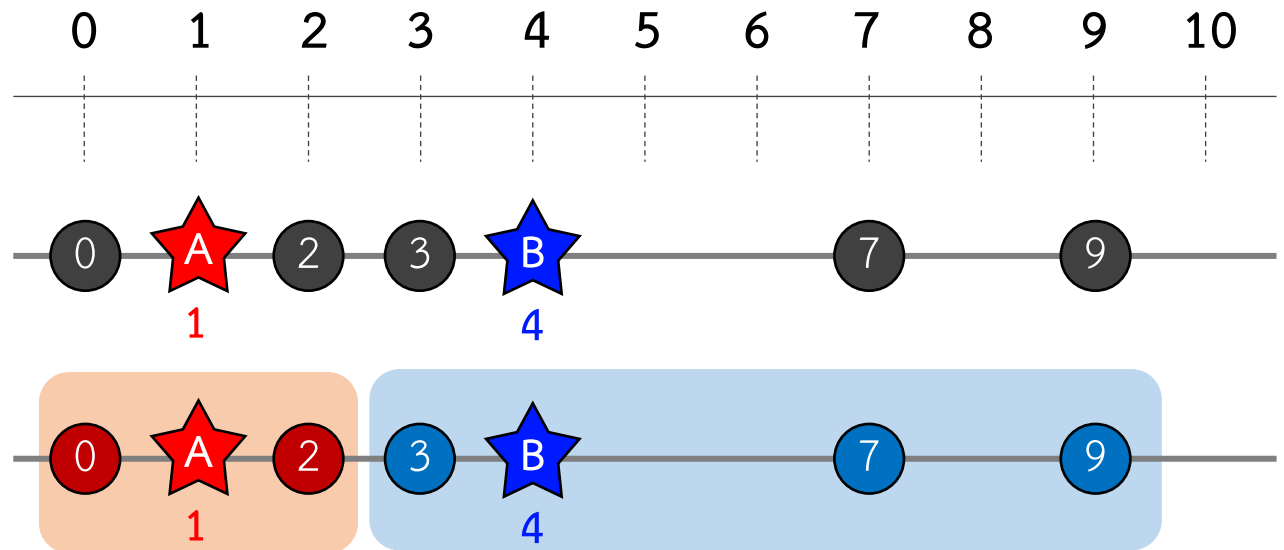
New Center of A =  $(0+2)/2 = 1.0$

New Center of B =  $(3+7+9)/3 = 6.3$

ค่าเฉลี่ยใหม่

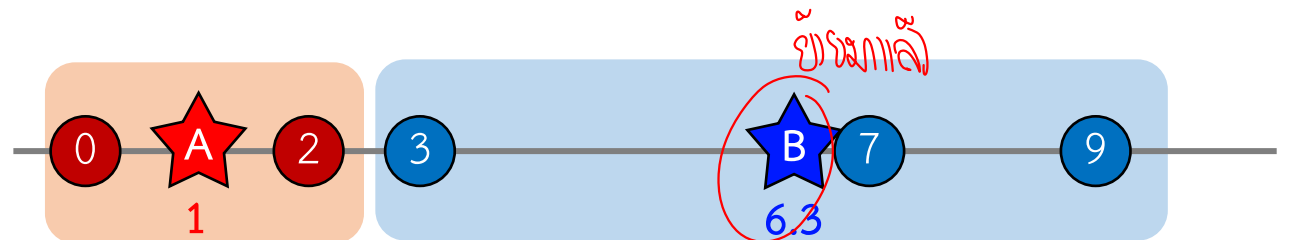
# K-means: Example Steps ( 1 feature )

Round 1

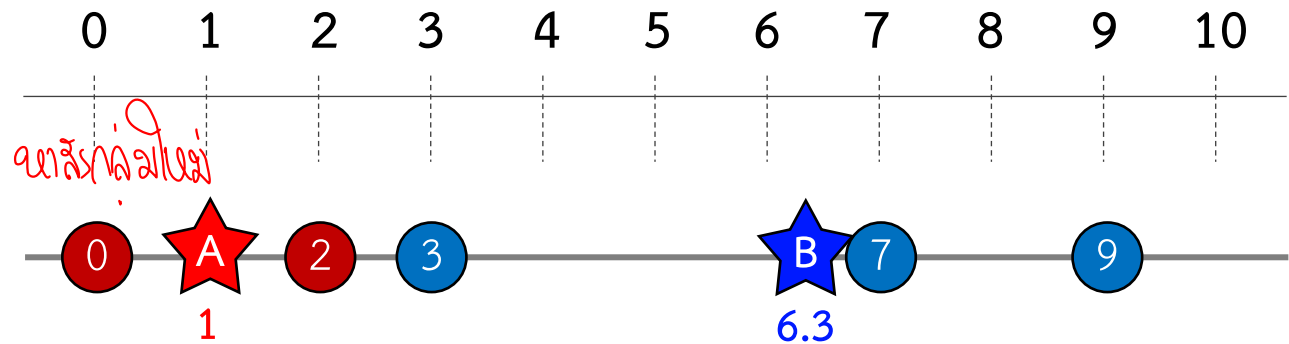


New Center of A =  $(0+2)/2 = 1.0$

New Center of B =  $(3+7+9)/3 = 6.3$



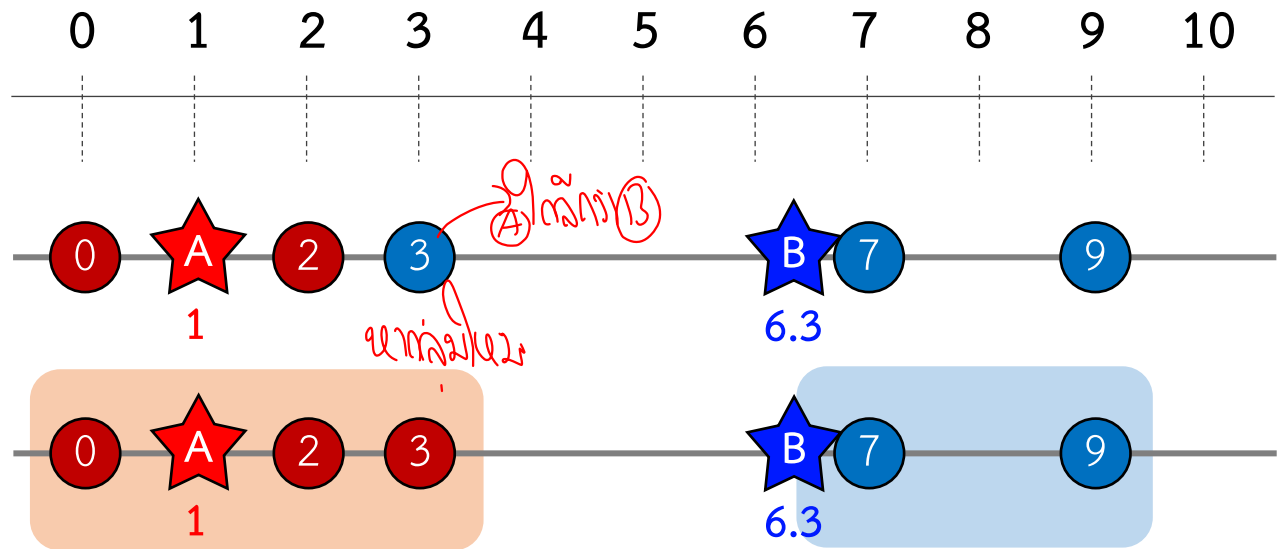
# K-means: Example Steps ( 1 feature )



