

COMP 2211 Exploring Artificial Intelligence K-Means Clustering Dr. Desmond Tsoi

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Problem

• Given the following simplified Mall customers dataset with attributes, age, income (in thousands) and expense score (1-100).

Person	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Age	19	67	35	60	65	49	70	70	57	68	23	65	27	47	57	43	56	40	37	34
1 Income $\times 1000$	15	19	24	30	38	42	46	49	54	59	62	63	67	71	75	78	79	87	97	103
Expense score 1-100	39	14	35	4	35	52	56	55	51	55	41	52	56	9	5	17	35	13	32	23

Note: Expense score goes from 1 (low spends) to 100 (high spends).

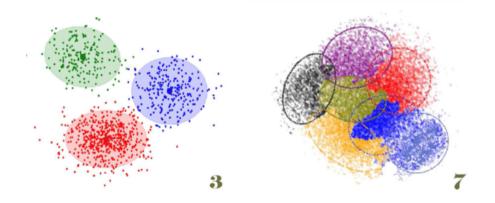
• Could we cluster the customers based on their characteristics (age, income & expense score) to determine the targeted advertisement and make the marketing budget more efficient?





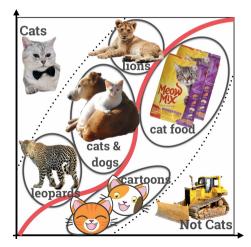
What is Clustering?

• Clustering is grouping things into "natural" categories when no class label is available.



Why Clustering?

- Labeling a large set of data samples can be costly.
- Clustering can be used for finding features that will be useful later for categorization.



What is Clustering for?

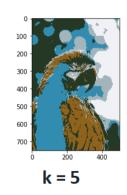
- Group people of similar sizes together to make "small", "medium", and "large" T-shirts.
 - Tailor-made for each person: too expensive
 - One-size-fits-all: Does not fit all
- Given a collection of text documents, we want to organize them according to their content similarities
 - To produce a topic hierarchy
- In marketing, segment customers according to their similarities
 - To do targeted marketing



What is Clustering for?

- Image segmentation
 - To partition a digital image into multiple image segments, also known as image regions.



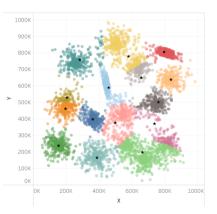




k = 20

K-Means Clustering

- K-Means clustering is an unsupervised learning algorithm.
- There is no labeled data for this clustering, unlike supervised learning.
- It performs the division of data into non-overlapping clusters that share similarities and are dissimilar to the data belonging to another cluster.
- The term 'K' is a number telling the system how many clusters you need to create. For example, if K = 3 refers to three clusters.



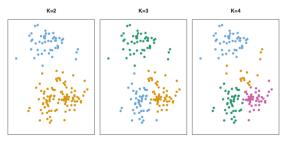
K-Means Clustering

• Let the set of data points

$$\mathsf{D} = \{\mathsf{x}_1, \mathsf{x}_2, \mathsf{x}_3, \dots, \mathsf{x}_n\}$$

where $\mathbf{x_i} = (x_{i1}, x_{i2}, \dots, x_{id})$ is the *i*th data point, and *d* is the number of dimensions.

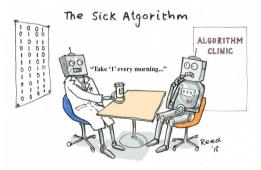
- To perform K-means clustering, we must first specify the desired number of clusters K.
- The K-Means algorithm partitions the given data points into K non-overlapped clusters:
 - Each cluster has a cluster center, called centroid



K-means Algorithm

Given K, the K-Means algorithm works as follows:

- 1. Choose K (random) data points (seeds) to be the initial centroids (cluster centers)
- 2. Find the distances between each data point in our training set with the K centroids
- 3. Assign each data point to the closest centroid according to the distance found
- 4. Re-compute the centroids using the current cluster memberships
- 5. If a convergence criterion is NOT met, repeat steps 2 to 4



Example

• Let's perform K-Means Clustering on the following simplified Mall customers dataset with attributes, age, income (in thousands) and expense score (1-100).

Person	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Age	19	67	35	60	65	49	70	70	57	68	23	65	27	47	57	43	56	40	37	34
1 Income $\times 1000$	15	19	24	30	38	42	46	49	54	59	62	63	67	71	75	78	79	87	97	103
Expense score 1-100	39	14	35	4	35	52	56	55	51	55	41	52	56	9	5	17	35	13	32	23

Note: Expense score goes from 1 (low spends) to 100 (high spends).





