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<https://github.com/pvateekul/ieat2026>



## Clustering

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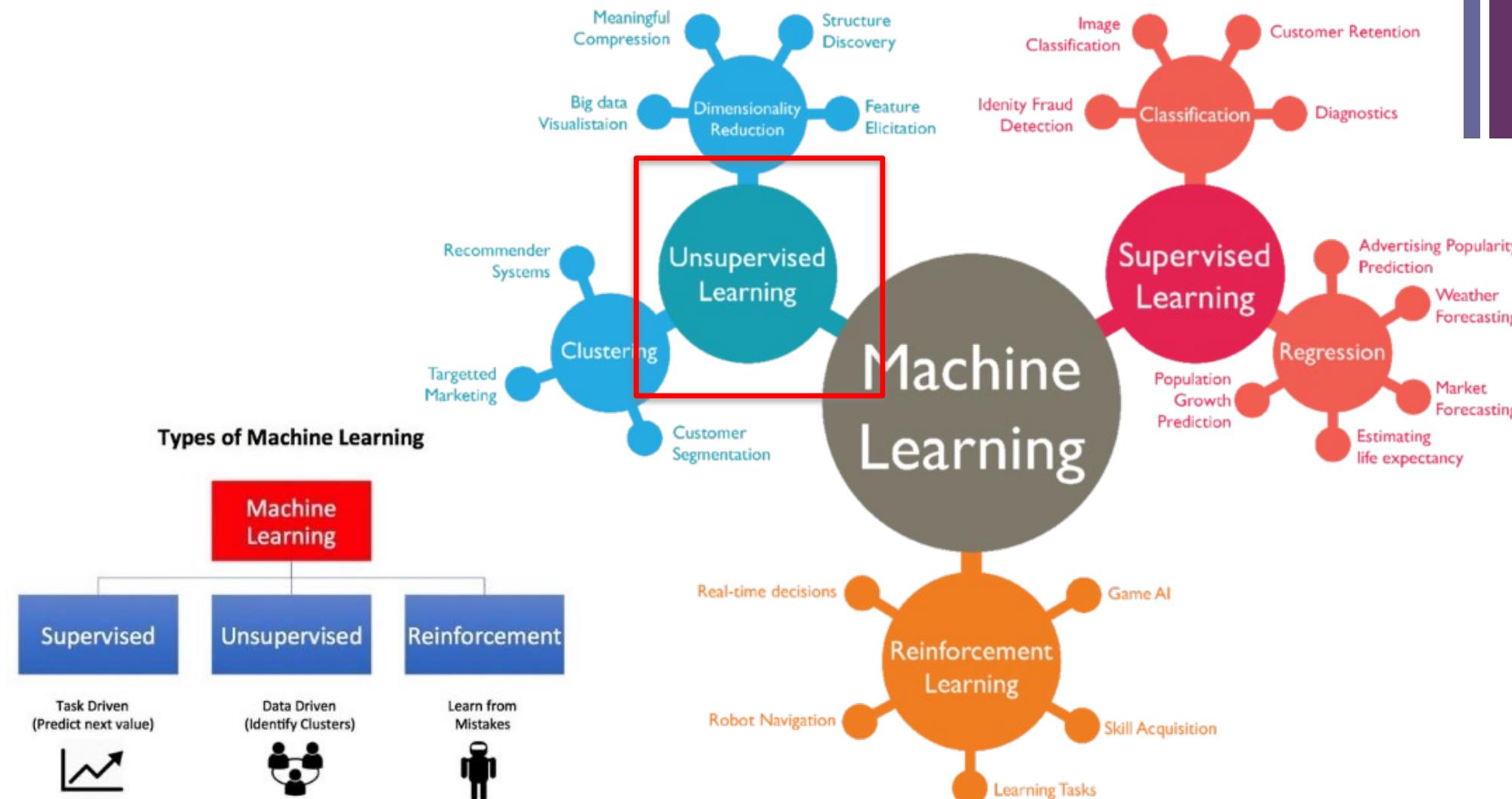
# Outlines

- Clustering
- LAB 4: Clustering



## Unsupervised Learning

# + Machine Learning (cont.)





## Task2: Unsupervised learning (descriptive task)

Training Data



Age	Income	Gender	Province	Purchase
25	25,000	Female	Bangkok	Yes
35	50,000	Female	Nontaburi	Yes
32	35,000	Male	Bangkok	No



INSIGHT

Testing Data



Age	Income	Gender	Province	Purchase
25	25,000	Female	Bangkok	?



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# Clustering



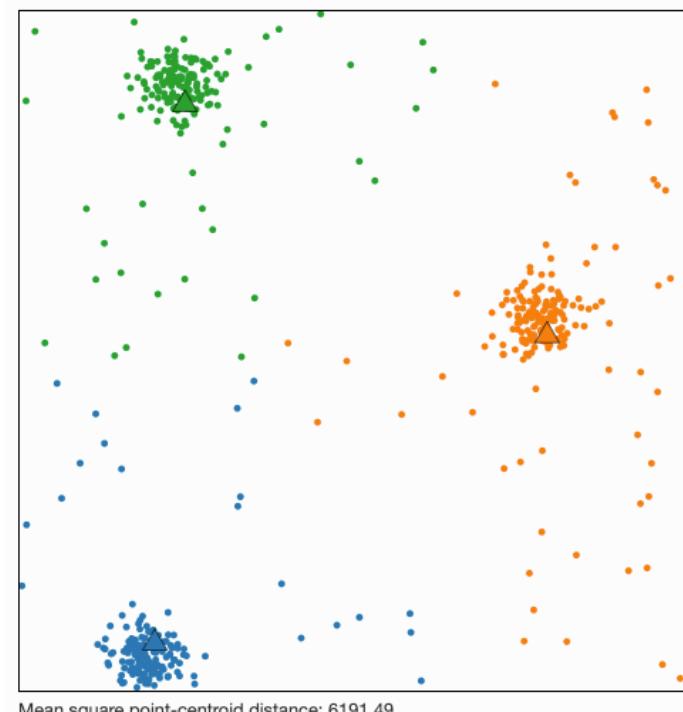
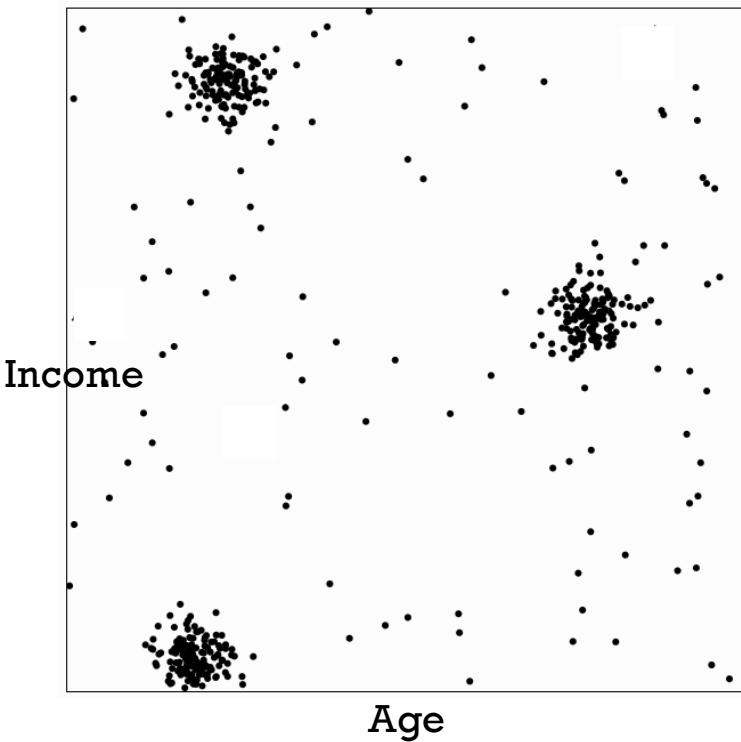
- In our class, there are many participants. Should we teach them using the same method?
- **May be not!** Since they may have different learning behaviors and backgrounds.
- Inputs
  - Education field
  - Level of English communication
  - Level of computer skills
  - Age range
  - Gender