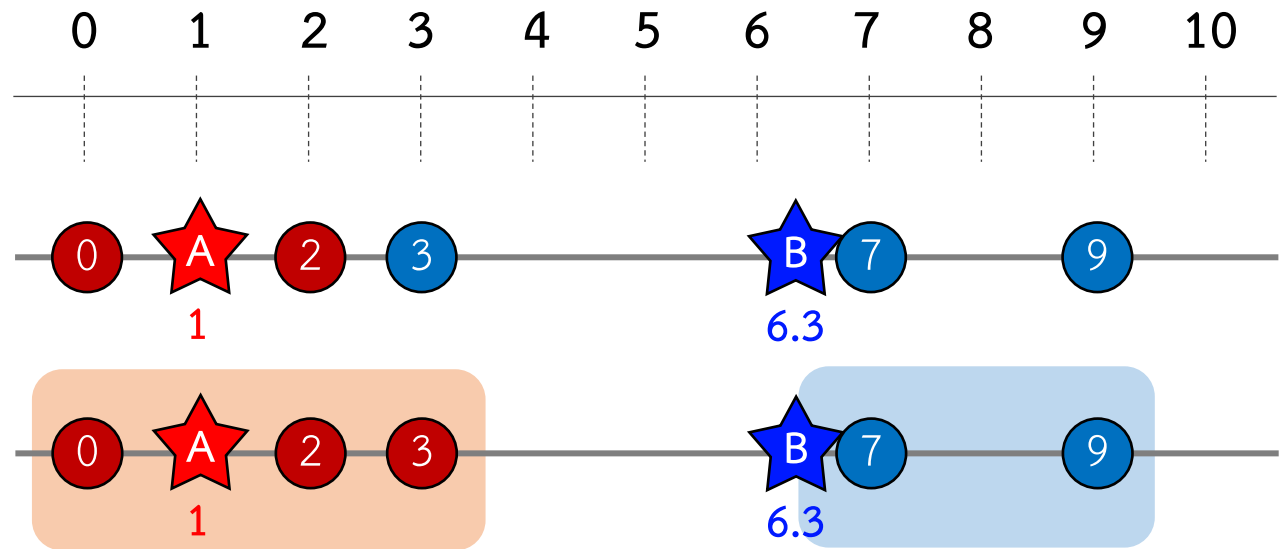
Round 2

# K-means: Example Steps ( 1 feature )

Round 2



# K-means: Example Steps ( 1 feature )



Round 2

$$\text{New Center of A} = (0+2+3)/3 = 1.7$$

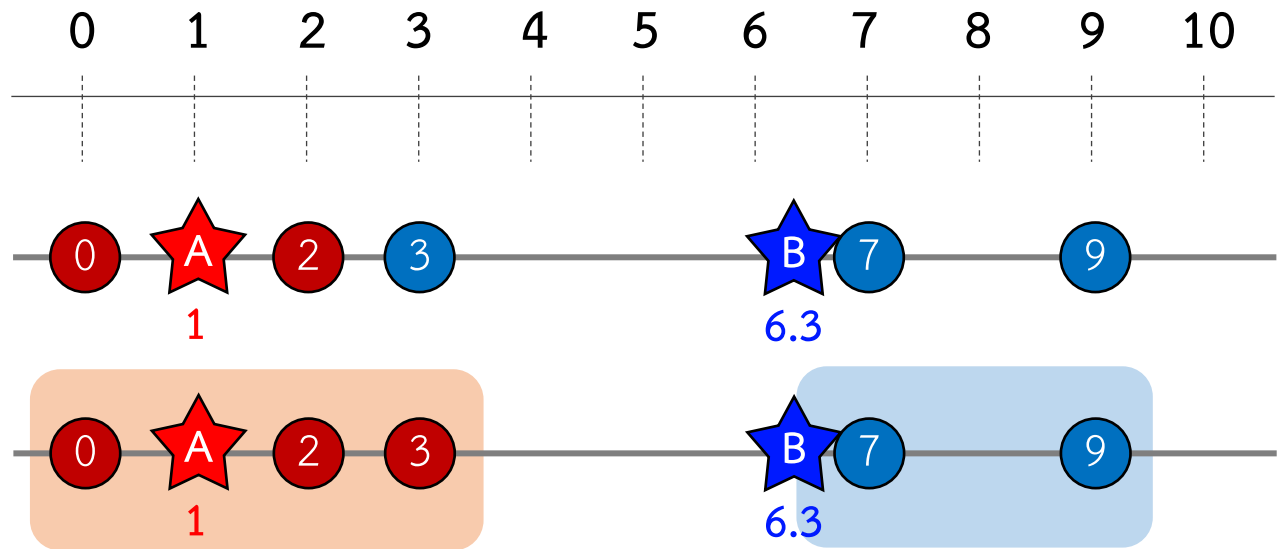
*for B*

$$\text{New Center of B} = (7+9)/2 = 8$$

*for A*

# K-means: Example Steps ( 1 feature )

Round 2

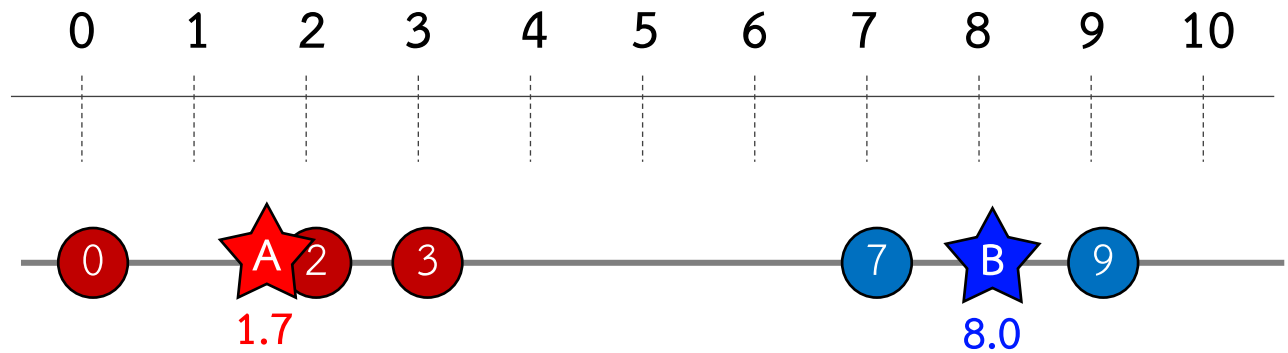


New Center of A =  $(0+2+3)/3 = 1.7$

New Center of B =  $(7+9)/2 = 8$

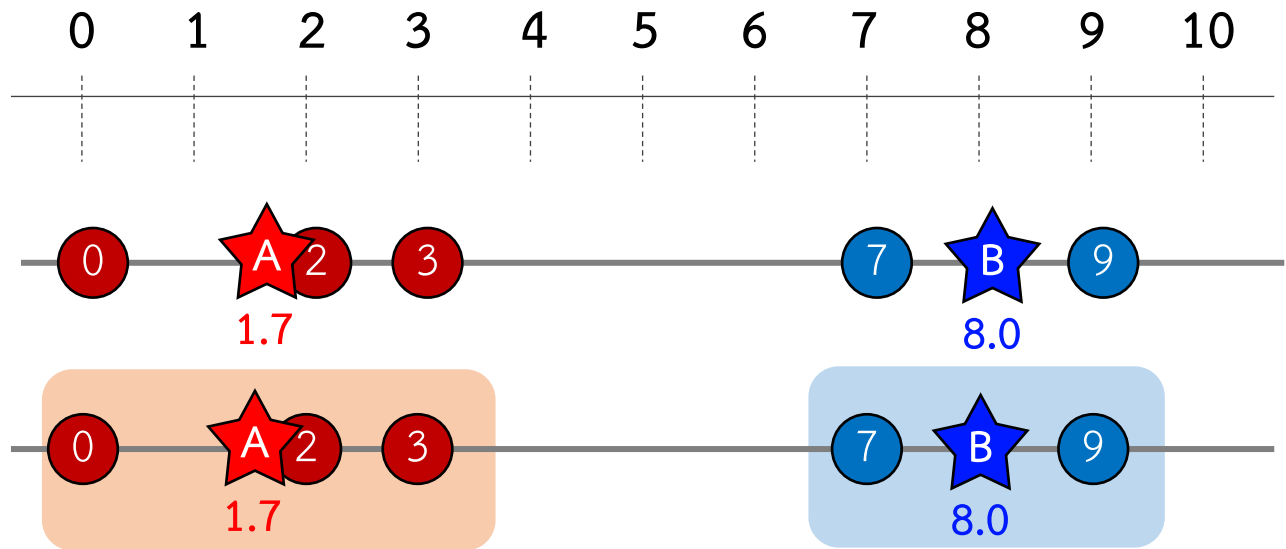


# K-means: Example Steps ( 1 feature )



Round 3

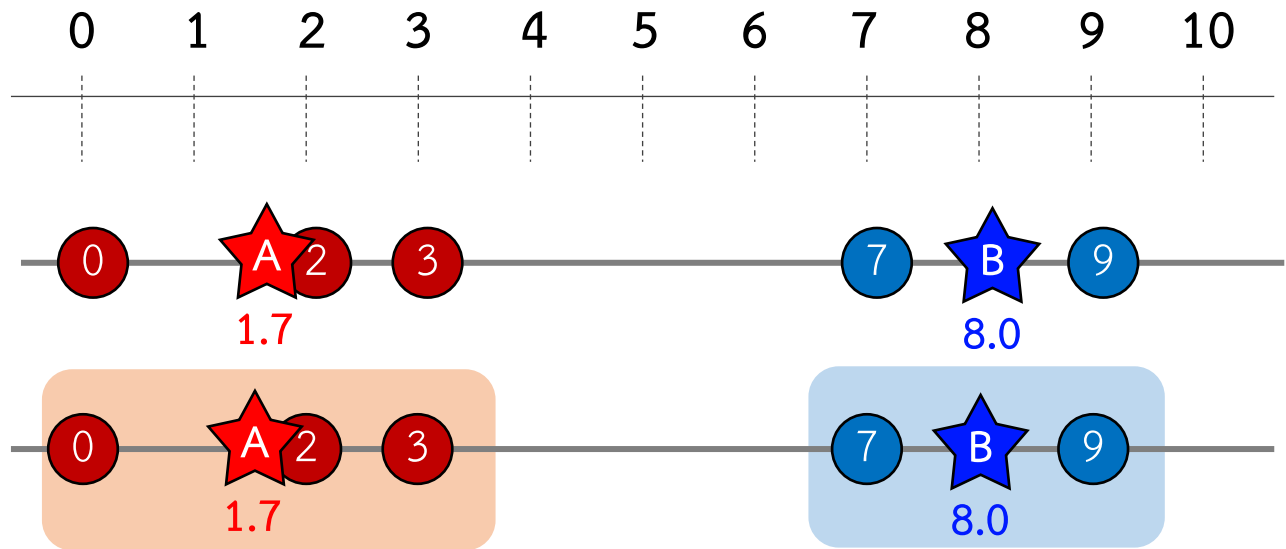
# K-means: Example Steps ( 1 feature )



Round 3



# K-means: Example Steps ( 1 feature )

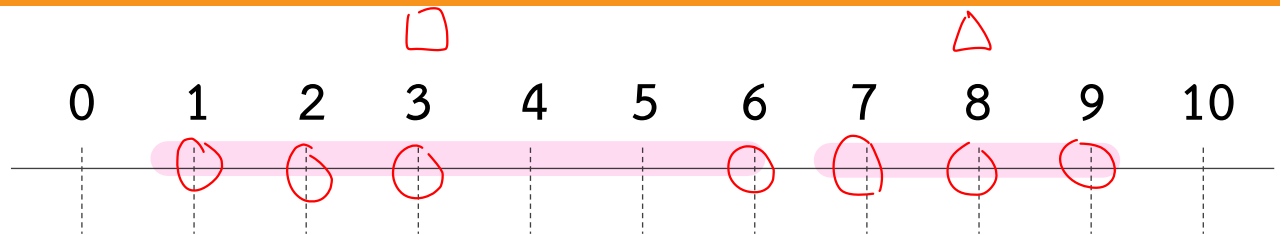


Round 3

$$\text{New Center of A} = (0+2+3)/3 = 1.7$$

$$\text{New Center of B} = (7+9)/2 = 8$$

# K-means: Example Steps ( 1 feature )



Round 3

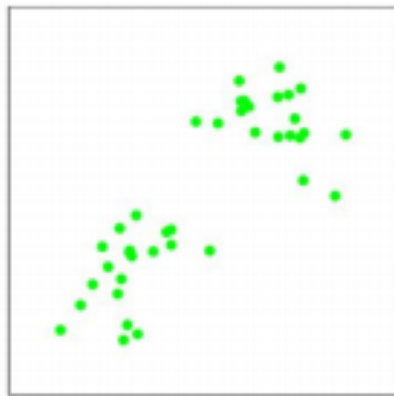


New Center of A =  $(0+2+3)/3 = 1.7$

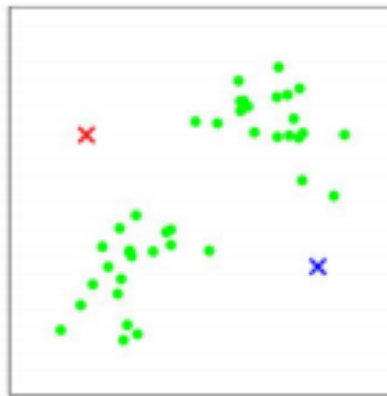
New Center of B =  $(7+9)/2 = 8$



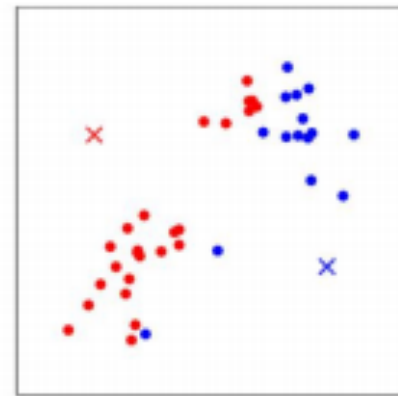
# K-means: Example (2 features)



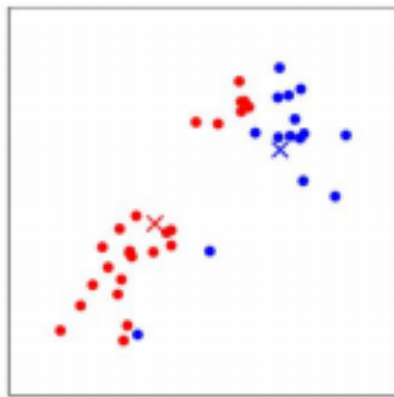
(a)



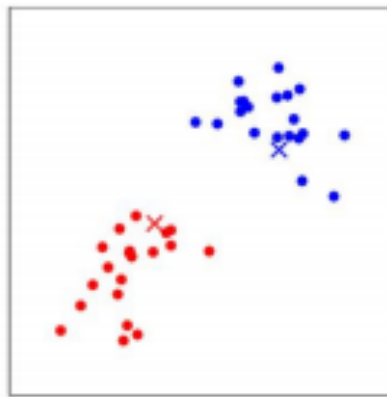
(b)



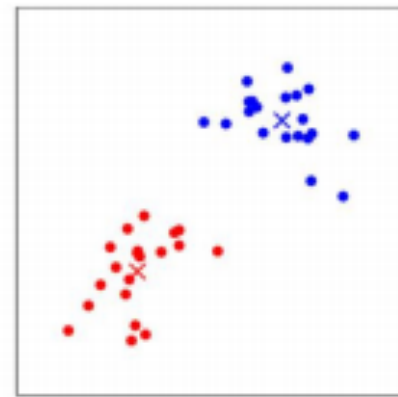
(c)



(d)



