# Basics of k-means clustering

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## Why k-means clustering?

- A critical drawback of hierarchical clustering: runtime
- K means runs significantly faster on large datasets

#### Step 1: Generate cluster centers

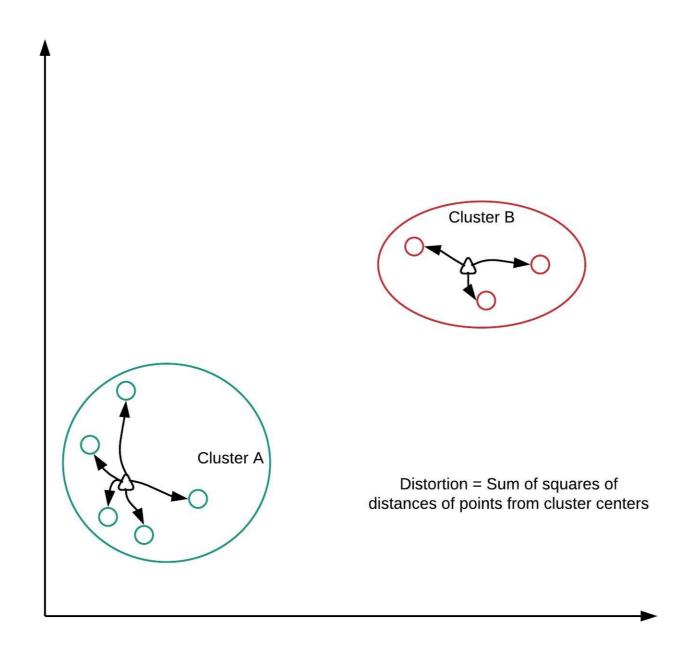
kmeans(obs, k\_or\_guess, iter, thresh, check\_finite)

- obs : standardized observations
- k\_or\_guess : number of clusters
- iter: number of iterations (default: 20)
- thres: threshold (default: 1e-05)
- check\_finite: whether to check if observations contain only finite numbers (default: True)

Returns two objects: cluster centers, distortion



#### How is distortion calculated?





#### Step 2: Generate cluster labels

```
vq(obs, code_book, check_finite=True)
```

- obs : standardized observations
- code\_book : cluster centers
- check\_finite: whether to check if observations contain only finite numbers (default: True)

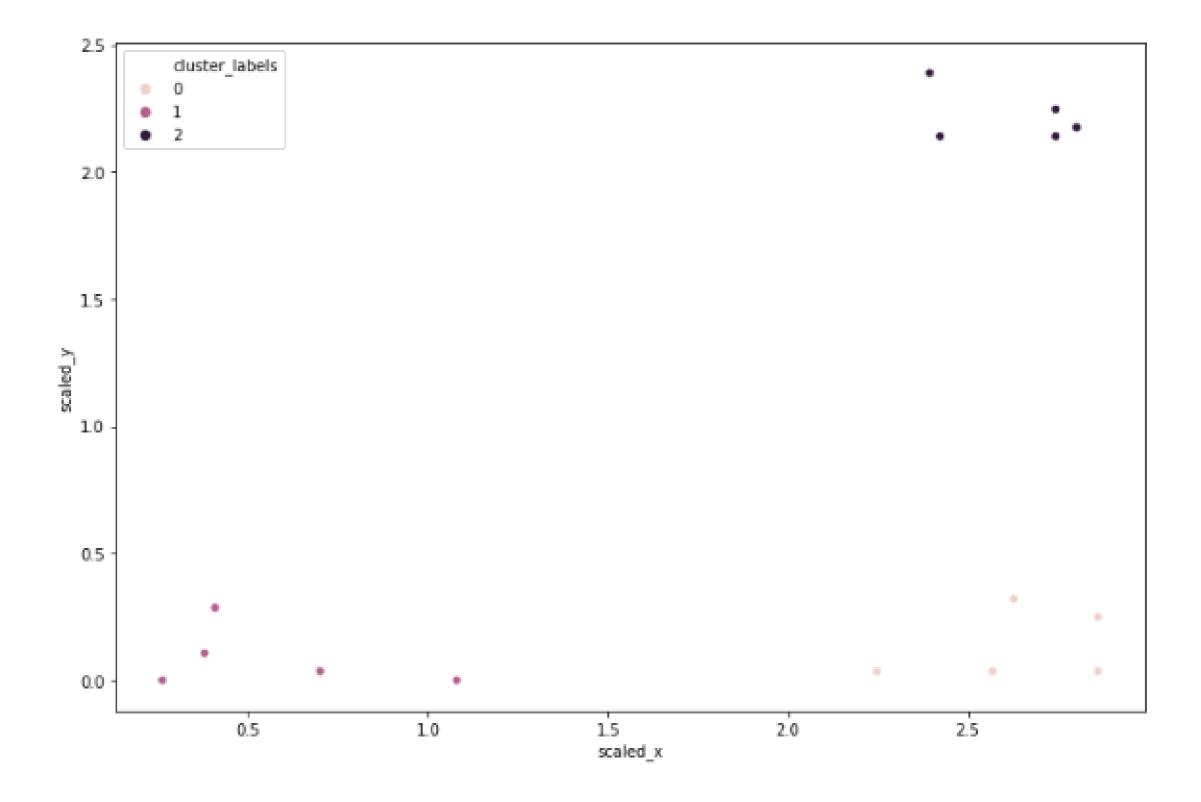
Returns two objects: a list of cluster labels, a list of distortions