# K-means Clustering

- <http://web.stanford.edu/class/ee103/visualizations/kmeans/kmeans.html>

## Visualizing K-Means Clustering



The  $k$ -means algorithm is an iterative method for clustering a set of  $N$  points (vectors) into  $k$  groups or clusters of points.

### Algorithm

Repeat until convergence:

#### Find closest centroid

Find the closest centroid to each point, and group points that share the same closest centroid.

#### Update centroid

Update each centroid to be the mean of the points in its group.

[Update centroid](#)

### Data

Clustered points  Random

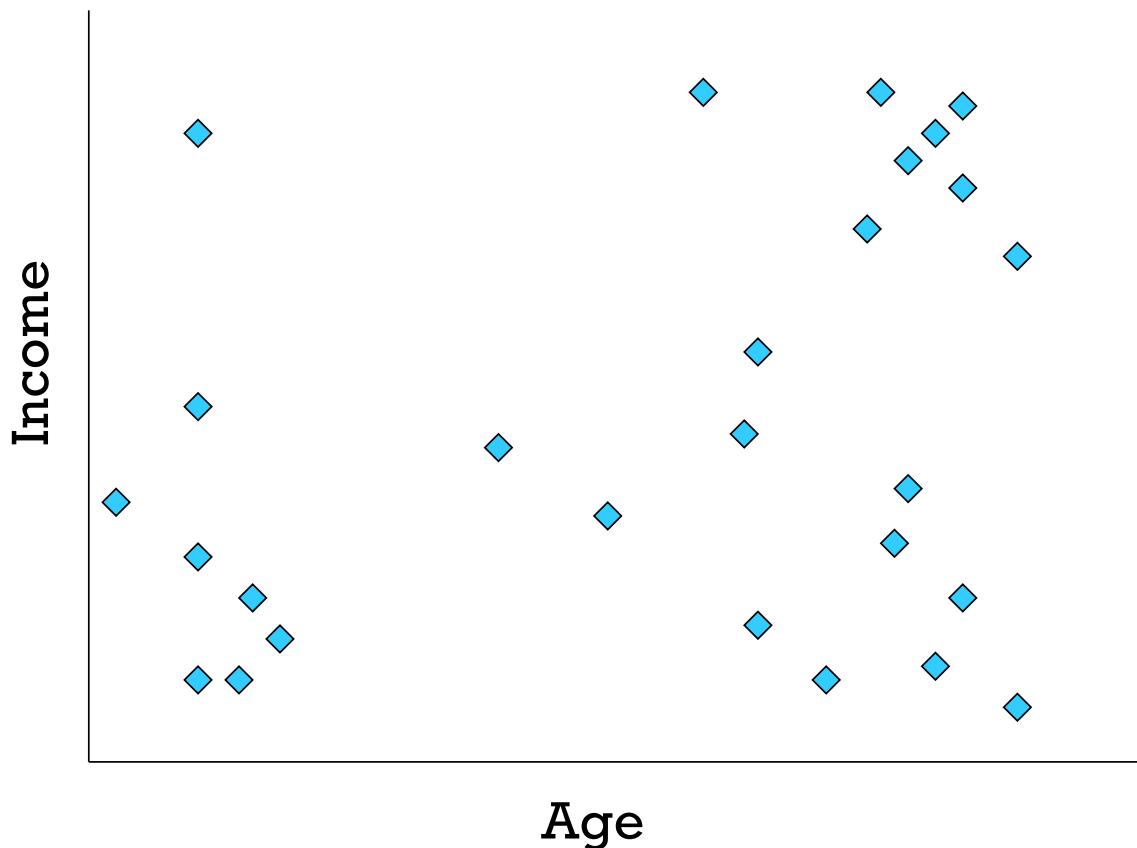
Number of clusters : 3

Number of centroids: 3

[New points](#)

[New centroids](#)