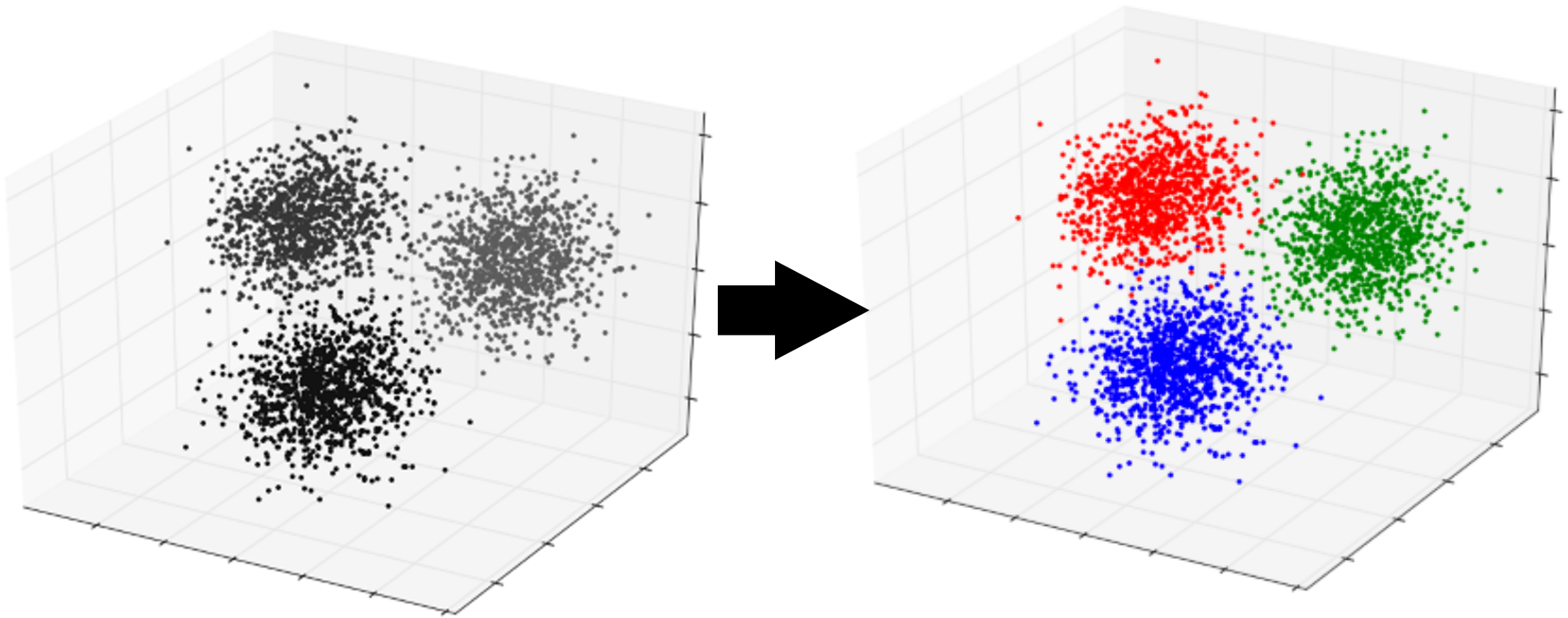
(e)



(f)

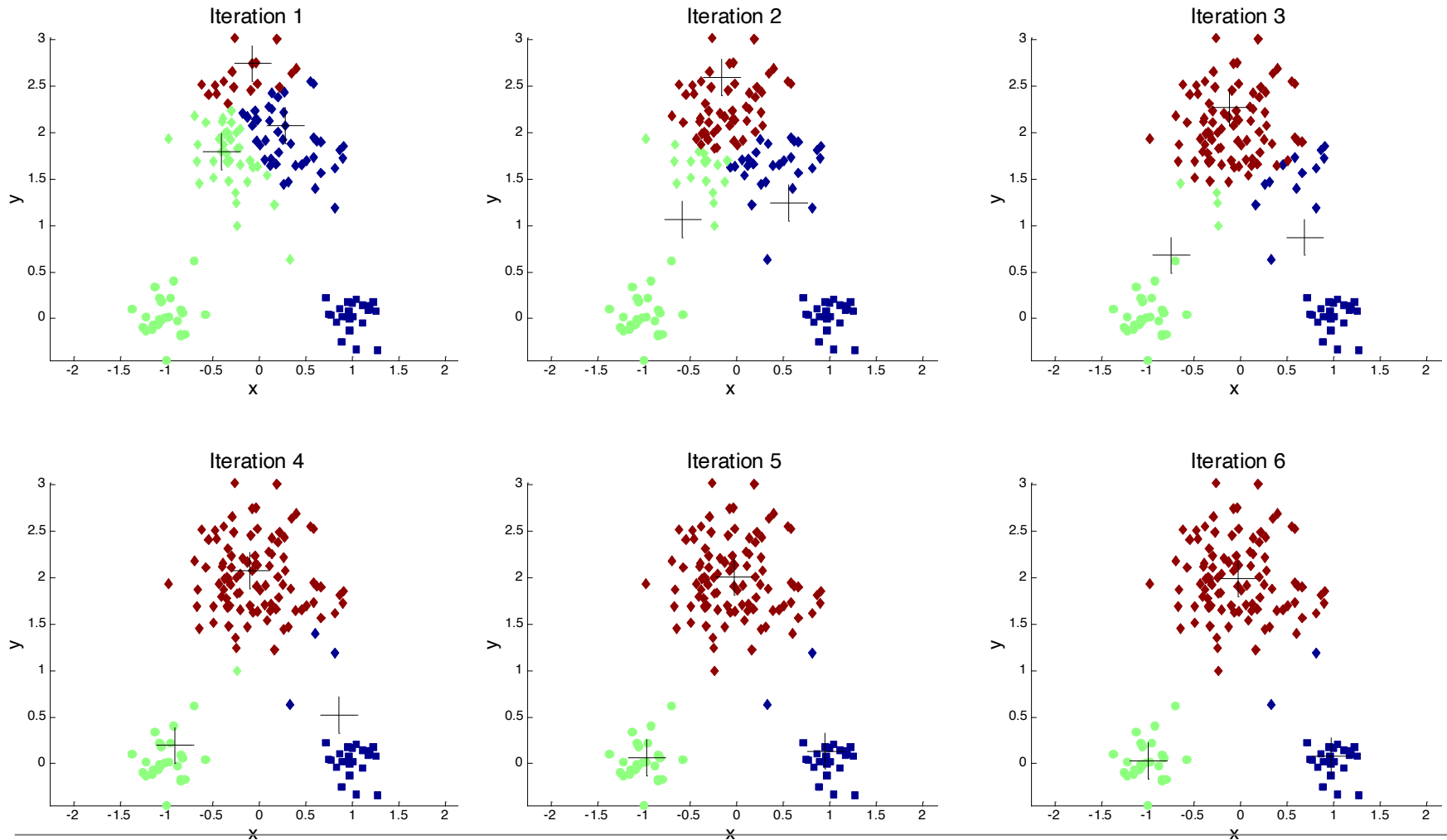
# K-means: Example (3 features)

သော feature သုံးချောင်း  $\rightarrow$  visualize စက်ကိတ်



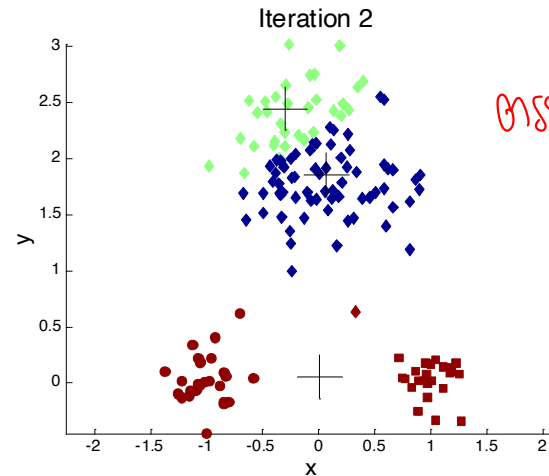
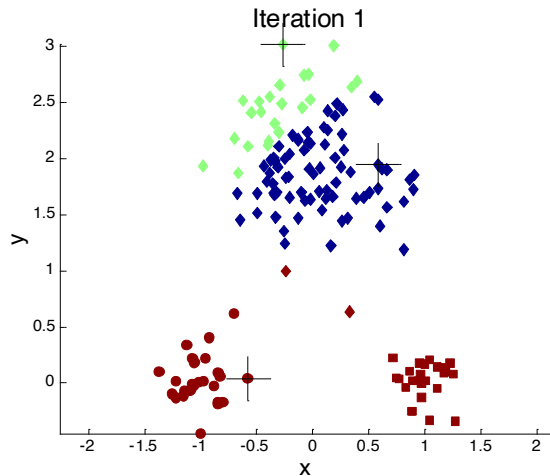
# Importance of Choosing Initial Centroids (A)

ถ้าเราเลือก initial centroids random ก็อาจมีปัญหา

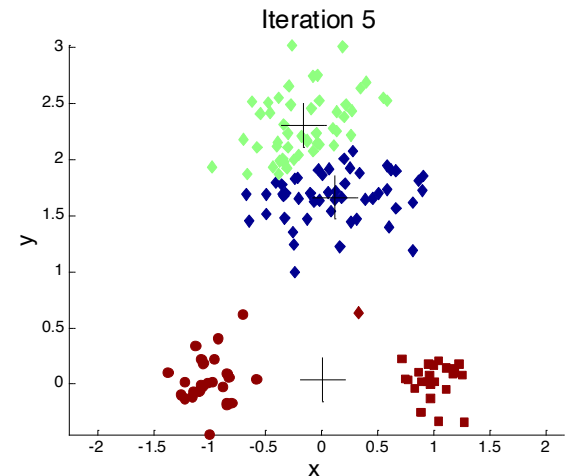
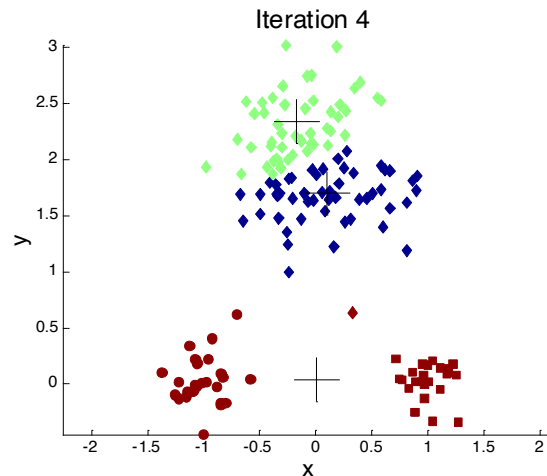
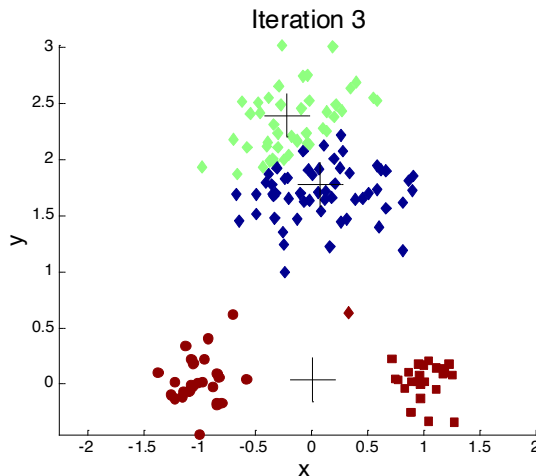


Ref: • Min-Hashing, "Locality Sensitive Hashing Clustering"

# Importance of Choosing Initial Centroids (B)



การสุ่มที่เปลี่ยนไป?  
↓  
random จากจุด  
ที่แน่นอน



# Dealing with Initialization

ถ้า → ① ทำซ้ำ  
② กำหนดค่าให้ตัวแปร

- Do **multiple runs** and select the clustering with the smallest error
- Select original set of points by methods other than random . E.g., pick the most distant (from each other) points as cluster centers

# Limitations of K-means

---

- K-means has problems when clusters are of different
  - Sizes
  - Densities
  - Non-globular shapes
- K-means has problems when the data contains outliers.

