## Practical implementation of k-means clustering

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#### Key steps

- Data pre-processing
- Choosing a number of clusters
- Running k-means clustering on pre-processed data
- Analyzing average RFM values of each cluster

#### Data pre-processing

We've completed the pre-processing steps and have these two objects:

- datamart\_rfm
- datamart\_normalized

```
import numpy as np
datamart_log = np.log(datamart_rfm)

from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
scaler.fit(datamart_log)

datamart_normalized = scaler.transform(datamart_log)
```

#### Methods to define the number of clusters

- Visual methods elbow criterion
- Mathematical methods silhouette coefficient
- Experimentation and interpretation

#### Running k-means

```
# Import package
from sklearn.cluster import KMeans
kmeans = KMeans(n_clusters=2, random_state=1)
# Compute k-means clustering on pre-processed data
kmeans.fit(datamart_normalized)
# Extract cluster labels from labels_ attribute
cluster_labels = kmeans.labels_
```

#### Analyzing average RFM values of each cluster

```
# Create a cluster label column in the original DataFrame
datamart_rfm_k2 = datamart_rfm.assign(Cluster = cluster_labels)
# Calculate average RFM values and size for each cluster
datamart_rfm_k2.groupby(['Cluster']).agg({
    'Recency': 'mean',
    'Frequency': 'mean',
    'MonetaryValue': ['mean', 'count'],
}).round(0)
```

#### Analyzing average RFM values of each cluster

The result of a simple 2-cluster solution:

	Recency	Frequency	MonetaryValue	
	mean	mean	mean	count
cluster				
0	137.0	5.0	92.0	2023
1	32.0	35.0	719.0	1620



## Let's practice running k-means clustering!

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### Choosing number of clusters

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#### Methods

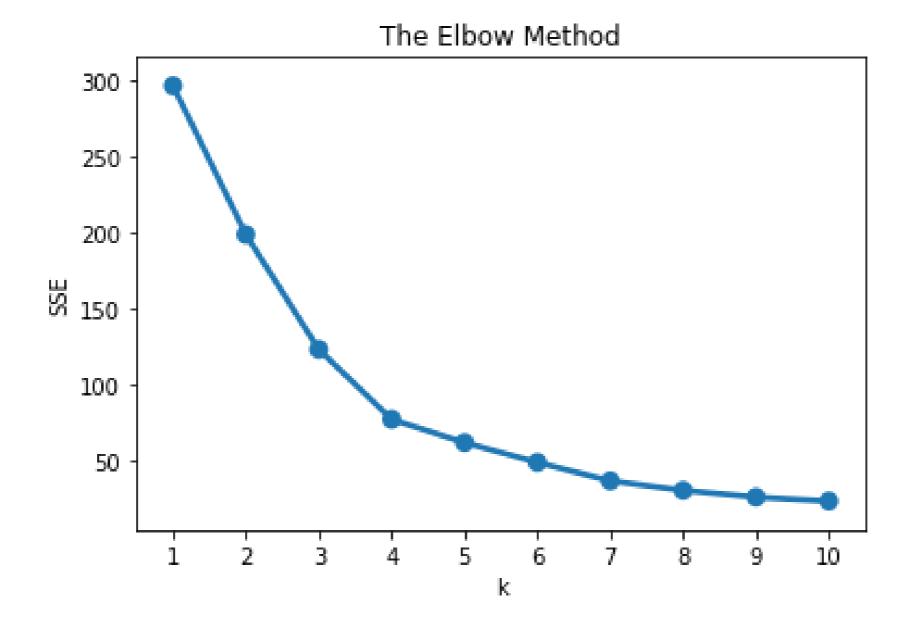
- Visual methods elbow criterion
- Mathematical methods silhouette coefficient
- Experimentation and interpretation

- Plot the number of clusters against within-cluster sum-of-squared-errors (SSE) sum of squared distances from every data point to their cluster center
- Identify an "elbow" in the plot
- Elbow a point representing an "optimal" number of clusters