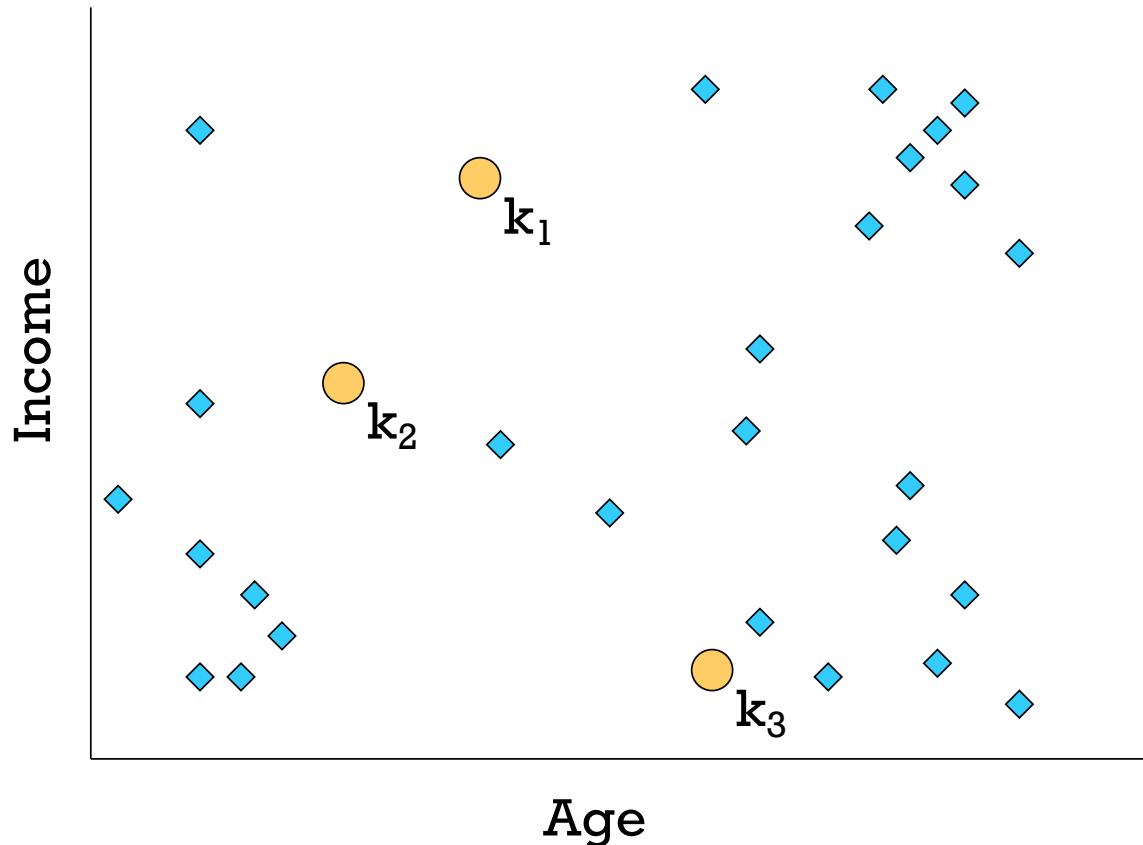
## + K-means: Step0





## K-means: Step 1

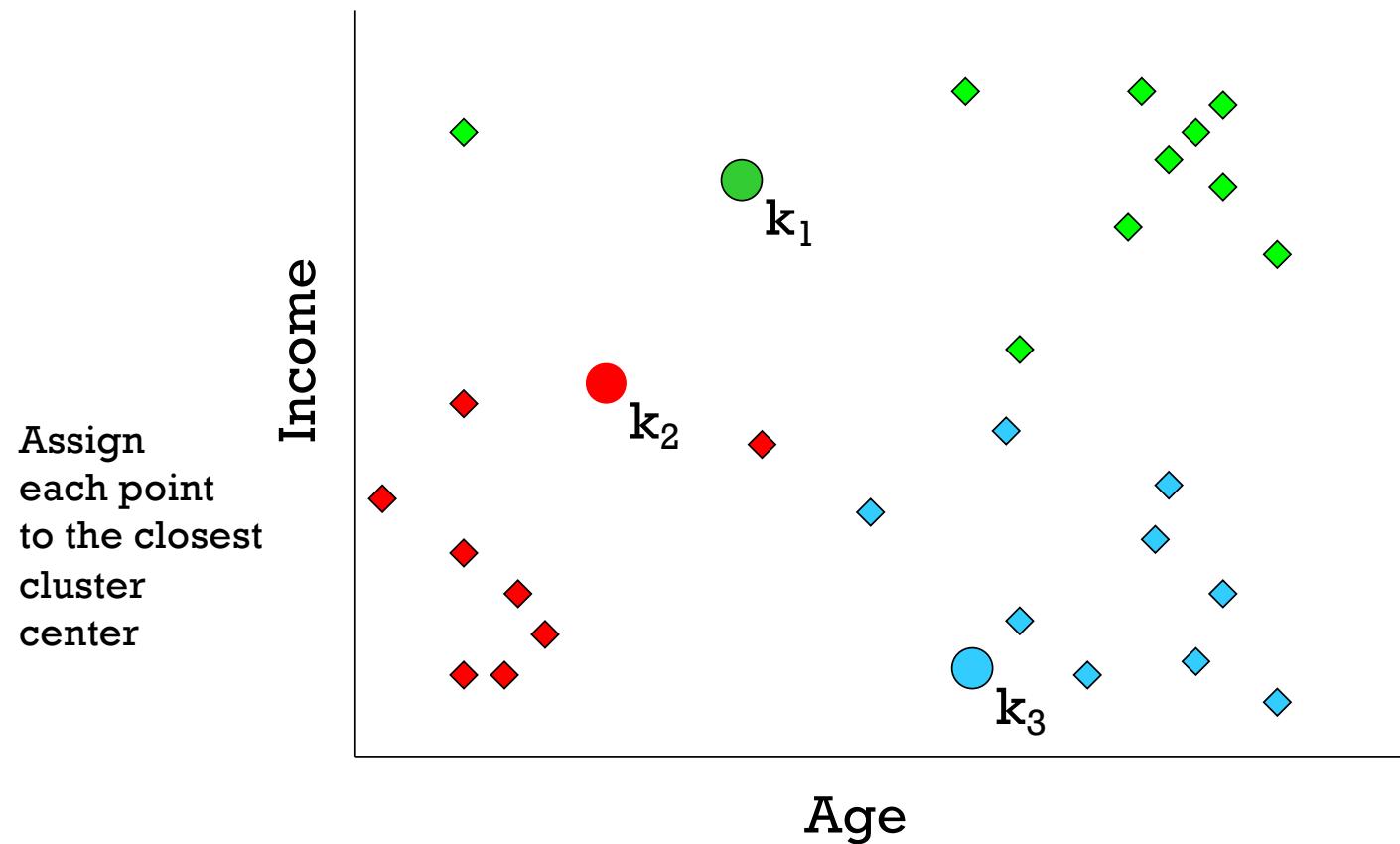
Pick 3  
initial  
cluster  
centers  
(randomly)