Step 1: Randomly Pick 3 Data Points as Initial Centroids

Centroids . Customer Segmentation - K-Means Clustering 50 (1-100) 40 Score (30 Expe 20 10 100 80 nelth 20 60 30 40 50 60 20 70

- 1st centroid: (70, 46, 56)
- 2nd centroid: (27, 67, 56)
- 3rd centroid: (37, 97, 32)

Step 2: Find the Distances Between Each Data Point with the 3 Centroids

- 1st centroid: (70, 46, 56)
- 2nd centroid: (27, 67, 56)
- 3rd centroid: (37, 97, 32)

Assume Euclidean distance is used.

Person	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Age	19	67	35	60	65	49	70	70	57	68	23	65	27	47	57	43	56	40	37	34
Income $\times 1000$	15	19	24	30	38	42	46	49	54	59	62	63	67	71	75	78	79	87	97	103
Expense																				
score 1-100	39	14	35	4	35	52	56	55	51	55	41	52	56	9	5	17	35	13	32	23
DC1	62	50	46	55	23	22	0	3	16	13	52	18	48	58	60	57	42	67	65	75
DC2	55	75	49	72	52	34	48	47	33	42	16	38	0	51	60	44	38	49	40	49
DC3	84	86	73	76	65	60	65	63	51	54	39	48	40	36	40	25	26	22	0	11

where DC1, DC2, DC3 are the distances between the data points and 1st centroid, 2nd centroid, and 3rd centroid, respectively.

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Step 3: Assign Each Data Point to the Closest Centroid

- 1st centroid: (70, 46, 56)
- 2nd centroid: (27, 67, 56)
- 3rd centroid: (37, 97, 32)

Person	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Age	19	67	35	60	65	49	70	70	57	68	23	65	27	47	57	43	56	40	37	34
Income $\times 1000$	15	19	24	30	38	42	46	49	54	59	62	63	67	71	75	78	79	87	97	103
Expense score 1-100	39	14	35	4	35	52	56	55	51	55	41	52	56	9	5	17	35	13	32	23
DC1	62	50	46	55	23	22	0	3	16	13	52	18	48	58	60	57	42	67	65	75
DC2	55	75	49	72	52	34	48	47	33	42	16	38	0	51	60	44	38	49	40	49
DC3	84	86	73	76	65	60	65	63	51	54	39	48	40	36	40	25	26	22	0	11
Cluster	2	1	1	1	1	1	1	1	1	1	2	1	2	3	3	3	3	3	3	3

where DC1, DC2, DC3 are the distances between the data points and 1st centroid, 2nd centroid, and 3rd centroid, respectively.

Step 4: Re-compute the Centroids Using the Current Cluster Memberships

Person	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
Age	19	67	35	60	65	49	70	70	57	68	23	65	27	47	57	43	56	40	37	34
Income $\times 1000$	15	19	24	30	38	42	46	49	54	59	62	63	67	71	75	78	79	87	97	103
Expense																				
score	39	14	35	4	35	52	56	55	51	55	41	52	56	9	5	17	35	13	32	23
1-100																				
DC1	62	50	46	55	23	22	0	3	16	13	52	18	48	58	60	57	42	67	65	75
DC2	55	75	49	72	52	34	48	47	33	42	16	38	0	51	60	44	38	49	40	49
DC3	84	86	73	76	65	60	65	63	51	54	39	48	40	36	40	25	26	22	0	11
Cluster	2	1	1	1	1	1	1	1	1	1	2	1	2	3	3	3	3	3	3	3

New 1st centroid:

 $\begin{aligned} x_1 &= (67+35+60+65+49+70+70+57+68+65)/10 = 60.6 \\ x_2 &= (19+24+30+38+42+46+49+54+59+63)/10 = 42.4 \\ x_3 &= (14+35+4+35+52+56+55+51+55+52)/10 = 40.9 \end{aligned}$

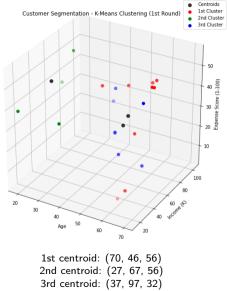
New 2nd centroid:

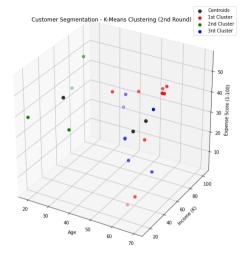
 $x_1 = (19 + 23 + 27)/3 = 23$ $x_2 = (15 + 62 + 67)/3 = 48$ $x_3 = (39 + 41 + 56)/3 = 45.33$