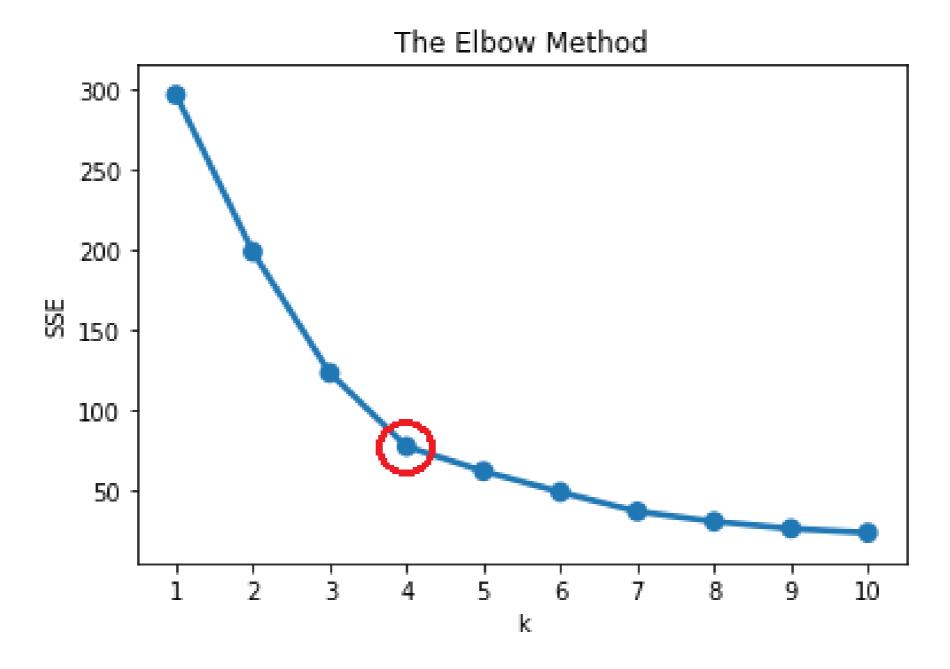
```
# Import key libraries
from sklearn.cluster import KMeans
import seaborn as sns
from matplotlib import pyplot as plt
# Fit KMeans and calculate SSE for each *k*
sse = {}
for k in range(1, 11):
    kmeans = KMeans(n_clusters=k, random_state=1)
    kmeans.fit(data_normalized)
    sse[k] = kmeans.inertia_ # sum of squared distances to closest cluster center
# Plot SSE for each *k*
plt.title('The Elbow Method')
plt.xlabel('k'); plt.ylabel('SSE')
sns.pointplot(x=list(sse.keys()), y=list(sse.values()))
plt.show()
```



The elbow criterion chart:

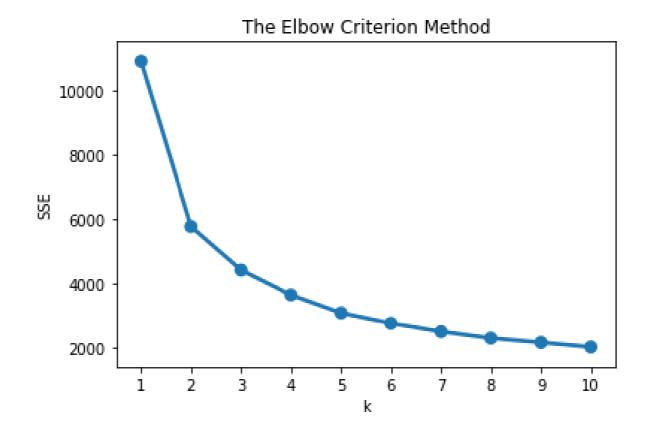


The elbow criterion chart:



#### Using elbow criterion method

- Best to choose the point on elbow, or the next point
- Use as a guide but test multiple solutions
- Elbow plot built on datamart\_rfm



#### Experimental approach - analyze segments

- Build clustering at and around elbow solution
- Analyze their properties average RFM values
- Compare against each other and choose one which makes most business sense



#### Experimental approach - analyze segments

	Recency	Frequency	MonetaryValue	
	mean	mean	mean	count
cluster				
0	137.0	5.0	92.0	2023
1	32.0	35.0	719.0	1620

- Previous 2-cluster solution
- 3-cluster solution on the same normalized RFM dataset

	Recency	Frequency	MonetaryValue	
	mean	mean	mean	count
cluster				
0	16.0	50.0	1051.0	901
1	167.0	3.0	53.0	1156
2	77.0	12.0	216.0	1586

### Let's practice finding the optimal number of clusters!

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### Profile and interpret segments

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#### Approaches to build customer personas

- Summary statistics for each cluster e.g. average RFM values
- Snake plots (from market research)
- Relative importance of cluster attributes compared to population



#### Summary statistics of each cluster

- Run k-means segmentation for several k values around the recommended value.
- Create a cluster label column in the **original** DataFrame:

```
datamart_rfm_k2 = datamart_rfm.assign(Cluster = cluster_labels)
```

Calculate average RFM values and sizes for each cluster:

```
datamart_rfm_k2.groupby(['Cluster']).agg({
    'Recency': 'mean',
    'Frequency': 'mean',
    'MonetaryValue': ['mean', 'count'],
}).round(0)
```

Repeat the same for k=3

#### Summary statistics of each cluster

• Compare average RFM values of each clustering solution

	Recency	Frequency	MonetaryValue	
	mean	mean	mean count	
cluster				
0	137.0	5.0	92.0	2023
1	32.0	35.0	719.0	1620
	Recency	Frequency	Moneta	ryValue
	Recency mean	Frequency mean	Moneta mean	ryValue count
cluster	_			-
cluster 0	_	mean		-
	mean	mean	mean	count



#### Snake plots to understand and compare segments

- Market research technique to compare different segments
- Visual representation of each segment's attributes
- Need to first normalize data (center & scale)
- Plot each cluster's average normalized values of each attribute