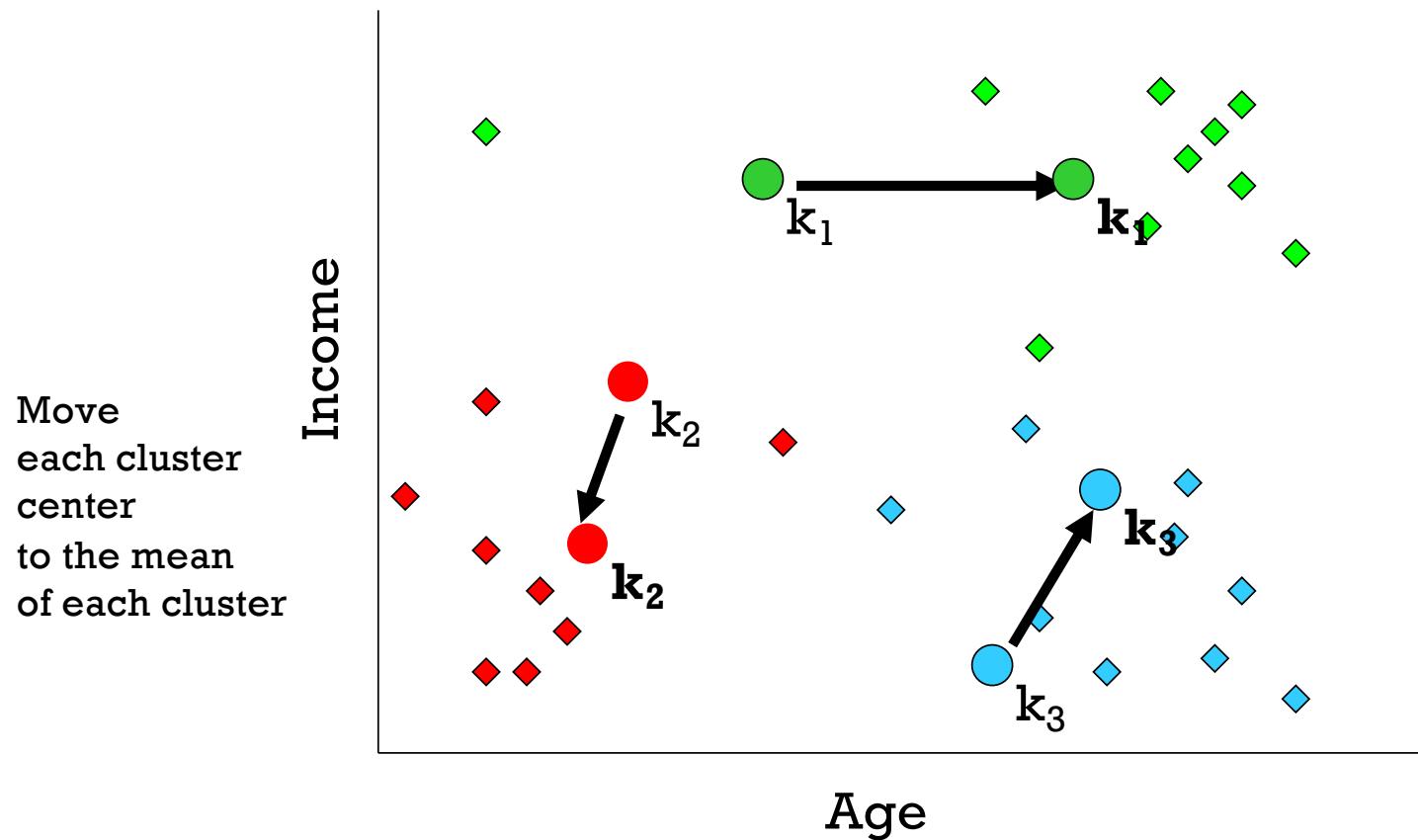
## K-means: Step2

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## K-means: Step3

