

Digital Technologies and Value Creation

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Learning objectives of today

Goals:

- Understand what clustering is
- Understand how K-means clustering works
- Understand how hierarchical clustering works

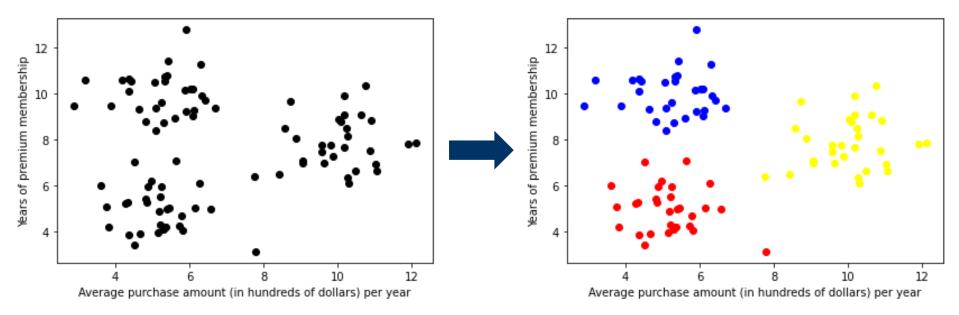
How will we do this?

- Learn the technical parts and how to use this in Python. In class, we'll make things more intuitive
- "Happiness Index" dataset: cluster countries based on some of their salient characteristics and plot on maps



What is clustering?

What is clustering



Each datapoint corresponds to a customer.

The goal of clustering is to group together points that have "similar" characteristics.

(Here, we could cluster to provide, e.g., specific promotional offers.)



Examples of clustering applications



Image segmentation (e.g., background vs foreground)



Detection of cancer (finding lumps in scans)



Segmentation of customers



Credit card or insurance claims fraud



Communities in social networks



Phylogeny (clustering of species)



... and many more

Clustering – an overview

- Clustering creates groups:
 - Within the same group: observations similar
 - Across groups: observations dissimilar
- **Similarity** = two observations are similar if the features of one observation are "close" to the features of the other
- What does "close" mean? → Many methods for clustering: we consider two very popular ones:
 - K-means
 - Hierarchical clustering



The happiness index

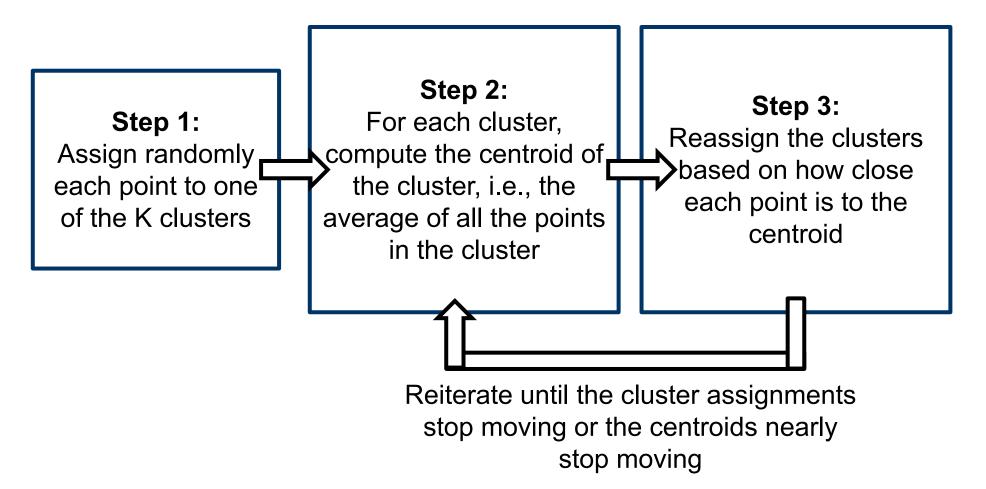
- The World Happiness Report is an annual publication of the United Nations Sustainable Development Solutions Network
- It contains attributes of different countries, feeding into one index, the happiness index.
- Goal: cluster these countries

	Country or region	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices	Generosity	Perceptions of corruption
0	Finland	1.340	1.587	0.986	0.596	0.153	0.393
1	Denmark	1.383	1.573	0.996	0.592	0.252	0.410
2	Norway	1.488	1.582	1.028	0.603	0.271	0.341
3	Iceland	1.380	1.624	1.026	0.591	0.354	0.118
4	Netherlands	1.396	1.522	0.999	0.557	0.322	0.298
		1	N		7.30		1
140	Rwanda	0.359	0.711	0.614	0.555	0.217	0.411
141	Tanzania	0.476	0.885	0.499	0.417	0.276	0.147
142	Afghanistan	0.350	0.517	0.361	0.000	0.158	0.025
143	Central African Republic	0.026	0.000	0.105	0.225	0.235	0.035
144	South Sudan	0.306	0.575	0.295	0.010	0.202	0.091