New 3rd centroid:

$x_1 = (47 + 57 + 43 + 56 + 40 + 37 + 34)/7 = 44.86$	
$x_2 = (71 + 75 + 78 + 79 + 87 + 97 + 103)/7 = 84.24$	9
$x_3 = (9 + 5 + 17 + 35 + 13 + 32 + 23)/7 = 19.14$	

The 1st and the 2nd Round

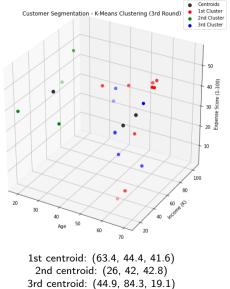


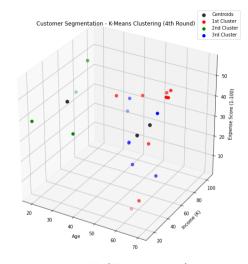


1st centroid: (60.6, 42.4, 40.9) 2nd centroid: (23, 48, 45.3) 3rd centroid: (44.9, 84.3, 19.1)

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The 3rd and the 4th Round





1st centroid: (63.4, 44.4, 41.6) 2nd centroid: (26, 42, 42.8) 3rd centroid: (44.9, 84.3, 19.1)

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Conclusion

• Cluster 1:

The average age is 63 years old, the average annual income is \$44K, and the average expense score is 42 of 100.

• Cluster 2:

The average age is 26 years old, the average annual income is 42K, and the average expense score is 43 of 100.

• Cluster 3:

The average age is 45 years old, the average annual income is \$84K, and the average expense score is 19 of 100.



K-Means Clustering Implementation using Scikit-Learn

Import the required libraries
import numpy as np
from sklearn.cluster import KMeans
import matplotlib.pyplot as plt

```
# Unlabled training data
data = np.array([[19, 15, 39], [67, 19, 14], [35, 24, 35], [60, 30, 4], [65, 38, 35],
            [49, 42, 52], [70, 46, 56], [70, 49, 55], [57, 54, 51], [68, 59, 55],
            [23, 62, 41], [65, 63, 52], [27, 67, 56], [47, 71, 9], [57, 75, 5],
            [43, 78, 17], [56, 79, 35], [40, 87, 13], [37, 97, 32], [34, 103, 23]])
```

Initial centroids
init_centroids = np.array([[70, 46, 56], [27, 67, 56], [37, 97, 32]])

```
# Create a KMeans object by specifying
# - Number of clusters (n_clusters) = 3, initial centroids (init) = init_centroids
# - Number of time the k-means algorithm will be run with different centroid seeds (n_init) = 1
# - Maximum number of iterations of the k-means algorithm for a single run (max_iter) = 4
kmeans = KMeans(n_clusters=3, init=init_centroids, n_init=1, max_iter = 4)
```

K-Means Clustering Implementation using Scikit-Learn

```
kmeans.fit(data)  # Compute k-means clustering
labels = kmeans.predict(data)  # Predict the closest cluster each sample in data belongs to
centroids = kmeans.cluster_centers_ # Get resulting centroids
fig = plt.figure(figsize = (10,10)) # Figure width = 10 inches, height = 10 inches
ax = fig.gca(projection='3d') # Defining 3D axes so that we can plot 3D data into it
```

Get boolean arrays representing entries with labels = 0, 1, and 2
a = np.array(labels == 0); b = np.array(labels == 1); c = np.array(labels == 2)

```
ax.set_title("Customer Segmentation - K-Means Clustering") # Put figure title
```

K-Means Stopping Criterion

- No/Minimum re-assignments of data points to different clusters, or
- No/Minimum change of centroids, or
- Minimum decrease in the sum of squared error (SSE) between successive iteration

$$SSE = \sum_{j=1}^{k} \sum_{\mathbf{x} \in C_j} dist(\mathbf{x}, \mathbf{m_j})^2$$

where

- C_j is the jth cluster
- **m**_j is the centroid of cluster *C*_j
- dist(x, m_j) is the distance between data point x and centroid m_j

Common Questions

1. How to choose K?

Answer: The following methods are used to find the optimal value of K for K-Means Clustering.

- Elbow method
- Silhouette method

These methods will be discussed in advanced level machine learning courses.

2. What distance metric should be used for K-Means?

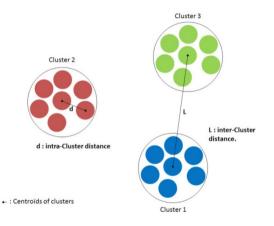
Answer: It depends on your data. Normally, K-Means uses Euclidean distance.