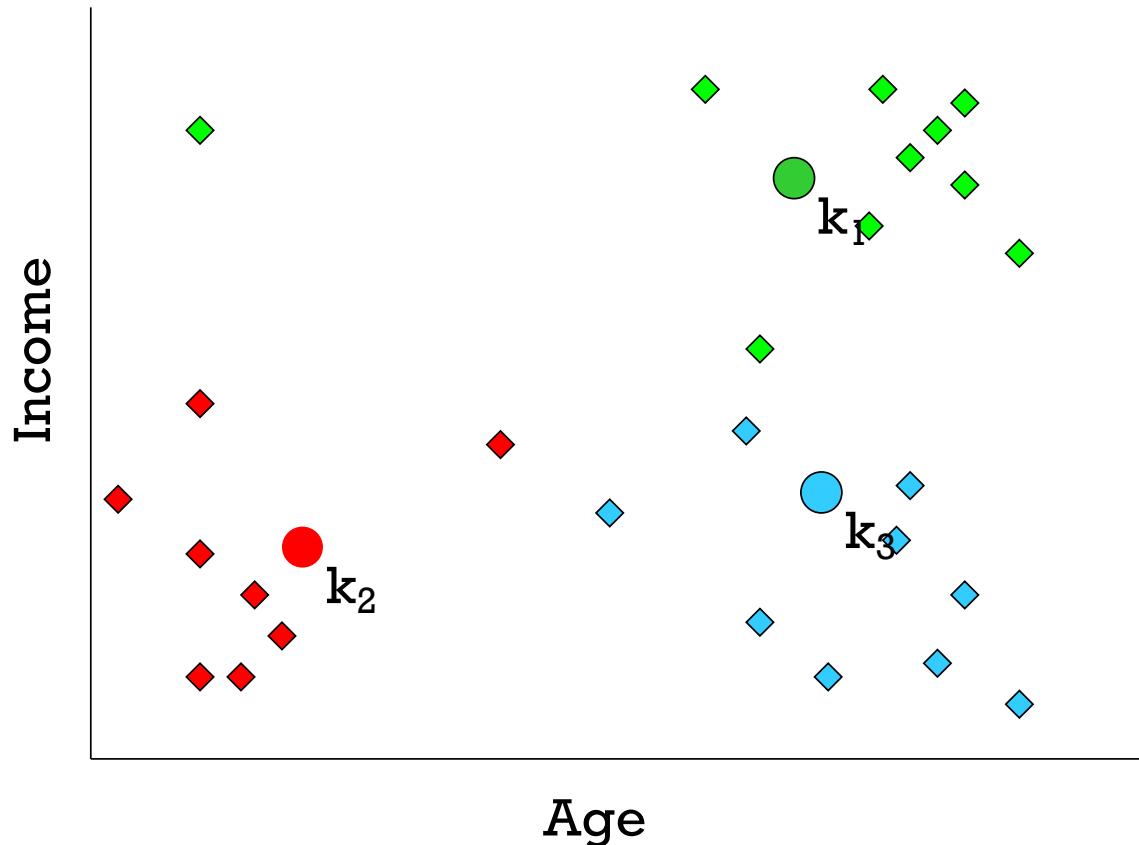




## K-means: Step4

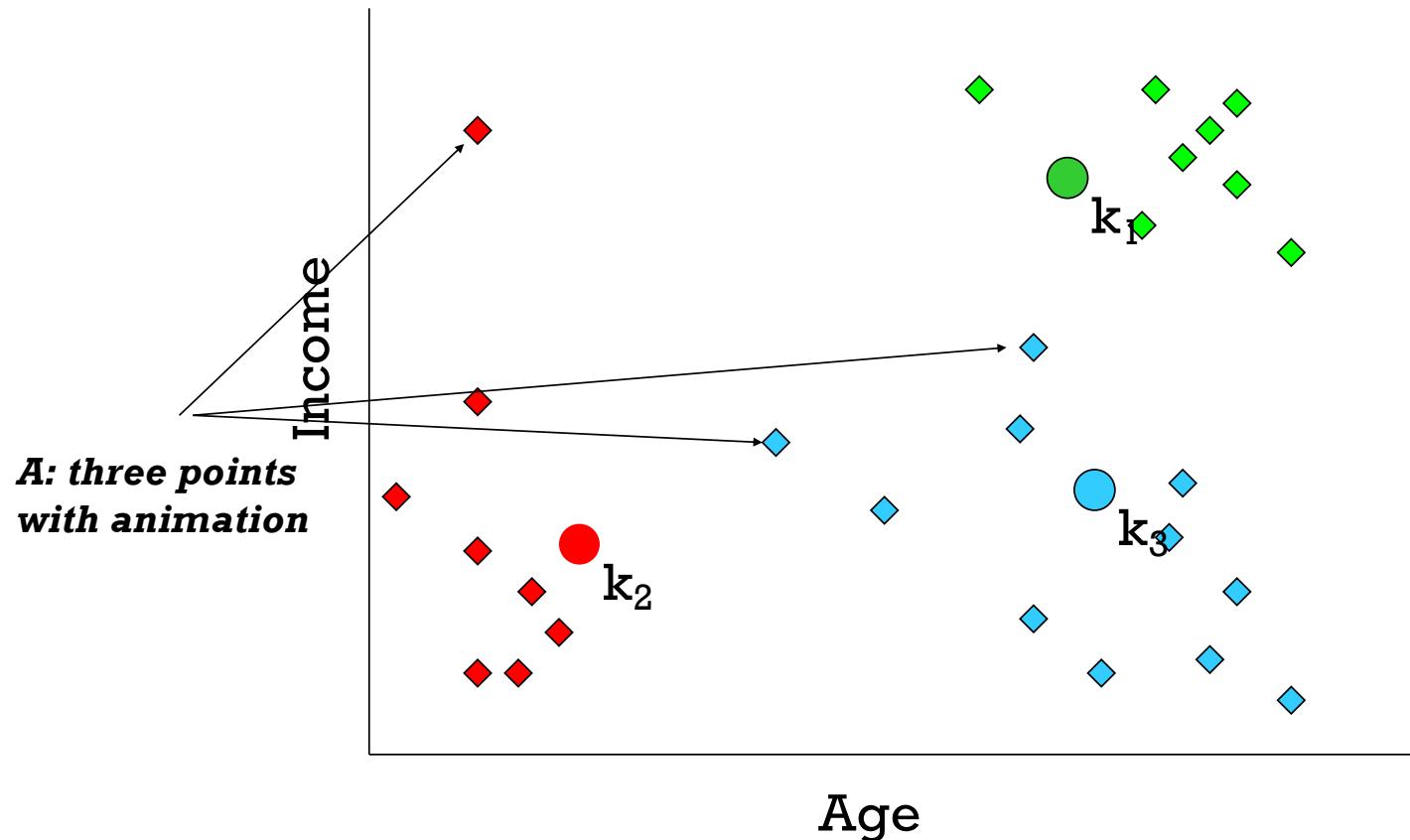
Reassign  
points  
closest to a  
different new  
cluster center