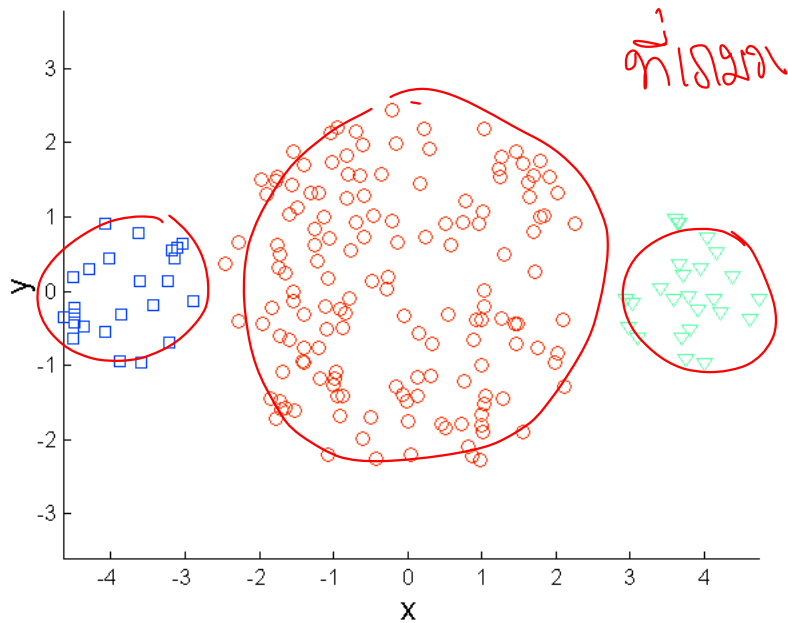
various

non-globular  
shapes

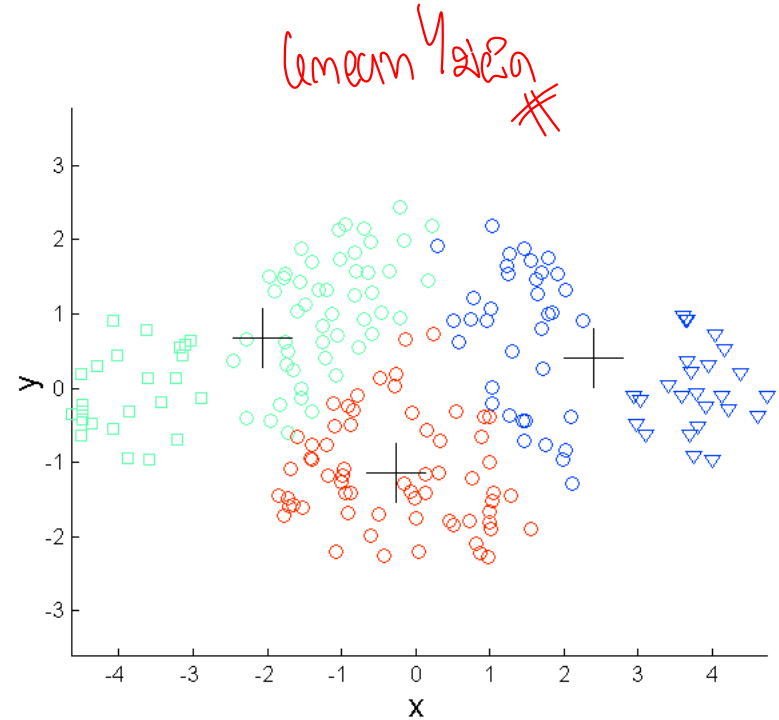
min outliers



# Limitations of K-means: Differing Sizes

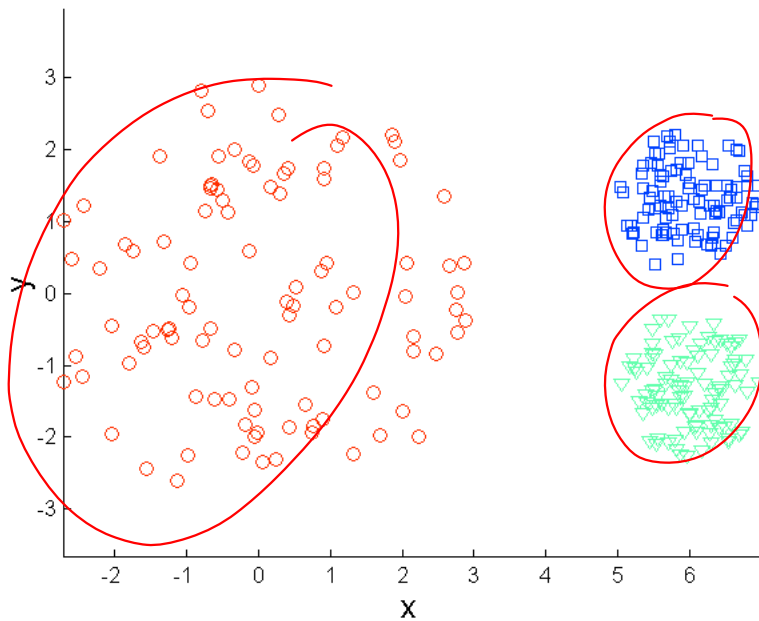


Original Points

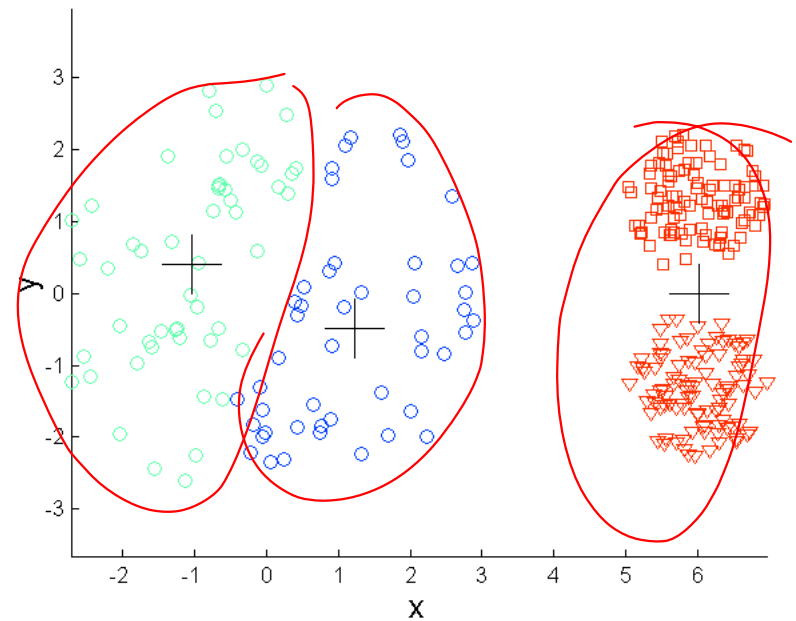


K-means (3 Clusters)

# Limitations of K-means: Differing Density



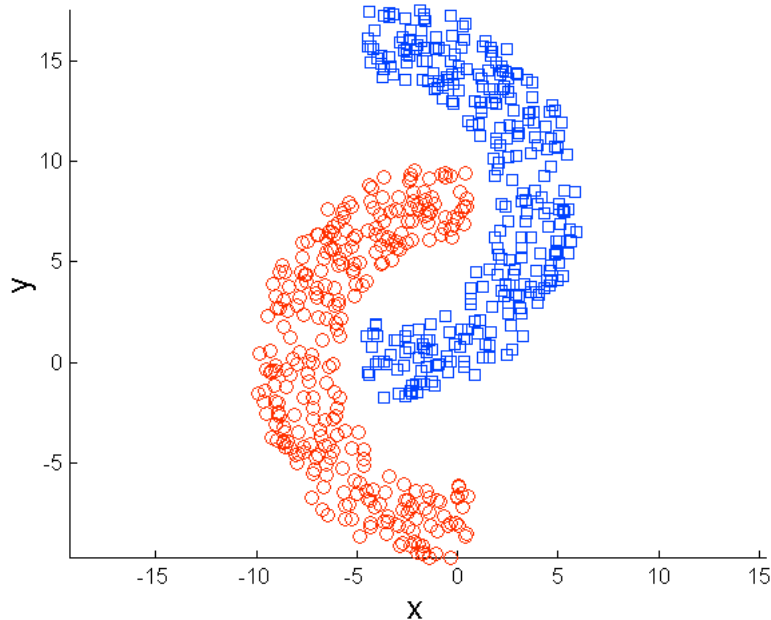
Original Points



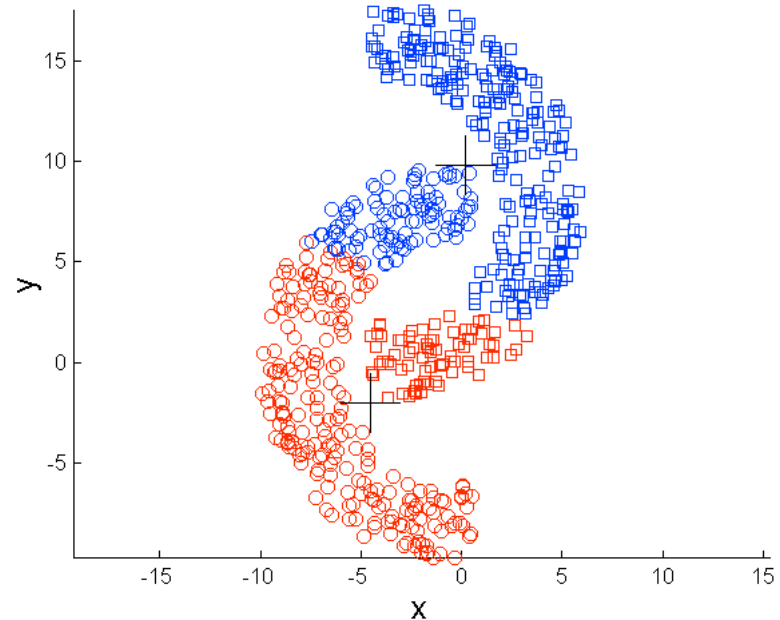
K-means (3 Clusters)

# Limitations of K-means: Non-globular Shapes

ຂໍ້ຈຳກັດຂອງ K-means (ຕາມໂຄງ) ຂາດໄປ



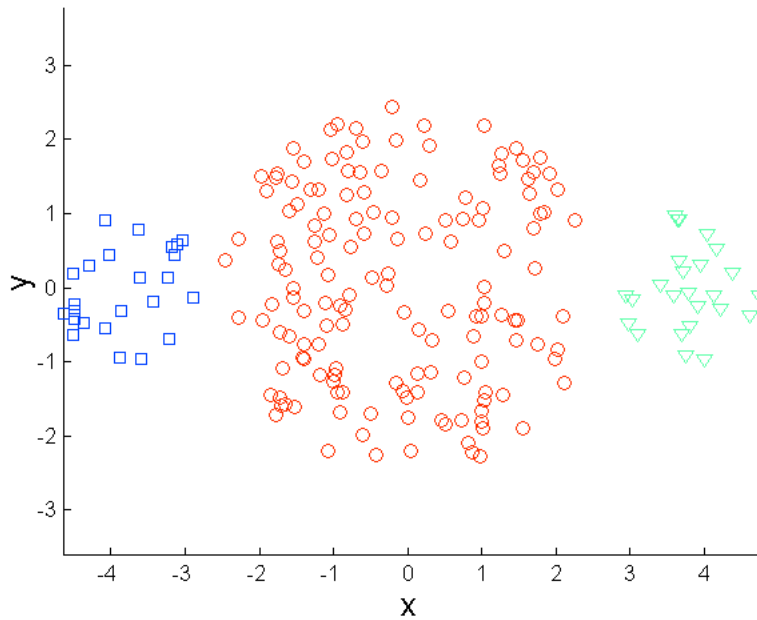
Original Points



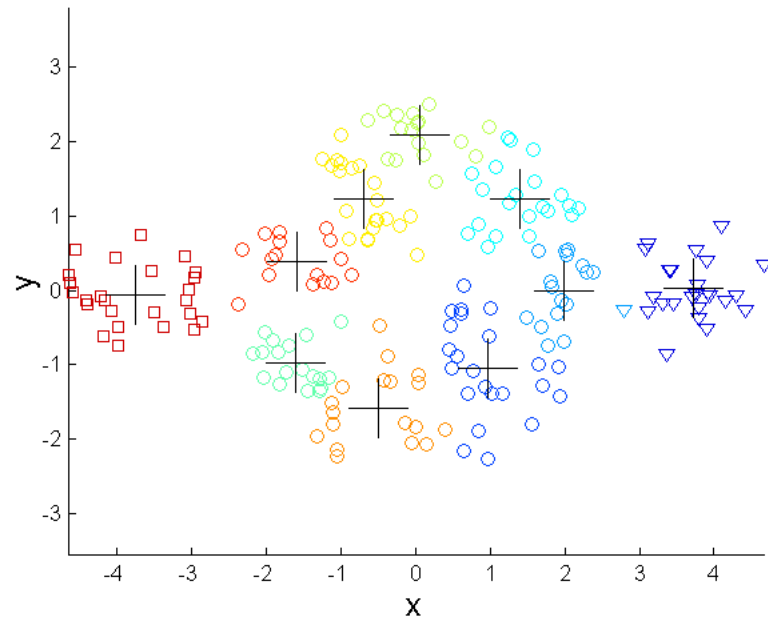
K-means (2 Clusters)