#### A note on distortions

- kmeans returns a single value of distortions
- vq returns a list of distortions.

#### Running k-means

```
# Import kmeans and vq functions
from scipy.cluster.vq import kmeans, vq
# Generate cluster centers and labels
cluster_centers, _ = kmeans(df[['scaled_x', 'scaled_y']], 3)
df['cluster_labels'], _ = vq(df[['scaled_x', 'scaled_y']], cluster_centers)
# Plot clusters
sns.scatterplot(x='scaled_x', y='scaled_y', hue='cluster_labels', data=df)
plt.show()
```



# Next up: exercises!

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## How many clusters?

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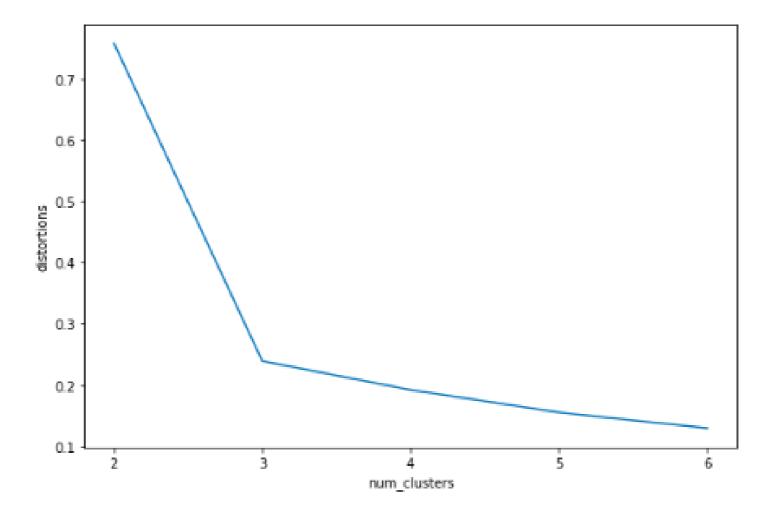


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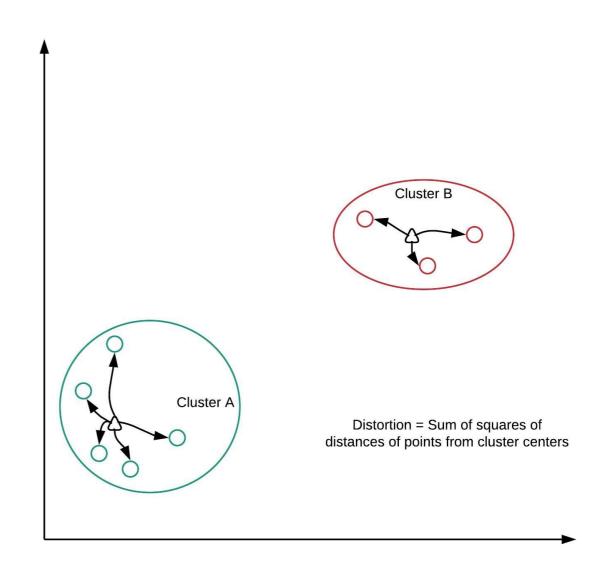
#### How to find the right k?