Clustering Quality

- High quality clustering
 - Maximizes inter-clusters distance (Isolation)
 - (i.e., distance between clusters)
 - Minimizes intra-clusters distance (Compactness)

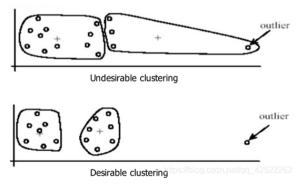
(i.e., distance between data points in the same cluster) $\label{eq:constraint}$



The quality of a clustering result depends on the algorithm, the distance function, and the application.

Weaknesses of K-Means

- It 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
 - Desirable and undesirable clustering with outliers



How to Deal with Outliers?

- Remove some data points in the clustering process that are much further away from the centroids than other data points.
 - To be safe, we may want to monitor these possible outliers over a few iterations and then decide to remove them.
- Perform random sampling. Since in sampling, we only choose a small subset of the data points, the chance of select an outlier is very small.
 - Assign the rest of the data points to the clusters by distance or similarity comparison, or classification.



Weaknesses of K-Means

• The algorithm is sensitive to initial seeds.

If we use seeds that are not-so-good: not-so-good results



(A). Random selection of seeds (centroids)



(B). Iteration 1



(C). Iteration 2

If we use different seeds: good results



(A). Random selection of k seeds (centroids)



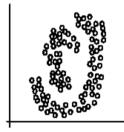
(B). Iteration 1



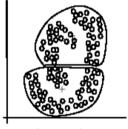
(C). Iteration 2

Weaknesses of K-Means

• The K-Means algorithm is not suitable for discovering clusters that are not hyper-ellipsoids (or hyper-spheres).



Two natural clusters



k-means clusters

Pros and Cons

• Pros:

- Easy to understand and implement
- It is efficient, given K and the number of iterations is small
- Cons:
 - It is only applicable if the mean is defined.
 - For categorical data, K-mode is used, i.e., the centroid is represent by most frequent values
 - $\bullet\,$ The user needs to specify K
 - It is sensitive to outliers



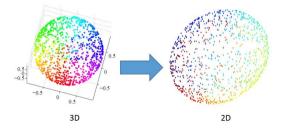
Summary

- Despite weaknesses, K-Means is still the most popular algorithm due to its simplicity, efficiency and other clustering algorithms have their own lists of weaknesses.
- No clear evidence that any other clustering algorithm performs better in general, although they may be more suitable for some specific types of data or applications.
- Comparing different clustering algorithms is a difficult task. No one knows the correct clusters.



Remark: Dimension Reduction using PCA

- Principal Component Analysis (PCA) is one of the easiest, most intuitive and most frequently used methods for dimensionality reduction, projecting data onto its orthogonal feature subspace.
 - The main idea of PCA is to reduce the dimensionality of a data set consisting of many variables correlated with each other, while retaining the variation present in the dataset, up to the maximum extent.
- It is a common practice to apply PCA before K-Means clustering is performed.
- It is believed that it improves the clustering results in practice (noise reduction).



Practice Problem

- Given 4 types of medicines and each has two attributes (weight and pH index).
- Group these medicines into K = 2 group using K-Means clustering.

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4

• Use (1, 1) and (2, 1) as the initial centroids (i.e., seeds) and performing K-Means Clustering until there is no re-assignments of data points to different clusters.



Step 1: Use the 2 Given Data Points as Initial Centroids Step 2: Find the Distances Between Each Data Point with the 2 Centroids

- 1st centroid: (1, 1)
- 2nd centroid: (2, 1)

Assume Euclidean distance is used.

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	1	0	2.8	4.2

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 3: Assign Each Data Point to the Closest Centroid

- 1st centroid: (1, 1)
- 2nd centroid: (2, 1)

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	1	0	2.8	4.2
Cluster	1	2	2	2

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 4: Re-compute the Centroids Using the Current Cluster Memberships

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	1	0	2.8	4.2
Cluster	1	2	2	2

• New 1st centroid:

 $x_1 = 1$ $x_2 = 1$

• New 2nd centroid:

$$x_1 = (2+4+5)/3 = 3.67$$

 $x_2 = (1+3+4)/3 = 2.67$

Step 5: Find the Distance Between Each Data Point with the 2 Centroids

- 1st centroid: (1, 1)
- 2nd centroid: (3.67, 2.67)

Assume Euclidean distance is used.