# Overcoming K-means Limitations

set to be using



Original Points

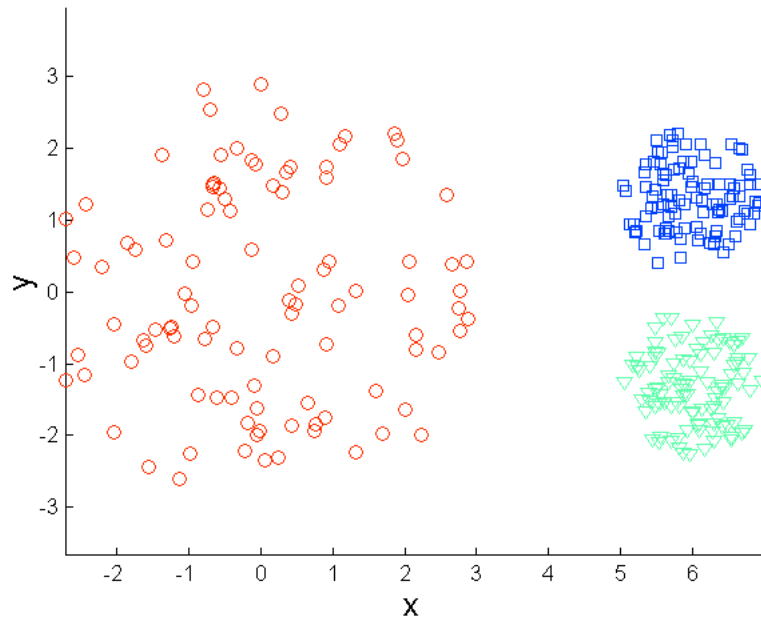


K-means Clusters

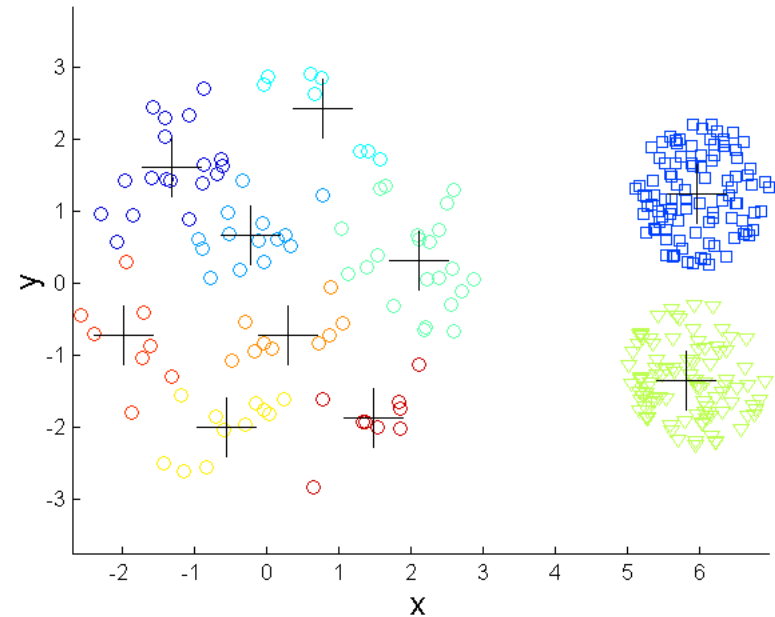
One solution is to use many clusters.

Find parts of clusters, but need to put together.

# Overcoming K-means Limitations

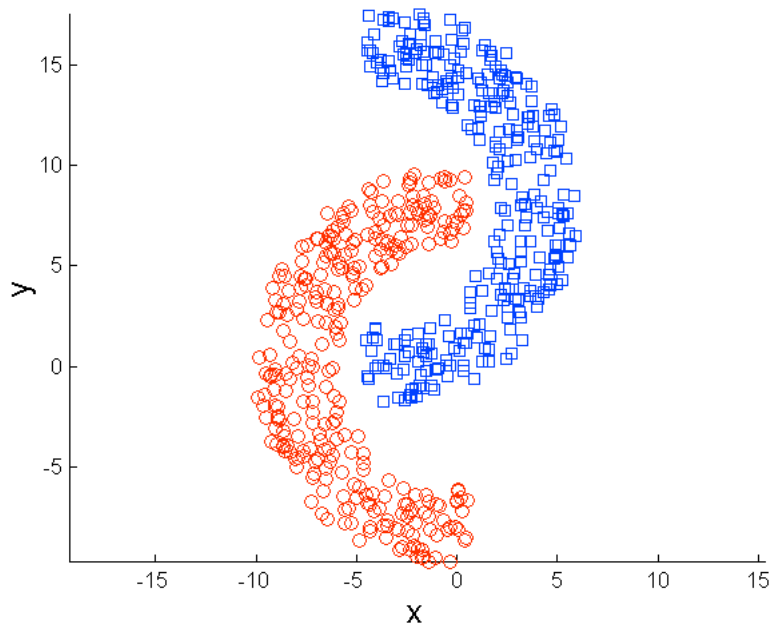


Original Points

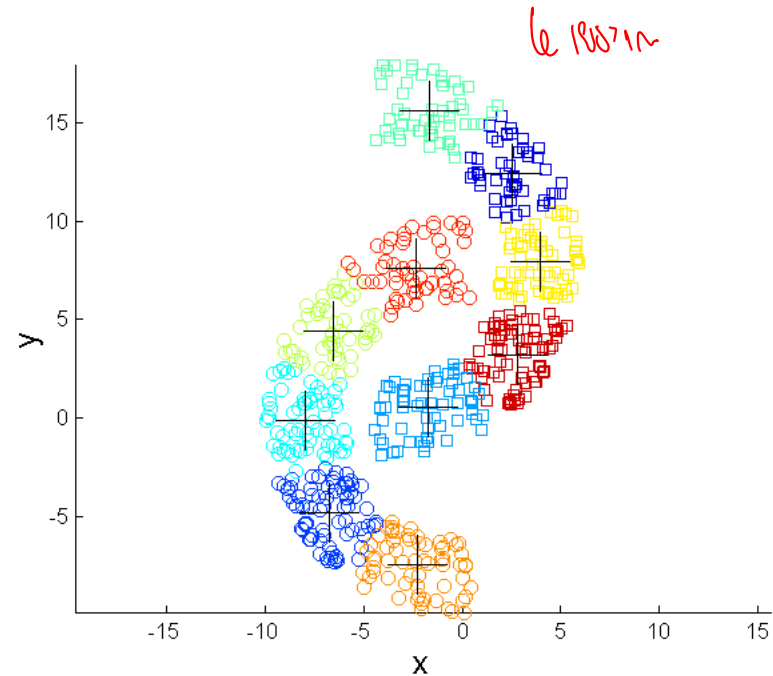


K-means Clusters

# Overcoming K-means Limitations



Original Points



K-means Clusters

# Variations

ខ្មែរ

ការបំបែកទិន្នន័យ

- **K-medoids** → ការបំបែកទិន្នន័យដោយយកចំណុចទៅជាកណ្តាល  
  - Similar problem definition as in K-means, but the centroid of the cluster is defined to be one of the points in the cluster (the **medoid**).
- **K-centers** → ការបំបែកទិន្នន័យដោយយកចំណុចទៅជាកណ្តាល  
  - Similar problem definition as in K-means, but the goal now is to minimize the maximum **diameter** of the clusters (diameter of a cluster is maximum distance between any two points in the cluster).

# Python

গিট বিব এন্ড লাইসেন্স

```
>>> from sklearn.cluster import KMeans
>>> import numpy as np
>>> X = np.array([[1, 2], [1, 4], [1, 0],
...               [4, 2], [4, 4], [4, 0]])
>>> kmeans = KMeans(n_clusters=2, random_state=0).fit(X)
>>> kmeans.labels_
array([0, 0, 0, 1, 1, 1], dtype=int32)
>>> kmeans.predict([[0, 0], [4, 4]])
array([0, 1], dtype=int32)
>>> kmeans.cluster_centers_
array([[ 1.,  2.],
       [ 4.,  2.]])
```



# DBSCAN

1999, Ester et al.