- No absolute method to find right number of clusters (k) in k-means clustering
- Elbow method



#### Distortions revisited

- Distortion: sum of squared distances of points from cluster centers
- Decreases with an increasing number of clusters
- Becomes zero when the number of clusters equals the number of points
- Elbow plot: line plot between cluster centers and distortion



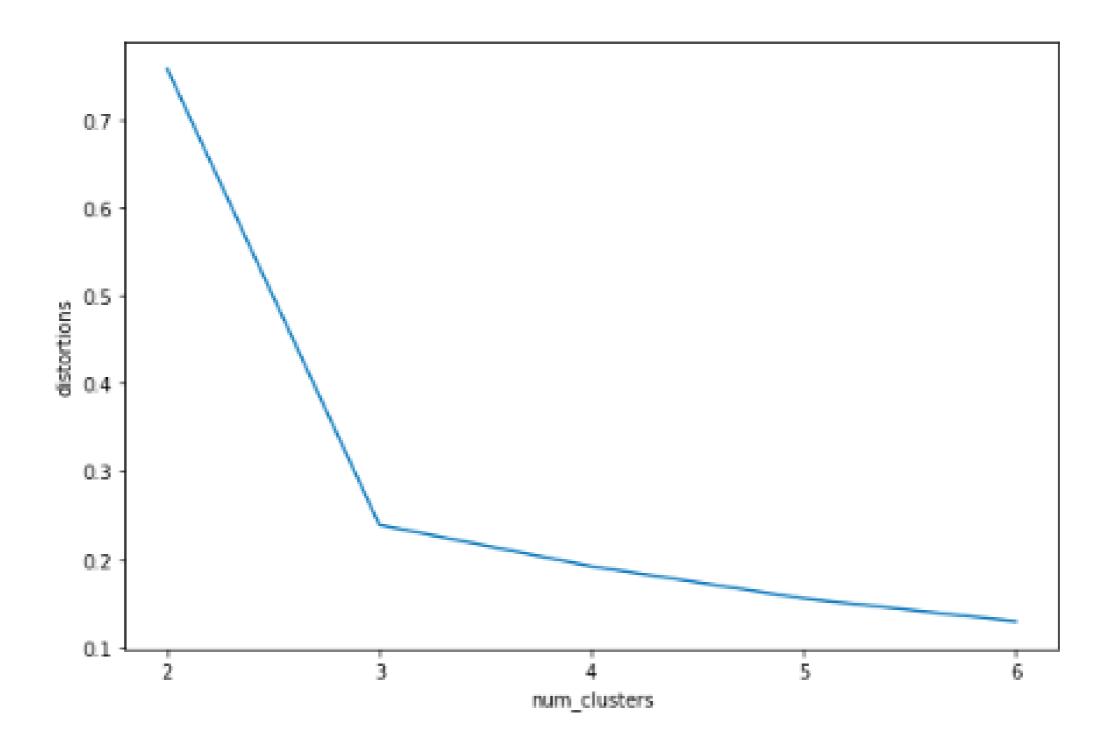
#### **Elbow method**

- Elbow plot: plot of the number of clusters and distortion
- Elbow plot helps indicate number of clusters present in data

## Elbow method in Python

```
# Declaring variables for use
distortions = []
num_clusters = range(2, 7)
# Populating distortions for various clusters
for i in num_clusters:
    centroids, distortion = kmeans(df[['scaled_x', 'scaled_y']], i)
    distortions.append(distortion)
# Plotting elbow plot data
elbow_plot_data = pd.DataFrame({'num_clusters': num_clusters,
                                'distortions': distortions})
sns.lineplot(x='num_clusters', y='distortions',
             data = elbow_plot_data)
plt.show()
```





## Final thoughts on using the elbow method

- Only gives an indication of optimal k (numbers of clusters)
- Does not always pinpoint how many k (numbers of clusters)
- Other methods: average silhouette and gap statistic