***Q: Which points  
are reassigned?***



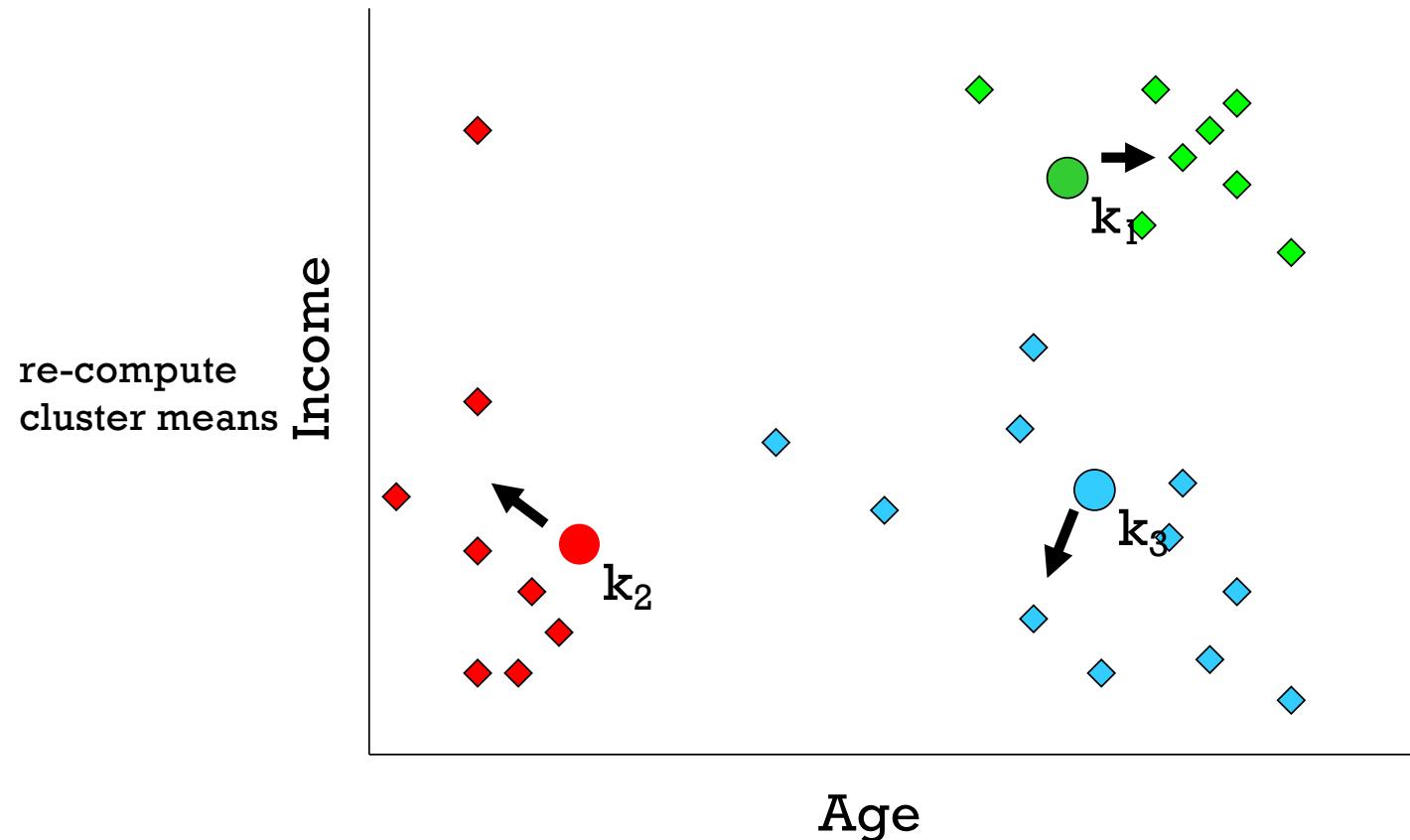


## K-means: Step4(a)





## K-means: Step5





## K-means: Step5(a)

### Cautions:

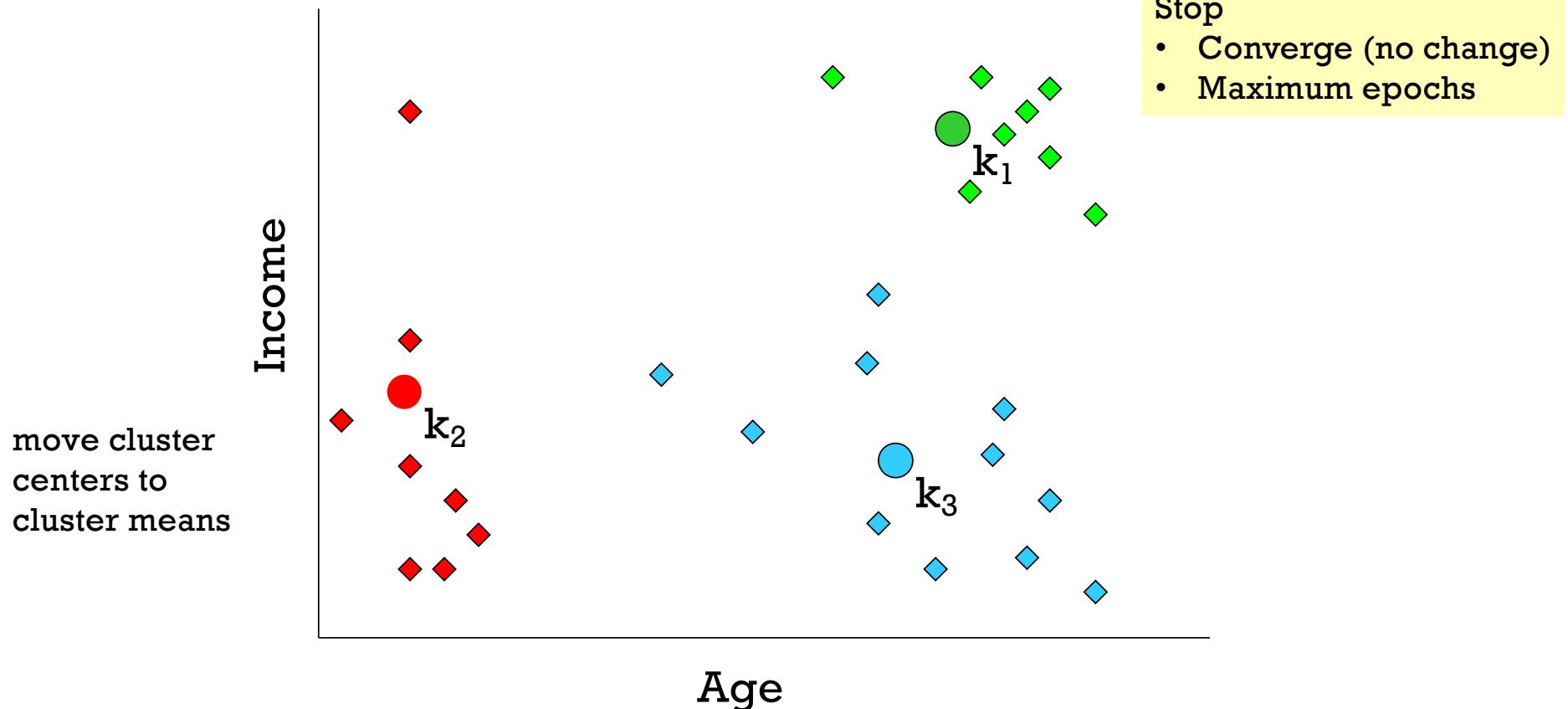
- Support only numerical variables
- Need to adjust variable range