↓  
ใช้วิธี k-means กับ data → 2 clusters  
↓  
↓  
noise and data



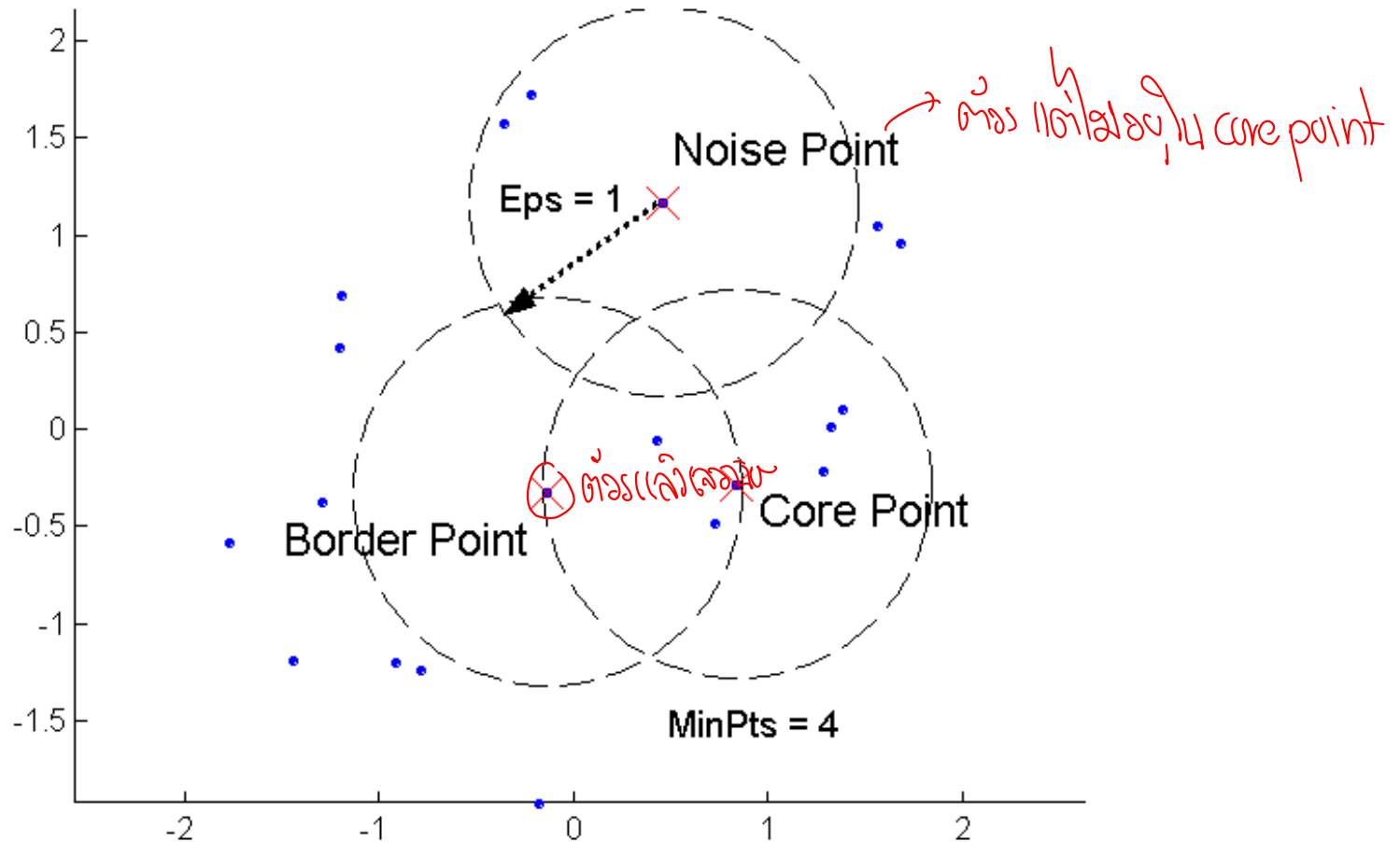
# DBSCAN: Density-Based Clustering

- DBSCAN is a Density-Based Clustering algorithm
  - Reminder: In density based clustering we partition points into dense regions separated by not-so-dense regions.
  - Important Questions:
    - How do we measure density?
    - What is a dense region?
  - DBSCAN:
    - Density at point  $p$ : number of points within a circle of radius  $Eps$
    - Dense Region: A circle of radius  $Eps$  that contains at least  $MinPts$  points
- หมายเหตุ: จุดที่อยู่ในก้อนเดียวกัน  
แล้วเชื่อมต่อกันได้
- ① เราใช้รัศมี Eps
- ② ผลลัพธ์คือ Eps  
แล้วใช้ MinPts  
เพื่อหาจุด

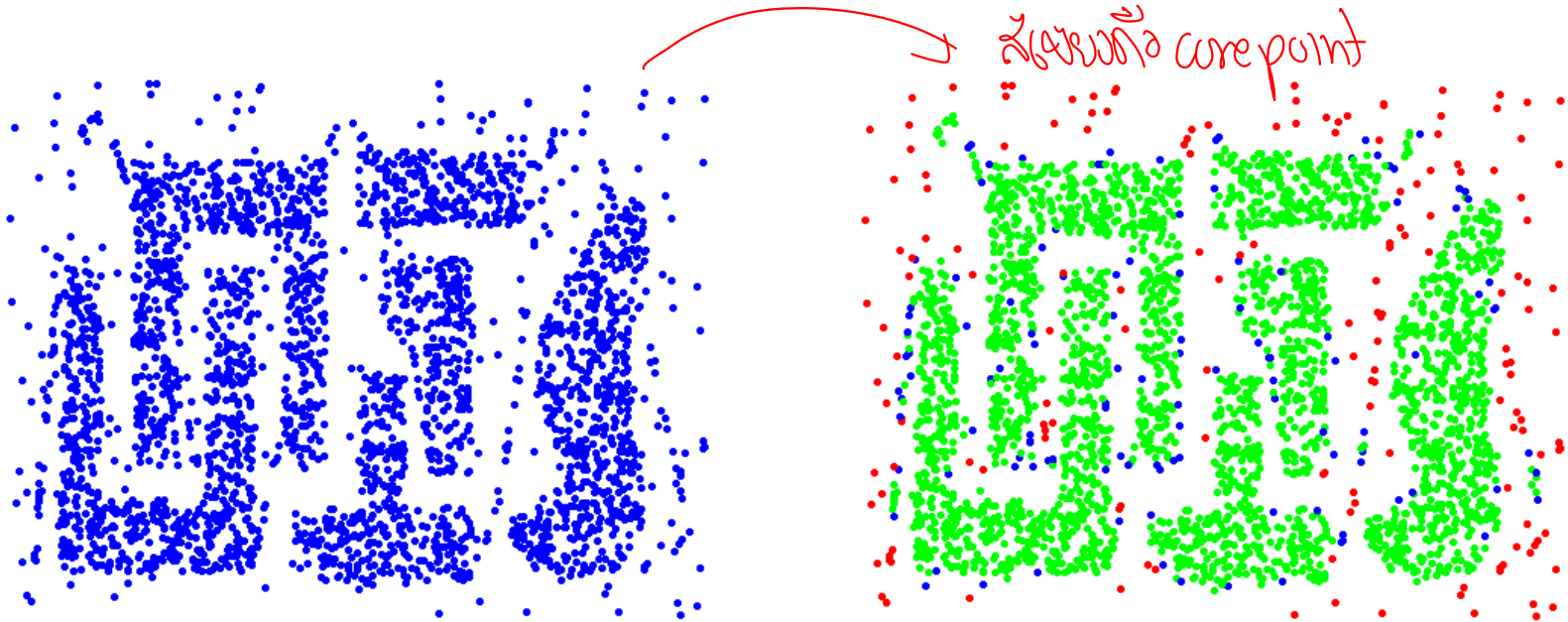
# DBSCAN

- Characterization of points ถ้าเราเลือกค่าของ  $\epsilon$  แล้ว  $\epsilon$  ก็คือรัศมีที่รอบๆ 3 แฉ
  - A point is a core point ① จำนวนรอบข้าง  $\geq \text{min\_points}$  if it has more than a specified number of points (**MinPts**) within **Eps**
    - These points belong in a **dense region** and are at the **interior** of a cluster
  - A border point ② จำนวนรอบข้าง  $< \text{min\_point}$  has fewer than **MinPts** within **Eps**, but is in the neighborhood of a **core point**.
  - A **noise point** is any point that is not a core point or a border point. ③ ไม่เจอข้างไหนเลย  $\rightarrow$  ตัดทิ้งได้

# DBSCAN: Core, Border, and Noise Points



# DBSCAN: Core, Border, and Noise Points



Original Points

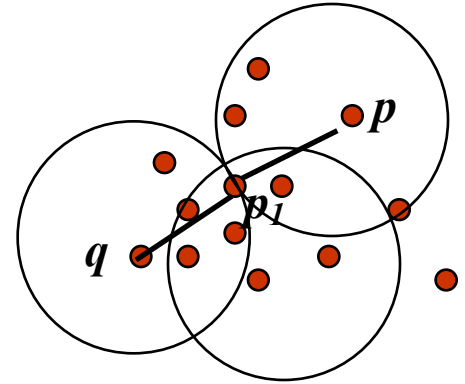
Point types: **core**,  
**border** and **noise**

Eps = 10, MinPts = 4

# Density-Connected points

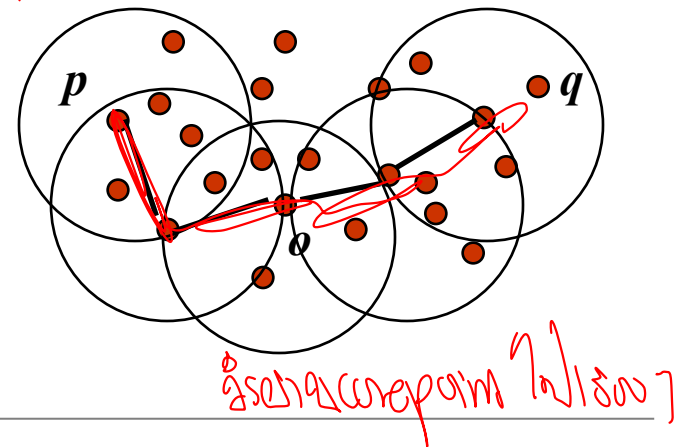
- Density edge

- We place an **edge** between two core points **q** and **p** if they are within distance **Eps**.



- Density-connected *ဒီပက်စီတီဆက်သွယ်မှုရှိသော နံပါတ်*

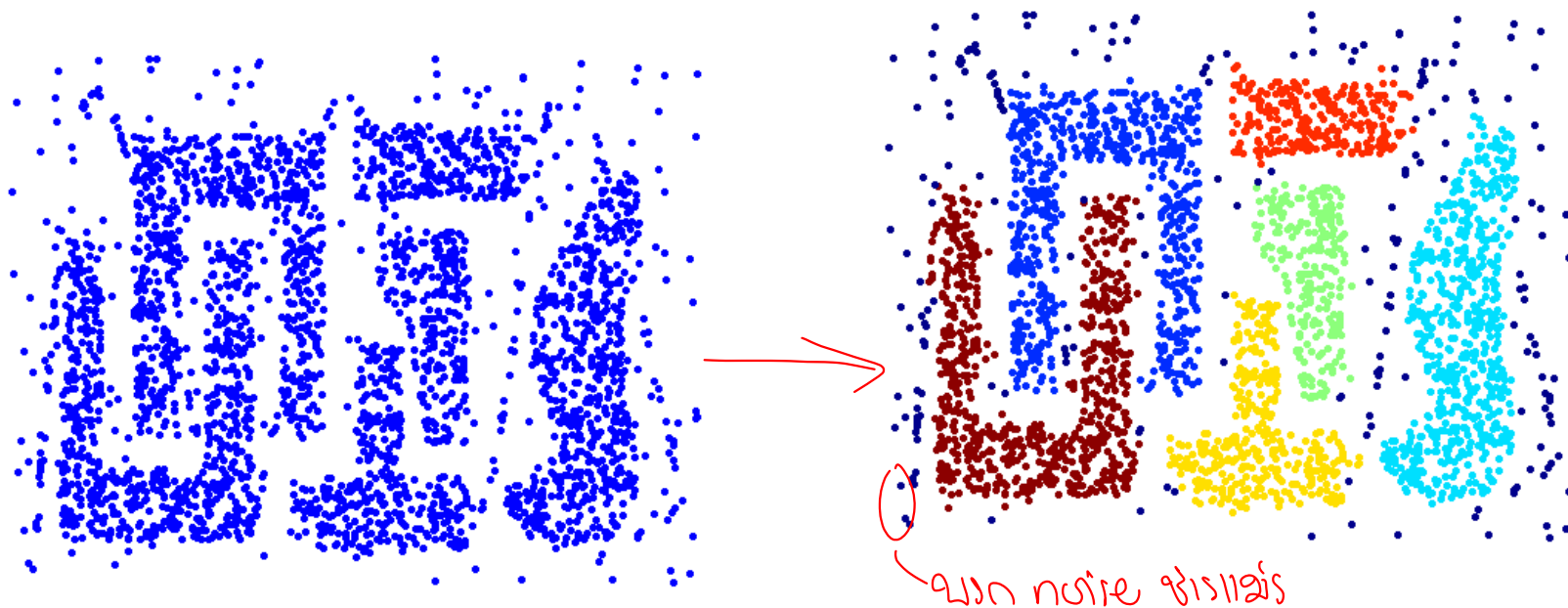
- A point **p** is **density-connected** to a point **q** if there is a **path of edges** from **p** to **q**



# DBSCAN Algorithm

- Label points as core, border and noise ① assign label for data
- Eliminate noise points ② ลบทิ้ง
- For every **core** point **p** that has not been assigned to a cluster  
core point ที่ยังไม่ได้ assign เป็น cluster ใด ๆ  
  - Create a new cluster with the point **p** and all the points that are **density-connected** to **p**.
- Assign **border** points to the cluster of the closest core point.