# Next up: exercises

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# Limitations of kmeans clustering

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## Limitations of k-means clustering

- How to find the right \_K\_ (number of clusters)?
- Impact of seeds
- Biased towards equal sized clusters

#### Impact of seeds

Initialize a random seed

from numpy import random
random.seed(12)

Seed: np.array(1000, 2000)

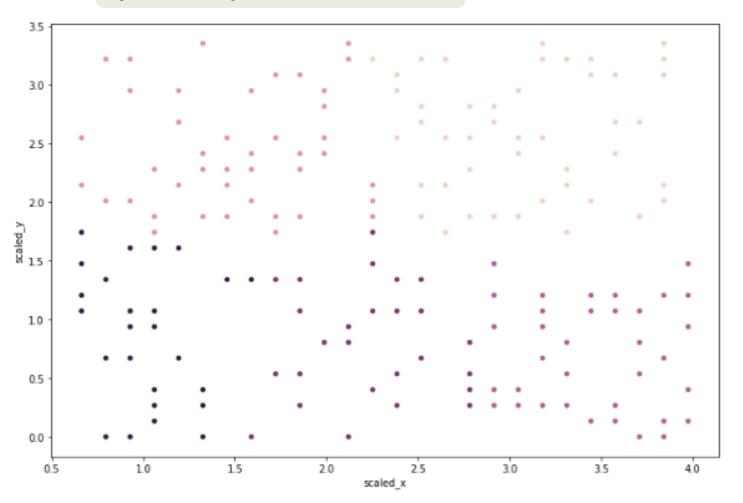
Cluster sizes: 29, 29, 43, 47, 52

Seed: np.array(1,2,3)

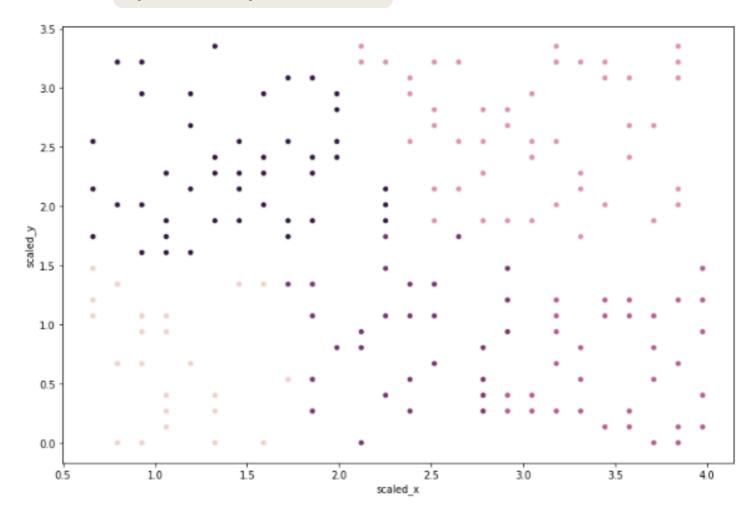
Cluster sizes: 26, 31, 40, 50, 53

#### Impact of seeds: plots

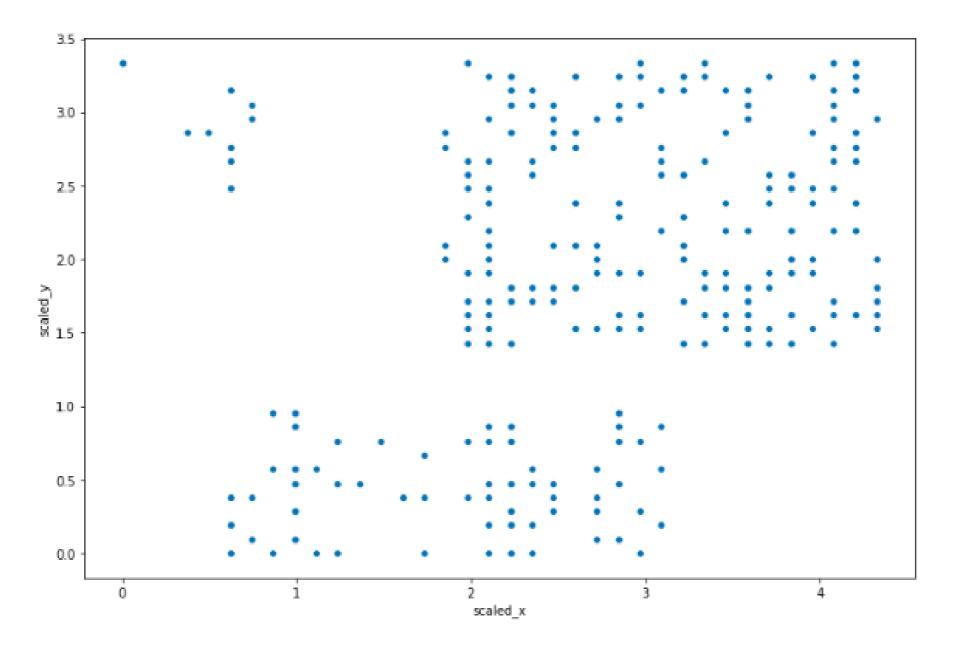
Seed: np.array(1000, 2000)