### Important Params:

- $k$ , Distance function
- Maximum epochs

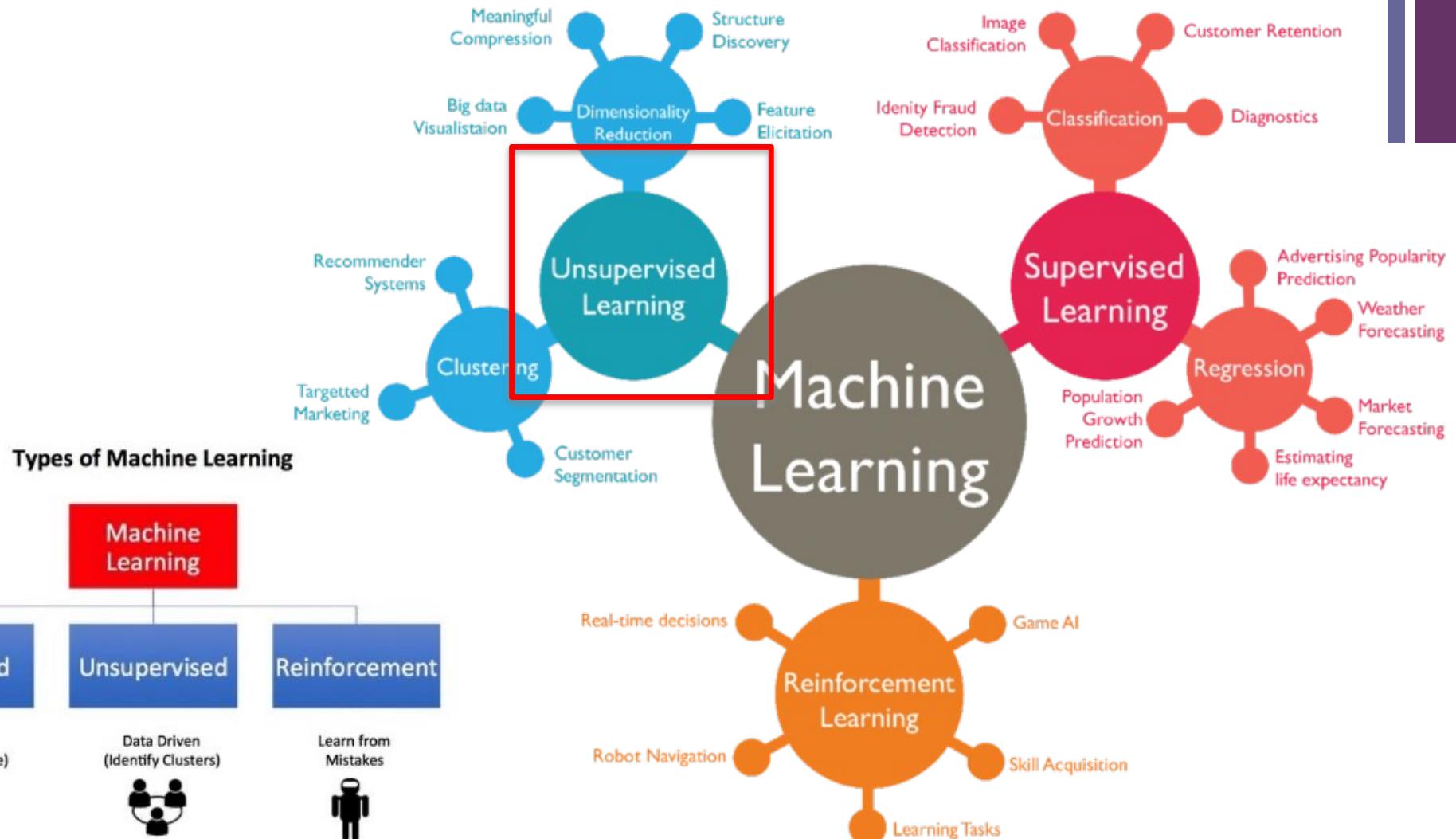
### Stop

- Converge (no change)
- Maximum epochs



# + Machine Learning

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scikit-learn 1.1.2

Other versions

Please [cite us](#) if you use the software.

sklearn.metrics.silhouette\_score  
Examples using  
sklearn.metrics.silhouette\_score

For the whole cluster

## sklearn.metrics.silhouette\_score

```
sklearn.metrics.silhouette_score(X, labels, *, metric='euclidean', sample_size=None, random_state=None, **kwds)
```

[\[source\]](#)

Compute the mean Silhouette Coefficient of all samples.

The Silhouette Coefficient is calculated using the mean intra-cluster distance (`a`) and the mean nearest-cluster distance (`b`) for each sample. The Silhouette Coefficient for a sample is  $(b - a) / \max(a, b)$ . To clarify, `b` is the distance between a sample and the nearest cluster that the sample is not a part of. Note that Silhouette Coefficient is only defined if number of labels is  $2 \leq n\_labels \leq n\_samples - 1$ .

This function returns the mean Silhouette Coefficient over all samples. To obtain the values for each sample, use `silhouette_samples`.

The best value is 1 and the worst value is -1. Values near 0 indicate overlapping clusters. Negative values generally indicate that a sample has been assigned to the wrong cluster, as a different cluster is more similar.

Read more in the [User Guide](#).

**Parameters:** `X : array-like of shape (n_samples_a, n_samples_a) if metric == "precomputed" or (n_samples_a, n_features) otherwise`

An array of pairwise distances between samples, or a feature array.

`labels : array-like of shape (n_samples,)`

Predicted labels for each sample.

`metric : str or callable, default='euclidean'`

The metric to use when calculating distance between instances in a feature array. If metric is a string, it must be one of the options allowed by `metrics.pairwise.pairwise_distances`. If `X` is the distance array itself, use `metric="precomputed"`.

`sample_size : int, default=None`

The size of the sample to use when computing the Silhouette Coefficient on a random subset of the data. If `sample_size is None`, no sampling is used.

[https://scikit-learn.org/stable/modules/generated/sklearn.metrics.silhouette\\_score.html](https://scikit-learn.org/stable/modules/generated/sklearn.metrics.silhouette_score.html)

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scikit-learn 1.3.0

Other versions

Please [cite us](#) if you use the software.[sklearn.metrics.silhouette\\_samples](#)[silhouette\\_samples](#)

Examples using

[sklearn.metrics.silhouette\\_samp](#)

## sklearn.metrics.silhouette\_samples

```
sklearn.metrics.silhouette_samples(X, labels, *, metric='euclidean', **kwds)
```

[\[source\]](#)

Compute the Silhouette Coefficient for each sample.

The Silhouette Coefficient is a measure of how well samples are clustered with samples that are similar to themselves. Clustering models with a high Silhouette Coefficient are said to be dense, where samples in the same cluster are similar to each other, and well separated, where samples in different clusters are not very similar to each other.

The Silhouette Coefficient is calculated using the mean intra-cluster distance (`a`) and the mean nearest-cluster distance (`b`) for each sample. The Silhouette Coefficient for a sample is  $(b - a) / \max(a, b)$ . Note that Silhouette Coefficient is only defined if number of labels is  $2 \leq n\_labels \leq n\_samples - 1$ .

This function returns the Silhouette Coefficient for each sample.

The best value is 1 and the worst value is -1. Values near 0 indicate overlapping clusters.

Read more in the [User Guide](#).