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	3.1	2.4	0.5	1.9

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 6: Assign Each Data Point to the Closest Centroid

- 1st centroid: (1, 1)
- 2nd centroid: (3.67, 2.67)

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	3.1	2.4	0.5	1.9
Cluster	1	1	2	2

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 7: Re-compute the Centroids Using the Current Cluster Memberships

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0	1	3.6	5
DC2	3.1	2.4	0.5	1.9
Cluster	1	1	2	2

• New 1st centroid:

$$x_1 = (1+2)/2 = 1.5$$

 $x_2 = (1+1)/2 = 1$

• New 2nd centroid:

$$x_1 = (4+5)/2 = 4.5$$

 $x_2 = (3+4)/2 = 3.5$

Step 8: Find the Distance Between Each Data Point with the 2 Centroids

- 1st centroid: (1.5, 1)
- 2nd centroid: (4.5, 3.5)

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0.5	0.5	3.2	4.6
DC2	4.3	2.4	0.5	1.9

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 9: Assign Each Data Point to the Closest Centroid

- 1st centroid: (1.5, 1)
- 2nd centroid: (4.5, 3.5)

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0.5	0.5	3.2	4.6
DC2	4.3	3.5	0.7	0.7
Cluster	1	1	2	2

where DC1, DC2 are the distances between the data points and 1st centroid, and 2nd centroid, respectively.

Step 7: Re-compute the Centroids Using the Current Cluster Memberships

Medicine	A	В	С	D
Weight Index	1	2	4	5
pH Index	1	1	3	4
DC1	0.5	0.5	3.2	4.6
DC2	4.3	3.5	0.7	0.7
Cluster	1	1	2	2

• New 1st centroid:

$$x_1 = (1+2)/2 = 1.5$$

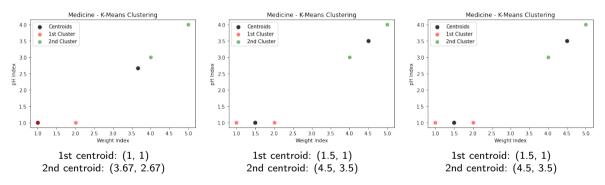
 $x_2 = (1+1)/2 = 1$

• New 2nd centroid:

$$x_1 = (4+5)/2 = 4.5$$

 $x_2 = (3+4)/2 = 3.5$

The 1st, the 2nd, and the 3rd Round



Conclusion

• Cluster 1:

The average weight index is 1.5, and the average pH index is 1.

• Cluster 2:

The average weight index is 4.5, and the average pH index is 3.5.



K-Means Clustering Implementation using Scikit-Learn

```
# Import the required libraries
import numpy as np
from sklearn.cluster import KMeans
import matplotlib.pyplot as plt
```

```
# Unlabled training data
data = np.array([[1,1], [2,1], [4,3], [5,4]])
```

```
# Initial centroids
init_centroids = np.array([[1,1], [2,1]])
```

```
# Create a KMeans object by specifying
# - Number of clusters (n_clusters) = 2, initial centroids (init) = init_centroids
# - Number of time the k-means algorithm will be run with different centroid seeds (n_init) = 1
# - Maximum number of iterations of the k-means algorithm for a single run (max_iter) = 3
kmeans = KMeans(n_clusters=2, init=init_centroids, n_init=1, max_iter = 3)
```

K-Means Clustering Implementation using Scikit-Learn

```
kmeans.fit(data)  # Compute k-means clustering
labels = kmeans.predict(data)  # Predict the closest cluster each sample in data belongs to
centroids = kmeans.cluster_centers_ # Get resulting centroids
fig, ax = plt.subplots()  # Defining 2D axes so that we can plot 2D data into it
```

```
# Get boolean arrays representing entries with labels = 0 and 1
a = np.array(labels == 0); b = np.array(labels == 1)
```

```
# Plot centroids with color = black, size = 50 units, transparency = 20%, and put label "Centroids"
ax.scatter(centroids[:,0], centroids[:,1], c="black", s=50, alpha=0.8, label="Centroids")
# Plot data in the different clusters (1st in red, 2nd in green) with transparency = 50%
ax.scatter(data[a,0], data[a,1], c="red", s=40, alpha=0.5, label="1st Cluster")
ax.scatter(data[b,0], data[b,1], c="green", s=40, alpha=0.5, label="2nd Cluster")
ax.legend() # Show legend
```

```
ax.set_xlabel("Weight Index")  # Put x-axis label "Weight Index"
ax.set_ylabel("pH Index")  # Put y-axis label "pH Index"
ax.set_title("Medicine - K-Means Clustering") # Put figure title
```

That's all! Any questions?