# When DBSCAN Works Well



Original Points

Clusters

- Resistant to Noise
- Can handle clusters of different shapes and sizes



# Cluster Validation

วัดความถูกต้องของ cluster



วัดภายในกลุ่มต้องกันติดกัน  
ต่างกับ ตัวไม่แนบกัน



# Different Aspects of Cluster Validation

---

1. Determining the **clustering tendency** of a set of data, i.e., distinguishing whether non-random structure actually exists in the data.
2. Comparing the results of a cluster analysis to externally known results, e.g., to externally given class labels.
3. Evaluating how well the results of a cluster analysis fit the data *without* reference to external information.
  - Use only the data
4. Comparing the results of two different sets of cluster analyses to determine which is better.
5. Determining the ‘correct’ number of clusters.

For 2, 3, and 4, we can further distinguish whether we want to evaluate the entire clustering or just individual clusters.

# Measures of Cluster Validity

- Numerical measures that are applied to judge various aspects of cluster validity, are classified into the following three types. *สามวิธี 3 แบบ*
  - External Index:** Used to measure the extent to which cluster labels match externally supplied class labels.
    - Entropy *↓ การเปรียบเทียบกับกลุ่มที่จับตามองอยู่แล้ว → entropy*
  - Internal Index:** Used to measure the goodness of a clustering structure *without* respect to external information.
    - Sum of Squared Error (SSE) *↓ วัดความเฉลี่ยภายในกลุ่มตัวเอง → SSE*
  - Relative Index:** Used to compare two different clusterings or clusters.
    - Often an external or internal index is used for this function, e.g., SSE or entropy
- Sometimes these are referred to as **criteria** instead of **indices** *↓ การเทียบกับเกณฑ์อื่น ดูว่าอันไหน*
  - However, sometimes criterion is the general strategy and index is the numerical measure that implements the criterion. *ได้ทั้ง SSE, entropy ทั้ง*

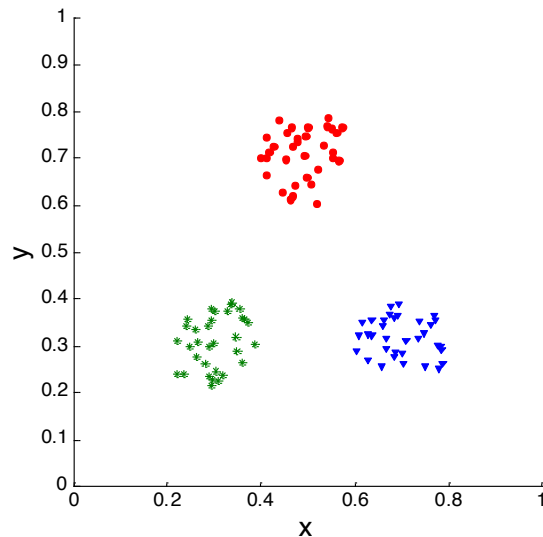
# Measuring Cluster Validity Via Correlation

---

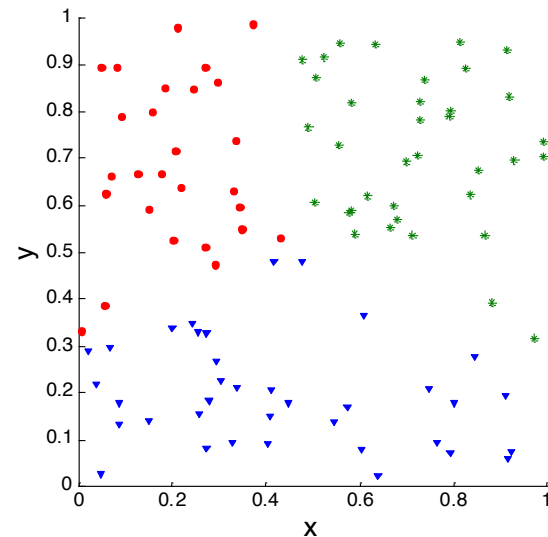
- Two matrices
  - Proximity Matrix
  - Ideal Similarity Matrix
    - One row and one column for each data point
    - An entry is 1 if the associated pair of points belong to the same cluster
    - An entry is 0 if the associated pair of points belongs to different clusters
- Compute the correlation between the two matrices
  - Since the matrices are symmetric, only the correlation between  $n(n-1) / 2$  entries needs to be calculated.
- High correlation indicates that points that belong to the same cluster are close to each other.
- Not a good measure for some density or contiguity based clusters.

# Measuring Cluster Validity Via Correlation

- Correlation of ideal similarity and proximity matrices for the K-means clusterings of the following two data sets.



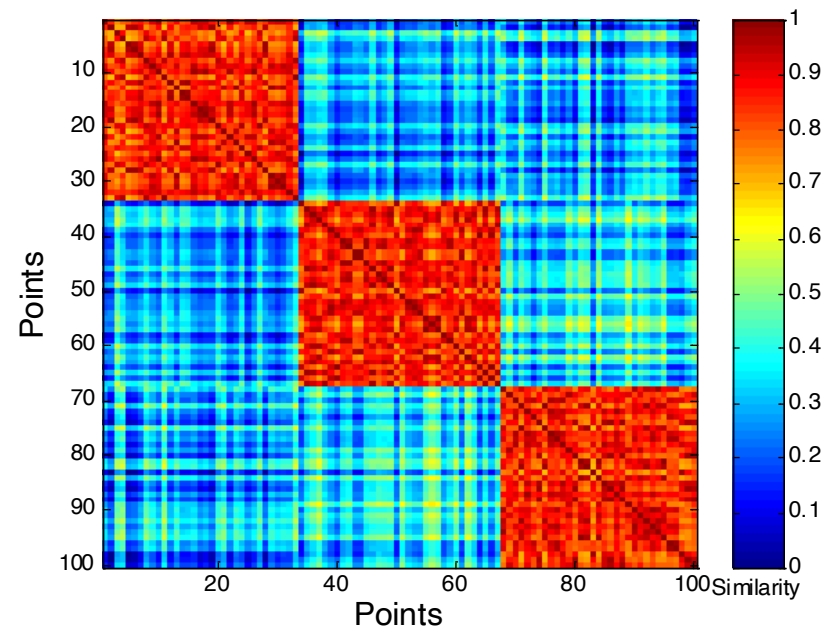
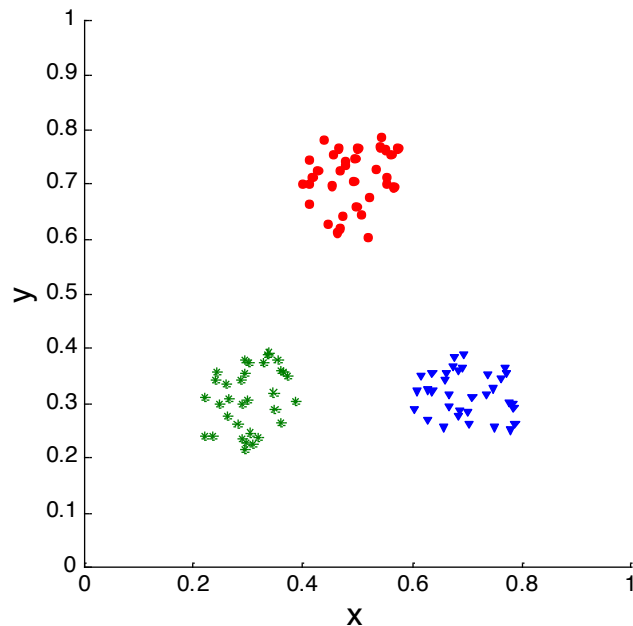
**Corr = -0.9235**



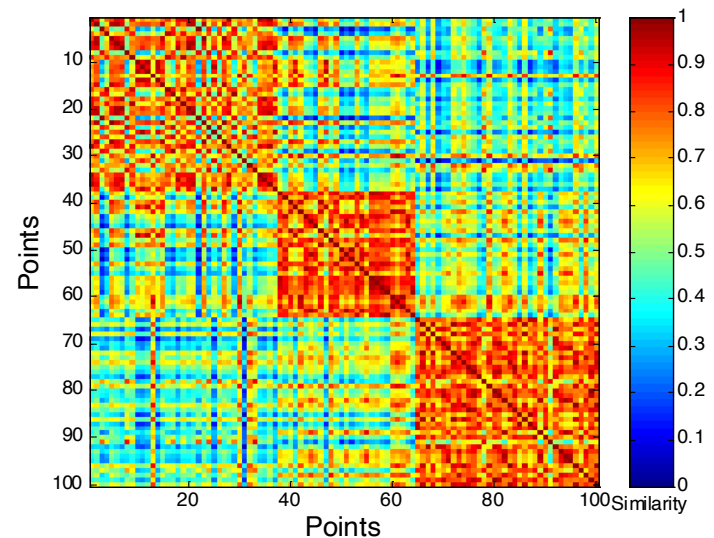
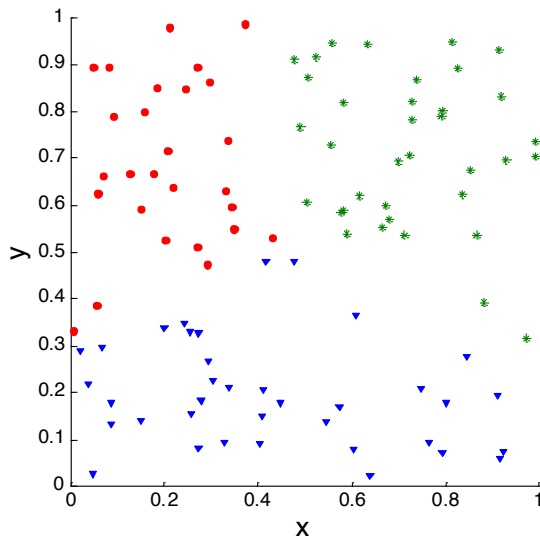
**Corr = -0.5810**

# Using Similarity Matrix for Cluster Validation

- Order the similarity matrix with respect to cluster labels and inspect visually.

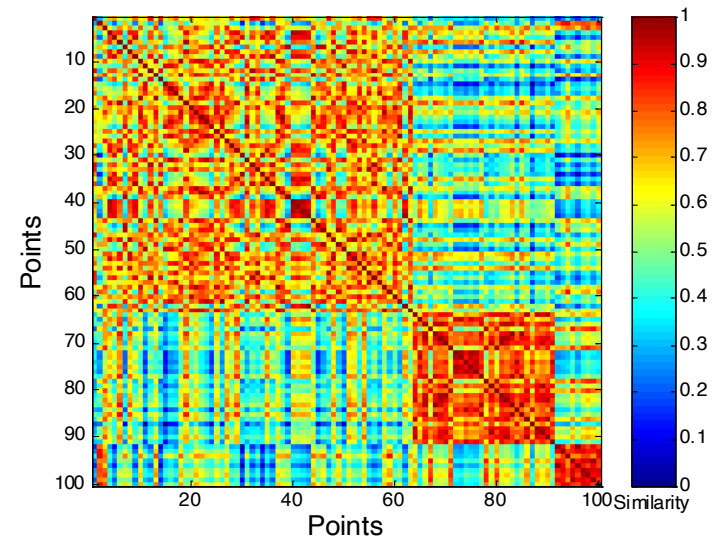
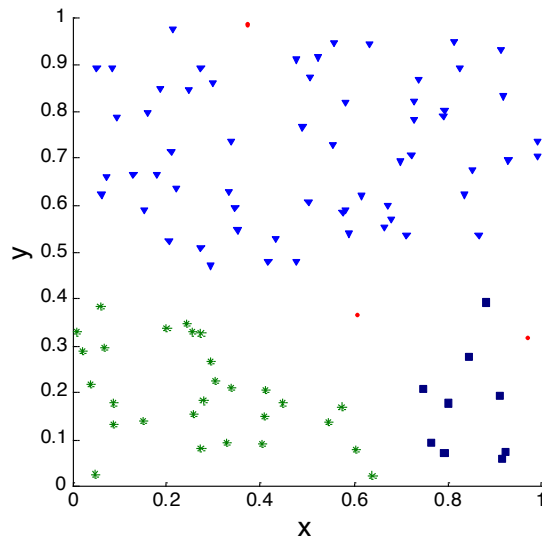