Seed: np.array(1,2,3)



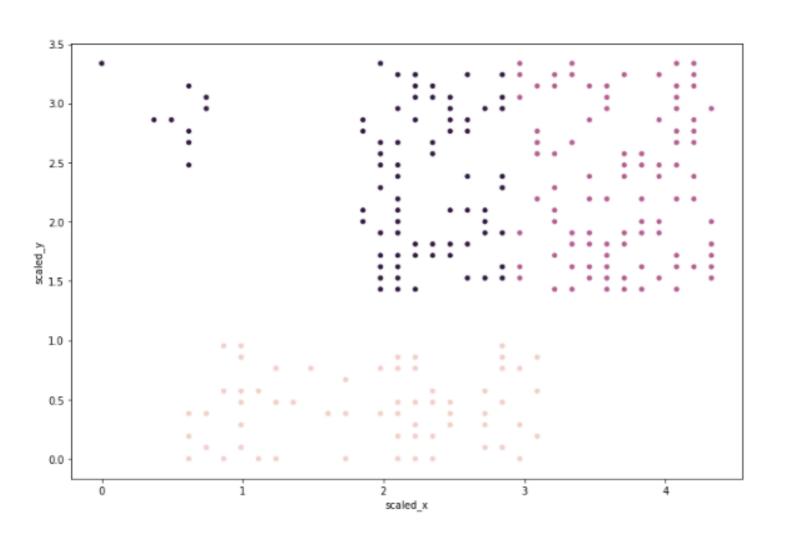
#### Uniform clusters in k means



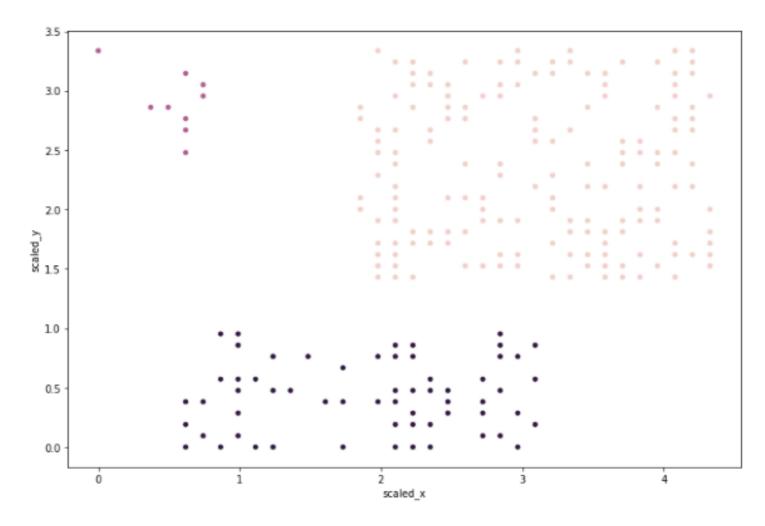


## Uniform clusters in k-means: a comparison

K-means clustering with 3 clusters



Hierarchical clustering with 3 clusters



## Final thoughts

- Each technique has its pros and cons
- Consider your data size and patterns before deciding on algorithm
- Clustering is exploratory phase of analysis



# Next up: exercises

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