K-means clustering

K-means clustering overview



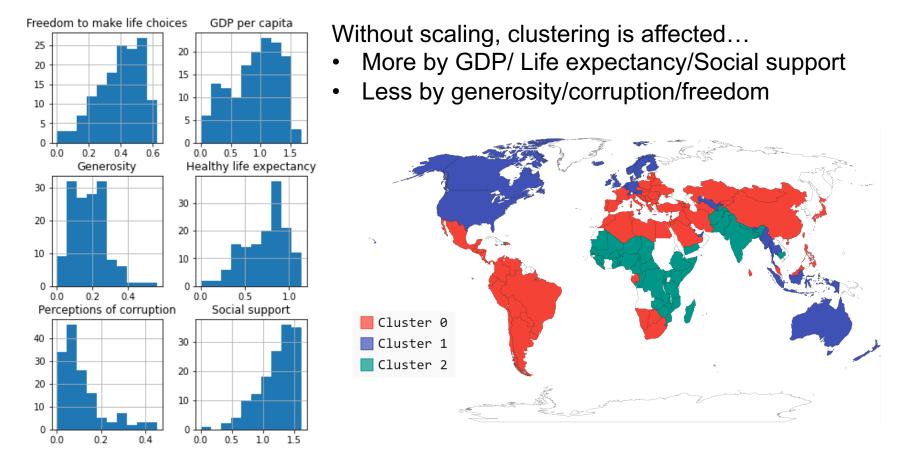


K-means clustering in Python





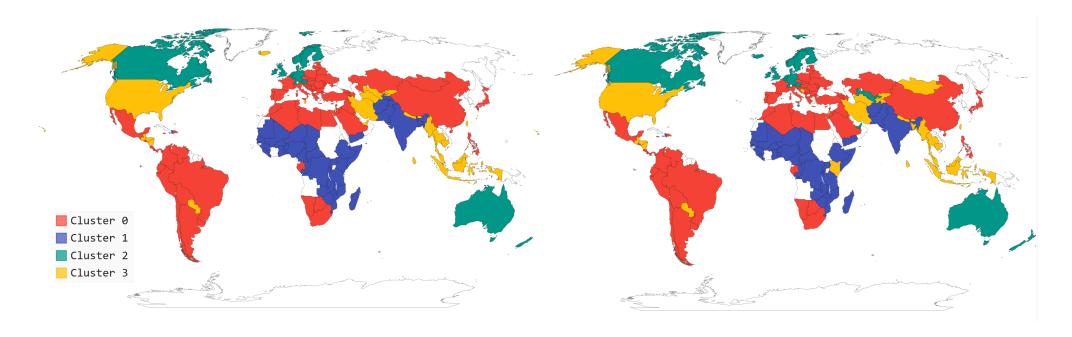
The importance of scaling



	Country or region	GDP per capita	Social support	Healthy life expectancy	Freedom to make life choices	Generosity	Perceptions of corruption	Code
27	Saudi Arabia	1.403	1.357	0.795	0.439	0.08	0.132	sa
14	United Kingdom	1.333	1.538	0.996	0.45	0.348	0.278	gb



The impact of random sampling



Exactly the same code is run both times, but we get different outputs.

Why?

Due to **Step 1:** each country is put into a cluster at random. When this changes, the end-result can also change.

How to mitigate this effect?



The impact of random sampling

Idea:

Run K-means many times (n_init is set to 10 by default)

```
kmeans = KMeans(n_clusters=4,n_init=1).fit(happiness_quant)
```

 Among all possible models obtained, pick the one that is best and work with that one.

What do we mean by "best"?

- For each cluster, compute the distance of each point in the cluster to the centroid of the cluster.
- Add all these distances squared together: inertia of the model
- The best model is the one which has the smallest inertia



How to pick K?

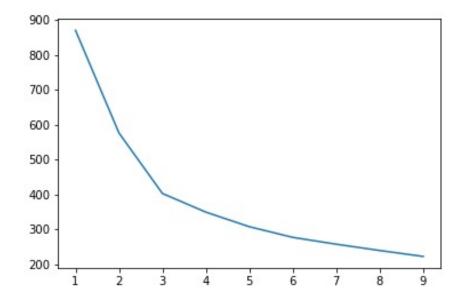
In all these examples, K was given to us. How to pick K?

- For different values of K, run K-means and retain the inertia each time
- Plot the inertia as a function of K: elbow plot

```
inertia_K=[]
K = range(1,10)
for k in K:
    kmeanModel = KMeans(n_clusters=k)
    kmeanModel.fit(happiness_quant)
    inertia_K.append(kmeanModel.inertia_)
```

Choose what is the "elbow" of the graph: here, K=3.