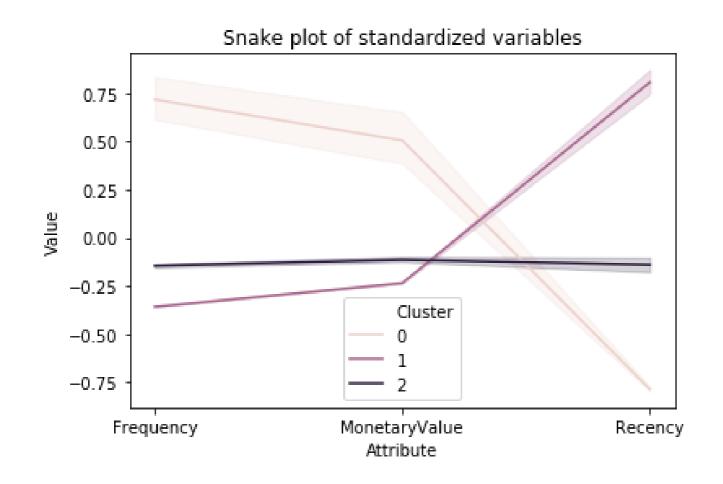
#### Prepare data for a snake plot

Transform datamart\_normalized as DataFrame and add a Cluster column

Melt the data into a long format so RFM values and metric names are stored in 1 column each

#### Visualize a snake plot

```
plt.title('Snake plot of standardized variables')
sns.lineplot(x="Attribute", y="Value", hue='Cluster', data=datamart_melt)
```



#### Relative importance of segment attributes

- Useful technique to identify relative importance of each segment's attribute
- Calculate average values of each cluster
- Calculate average values of population
- Calculate importance score by dividing them and subtracting 1 (ensures 0 is returned when cluster average equals population average)

```
cluster_avg = datamart_rfm_k3.groupby(['Cluster']).mean()
population_avg = datamart_rfm.mean()
relative_imp = cluster_avg / population_avg - 1
```

#### Analyze and plot relative importance

• As a ratio moves away from 0, attribute importance for a segment (relative to total pop.) increases.

```
relative_imp.round(2)
```

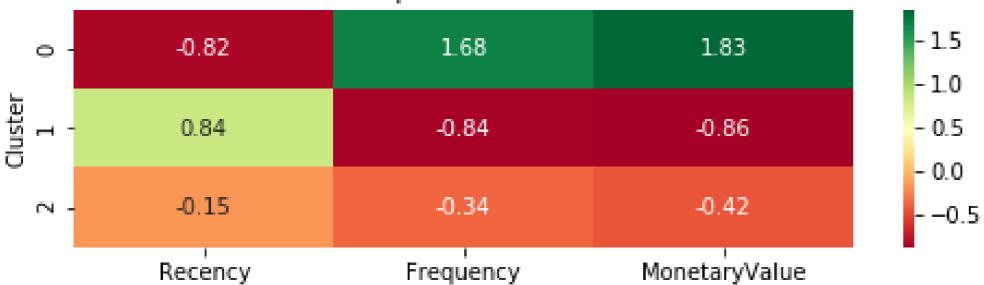
```
Recency Frequency MonetaryValue
Cluster
0 -0.82 1.68 1.83
1 0.84 -0.84 -0.86
2 -0.15 -0.34 -0.42
```

```
# Plot heatmap
plt.figure(figsize=(8, 2))
plt.title('Relative importance of attributes')
sns.heatmap(data=relative_imp, annot=True, fmt='.2f', cmap='RdYlGn')
plt.show()
```



#### Relative importance heatmap





	Recency	Frequency	MonetaryValue
Cluster			
0	-0.82	1.68	1.83
1	0.84	-0.84	-0.86
2	-0.15	-0.34	-0.42

# Your time to experiment with different customer profiling techniques!

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#### Implement end-toend segmentation solution

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#### Key steps of the segmentation project

- Gather data updated data with an additional variable
- Pre-process the data
- Explore the data and decide on the number of clusters
- Run k-means clustering
- Analyze and visualize results



#### **Updated RFM data**

- Same RFM values plus additional Tenure variable
- Tenure time since the first transaction
- Defines how long the customer has been with the company

	Recency	Frequency	MonetaryValue	Tenure
CustomerID				
12747	3	25	948.70	362
12748	1	888	7046.16	365
12749	4	37	813.45	214
12820	4	17	268.02	327
12822	71	9	146.15	88



#### Goals for this project

- Remember key pre-processing rules
- Apply data exploration techniques
- Practice running several k-means iterations
- Analyze results quantitatively and visually

#### Let's dig in!

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#### Final thoughts

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#### What you have learned

- Cohort analysis and visualization
- RFM segmentation
- Data pre-processing for k-means
- Customer segmentation with k-means
  - Evaluating number of clusters
  - Reviewing and visualizing segmentation solutions



#### Congratulations!

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