# Using Similarity Matrix for Cluster Validation



**K-means**

# Using Similarity Matrix for Cluster Validation



**DBSCAN**



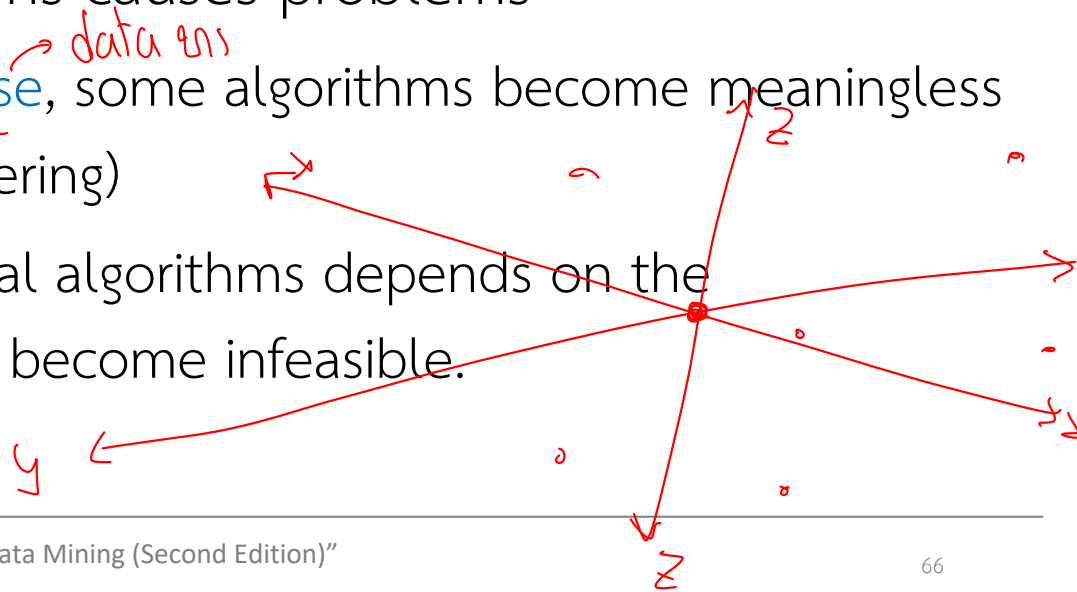
# Dimensionality Reduction

ရွှေ့ပြောင်း feature ခဏ



# The curse of dimensionality

- Real data usually have **thousands**, or **millions** of dimensions
  - E.g., web documents, where the dimensionality is the vocabulary of words
  - Facebook graph, where the dimensionality is the number of users
- Huge number of dimensions causes problems
  - Data becomes very **sparse**, some algorithms become meaningless (e.g. density based clustering)
  - The **complexity** of several algorithms depends on the dimensionality and they become infeasible.



# Dimensionality Reduction

---

- Usually the data can be described with fewer dimensions, without losing much of the meaning of the data.
  - The data **reside** in a space of lower dimensionality
- Essentially, we assume that some of the data is noise, and we can approximate the useful part with a lower dimensionality space.
  - Dimensionality reduction does not just reduce the amount of data, it often brings out the **useful** part of the data

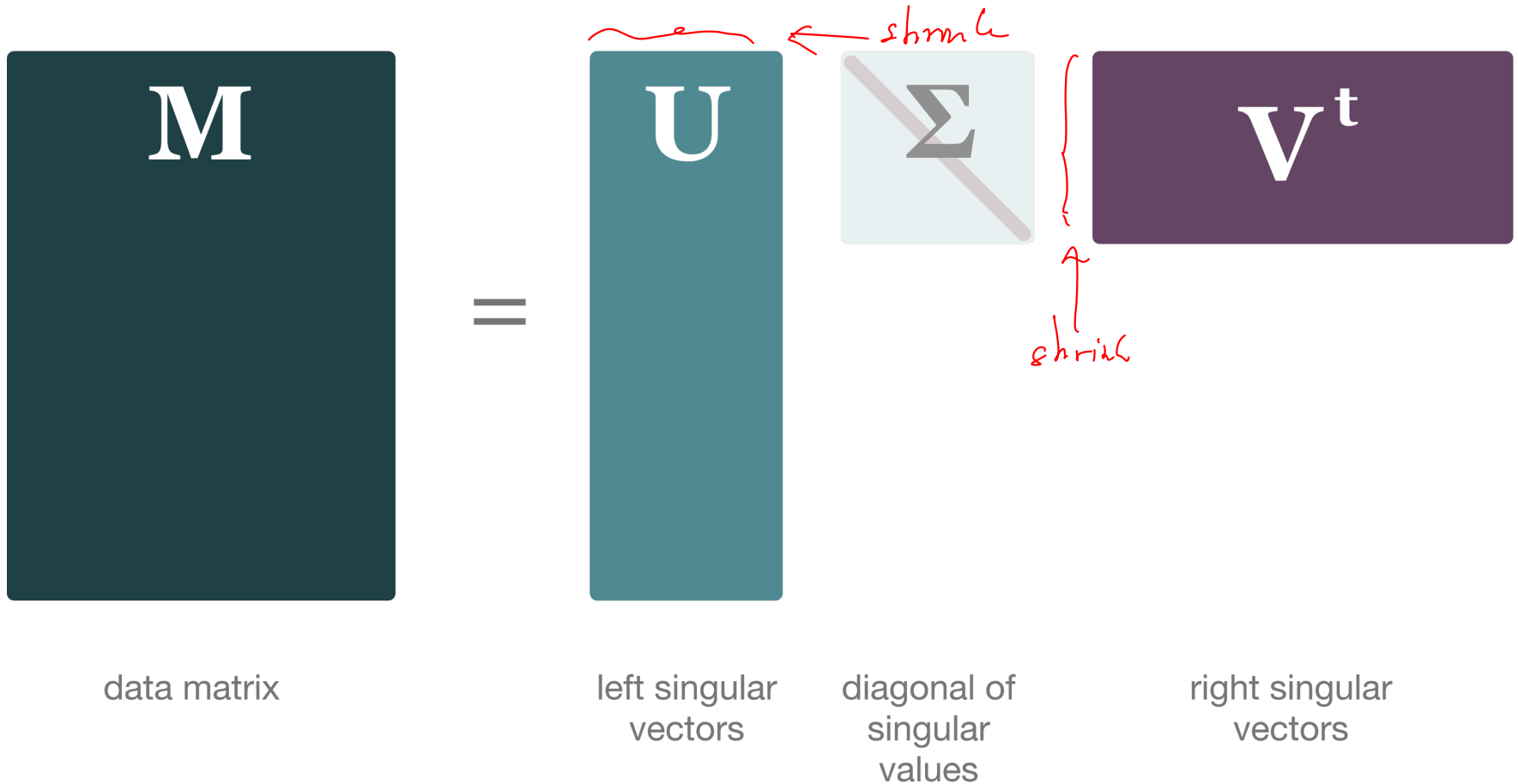
# Latent factor model

---

- Rows (columns) are linear combinations of  $k$  latent factors
  - E.g., in our extreme document example there are two factors
- Some noise is added to this rank- $k$  matrix resulting in higher rank
- SVD retrieves the latent factors (hopefully).

# SVD (Singular Value Decomposition)

Handwritten note:  $\rightarrow$  matrix  $U$  and  $V$  are  $n \times n$



# SVD

- Example

จัด cluster ใกล้เคียง แล้วจัดที่น้อยลง

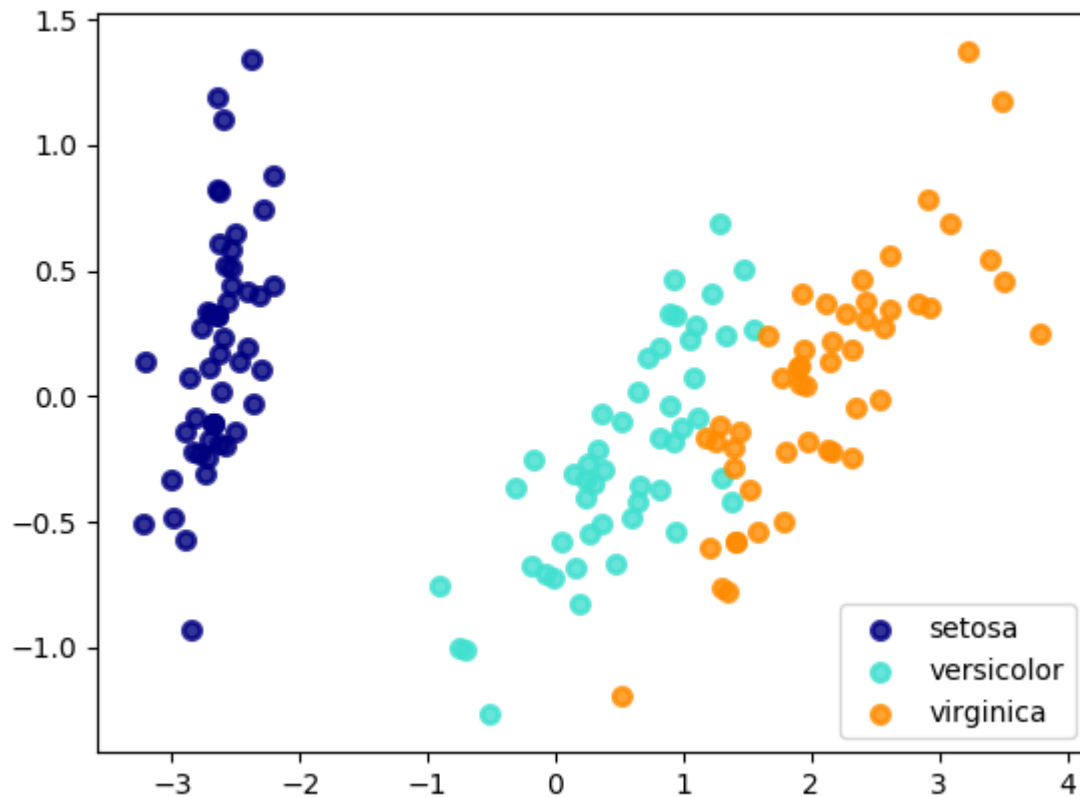
$$\begin{bmatrix} 1 & 1 & 1 & 0 & 0 \\ 2 & 2 & 2 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 \\ 5 & 5 & 5 & 0 & 0 \\ 0 & 0 & 0 & 2 & 2 \\ 0 & 0 & 0 & 3 & 3 \\ 0 & 0 & 0 & 1 & 1 \end{bmatrix} = \begin{bmatrix} 0.18 & 0 \\ 0.36 & 0 \\ 0.18 & 0 \\ 0.90 & 0 \\ 0 & 0.53 \\ 0 & 0.80 \\ 0 & 0.27 \end{bmatrix} \times \begin{bmatrix} 9.64 & 0 \\ 0 & 5.29 \end{bmatrix} \times \begin{bmatrix} 0.58 & 0.58 & 0.58 & 0 & 0 \\ 0 & 0 & 0 & 0.71 & 0.71 \end{bmatrix}$$

↓  
ของบิต

# Plot

---

- Dimensionality Reduction of the Iris Dataset



# Python

กำลังจะไปดูแถวๆ เหนือ เหนือๆ กันแล้ว

```
>>> import numpy as np
>>> from sklearn.decomposition import PCA
>>> X = np.array([[-1, -1], [-2, -1], [-3, -2], [1, 1], [2, 1], [3, 2]])
>>> pca = PCA(n_components=2)
>>> pca.fit(X)
PCA(copy=True, iterated_power='auto', n_components=2, random_state=None,
      svd_solver='auto', tol=0.0, whiten=False)
>>> print(pca.explained_variance_ratio_)
[ 0.99244...  0.00755...]
>>> print(pca.singular_values_)
[ 6.30061...  0.54980...]
```



“

Thanks to big data, machines can now be  
programmed to the next thing right.  
But only humans can do the next right thing.

”

Dov Seidman