"Best return for investment"

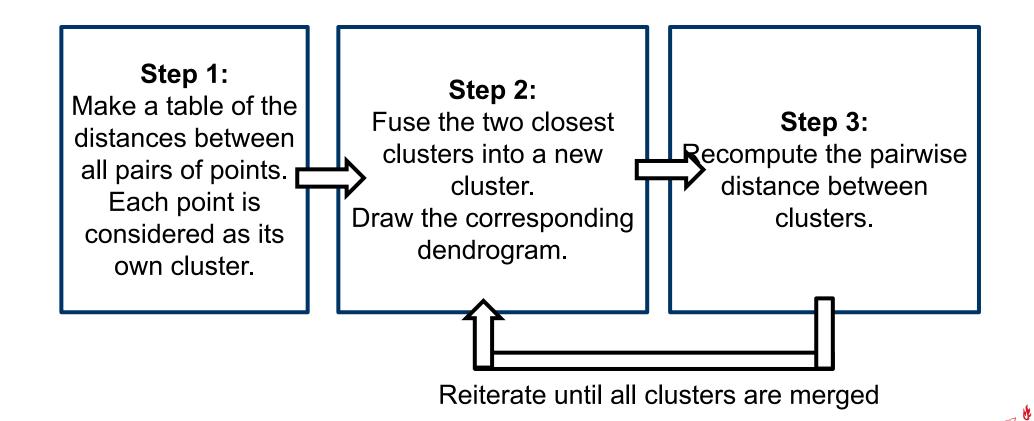




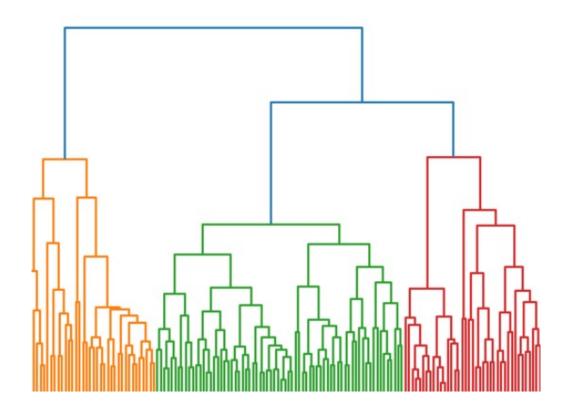


Hierarchical clustering

Hierarchical clustering overview



A dendogram





Hierarchical clustering in Python

- we use scipy instead of scikit-learn (dendrogram easier to obtain)
- Scaling is important for hierarchical clustering



Different linkages

- We build the dendogram based on "closeness" of clusters
- It's easy to measure the distance between two observations
- But when are clusters containing multiple observations "close" to each other?
 - Single-linkage: shortest distance between any pair of observations within the two clusters
 - Complete-linkage: distances between the farthest of any pair of observations within the two clusters
 - Average-linkage: distances between pairs of observations within the two clusters are averaged
 - Ward linkage: weighted distance between the centers of each cluster
 - ...



Hierarchical clustering in Python

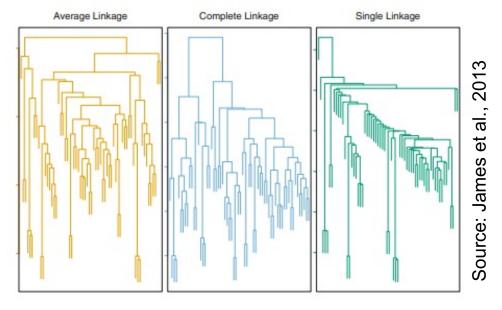




Different linkages give different results

Same dataset with different types of linkages can give completely different

results:



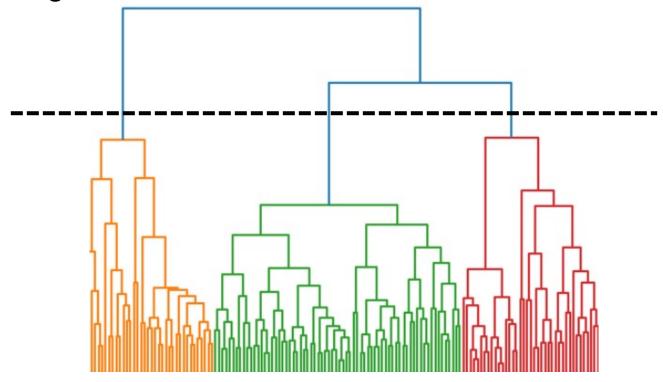
In general:

- Complete, average and Ward linkage tend to yield evenly sized clusters
- Single linkage tends to yield extended clusters to which single leaves are fused one by one
- Rule of thumb: complete or average linkage or Ward. Go with the dendrogram.



How to get clusters from dendograms?

Cut the final dendrogram at some level:



• In general: try and cut where there's a "jump" on the dendrogram. There may be multiple options and the best answer depends on the context.



Further study

- Part 3 of the video-exercise notebook contains another example of clustering, this time on newspaper articles
- It also highlights the aspect of interpreting clusters