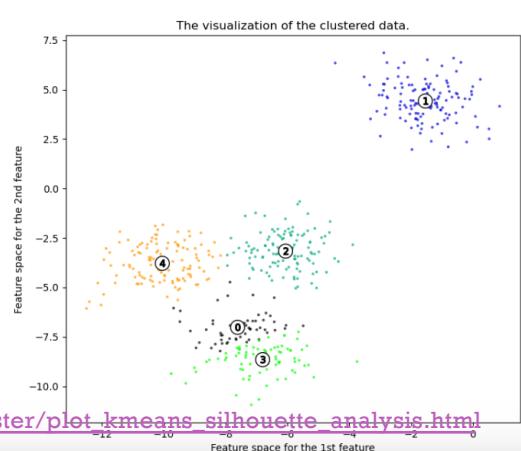
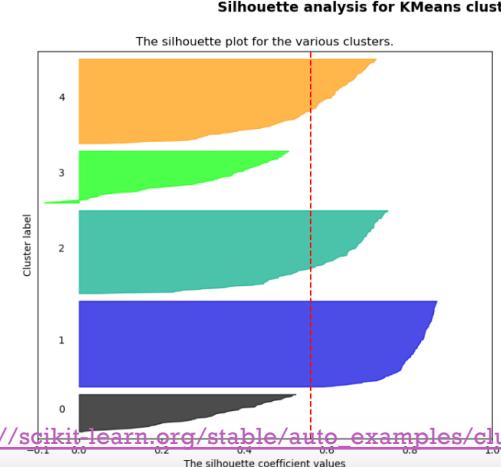
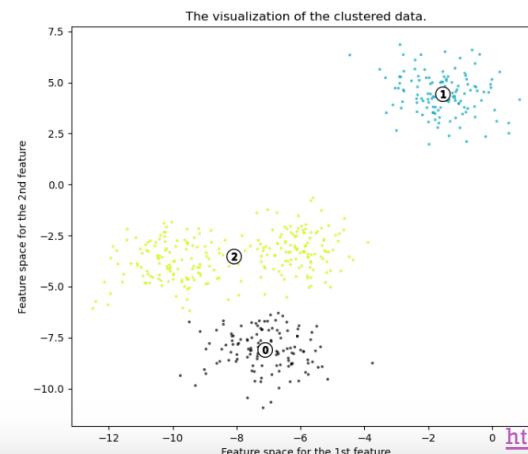
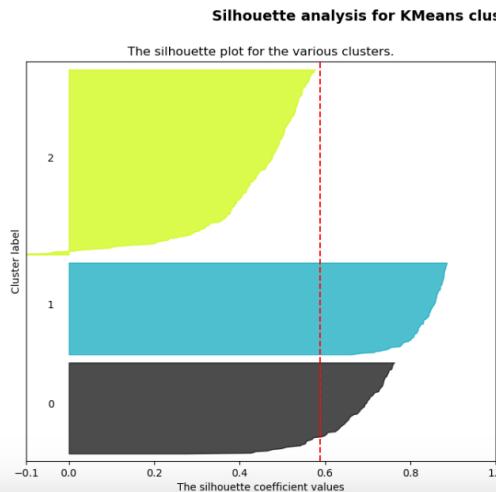
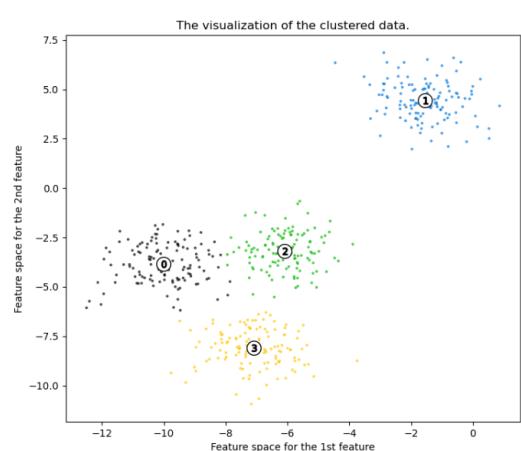
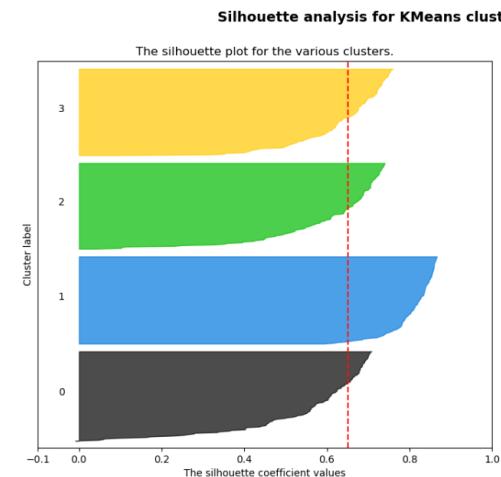
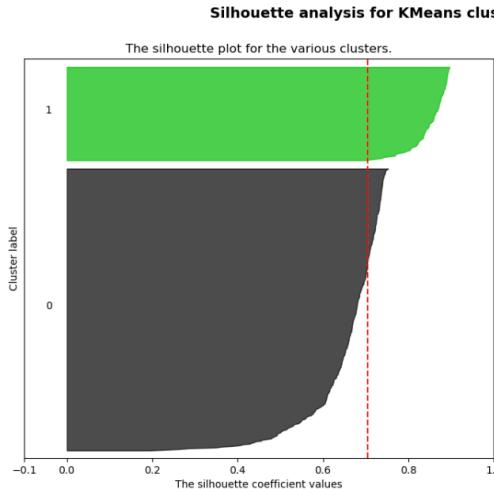
**Parameters:** `X : {array-like, sparse matrix} of shape (n_samples_a, n_samples_a) if metric == "precomputed" or (n_samples_a, n_features) otherwise`

An array of pairwise distances between samples, or a feature array. If a sparse matrix is provided, CSR format should be favoured avoiding an additional copy.

# How to choose n\_clusters? Chosen is 2 or 4.

Out:

For n\_clusters = 2 The average silhouette\_score is : 0.7049787496083262  
For n\_clusters = 3 The average silhouette\_score is : 0.5882004012129721  
For n\_clusters = 4 The average silhouette\_score is : 0.6505186632729437  
For n\_clusters = 5 The average silhouette\_score is : 0.561464362648773  
For n\_clusters = 6 The average silhouette\_score is : 0.4857596147013469



[https://scikit-learn.org/stable/auto\\_examples/cluster/plot\\_kmeans\\_silhouette\\_analysis.html](https://scikit-learn.org/stable/auto_examples/cluster/plot_kmeans_silhouette_analysis.html)



## LAB 4: Clustering

# LAB 4: Clustering

## Data Description

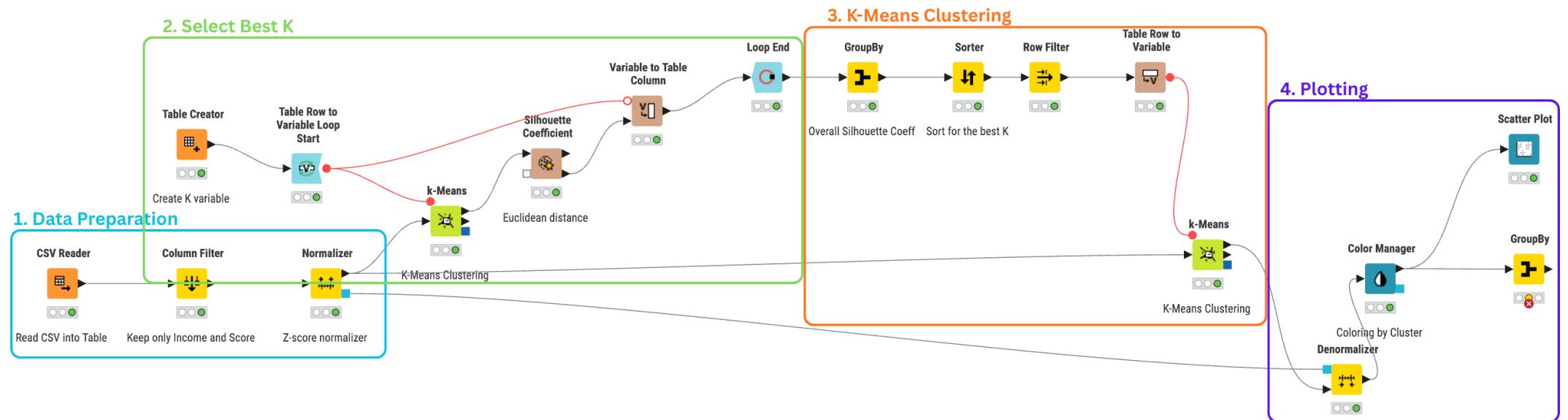
- The Mall Customer dataset contains 200 customers. Each row is one subject. The goal is to predict the Spending Class (High or Low Spender) of the customer.
- Target Variable: High Spender: “yes” if spending score  $\geq 60$ ; else “no”

## Input:

- CutomerID
- Gender: Male/Female
- Age: (years)
- Annual Income: (k\$)
- Spending Score: Derived Metric (0-100)



# LAB 4: Clustering



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Thank you & any questions