

#### CASA0006

- 1 Introduction to Module
- 2 Supervised Machine Learning
- 3 Tree-based Methods
- 4 Artificial Neural Networks
- **5** Analysis Workflow

- 6 Panel Regression
- 7 Difference in Difference
- 8 Regression Discontinuity
- **9** Dimensionality Reduction
- 10 Spatial Clustering



# Connecting with CASA0007 (T1) Clustering: Plan of Attack

#### **Standardisation Methods**

Z-Score (roughly symmetrical data)

Min-Max rescaling (asymmetric data)

IDR rescaling (data with significant outliers)

**Explicit rescaling** 

## **Clustering Methods**

K-Means

Hierarchical

### **Clustering Quality**

SSE

Silhouette Analysis

#### **Visualisation**

Elbow Diagram

Silhouette Plot

Dendrogram

Scatter Plots

### **Follow Up**

Examine cluster centroids

Describe cluster characteristics

Compare against unconsidered variables

/ categories / geography

Consider analysing clusters separately



## Connecting with CASA0013 (T1, W10)

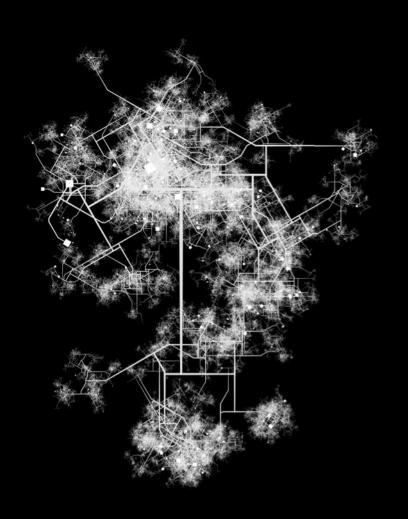
- Geodemographics
  - Booth map
  - London Output Area Classification
- Clustering methods

#### **Different Approaches**

Algorithm	Pros	Cons	Geographically Aware?
k-Means	Fast. Deterministic.	Every observation to cluster.	N.
DBSCAN	Allows for clusters <i>and</i> outliers.	Slower. Choice of \$\$\epsilon\$\$ critical. Can end up with all outliers.	N, but implicit in \$\$\epsilon\$\$.
OPTICS	Fewer parameters than DBSCAN.	Even slower.	N, but implicit in \$\$\epsilon\$\$.
Hierarchical	Can cut at any number of clusters.	No 'ideal' solution.	Y, with connectivity parameter
ADBSCAN	Scales. Confidence levels.	May need large data set to be useful. Choice of \$\$\epsilon\$\$ critical.	Υ.
Мах-р	Coherent regions returned.	Very slow if model poorly specified.	Υ.



# Outline



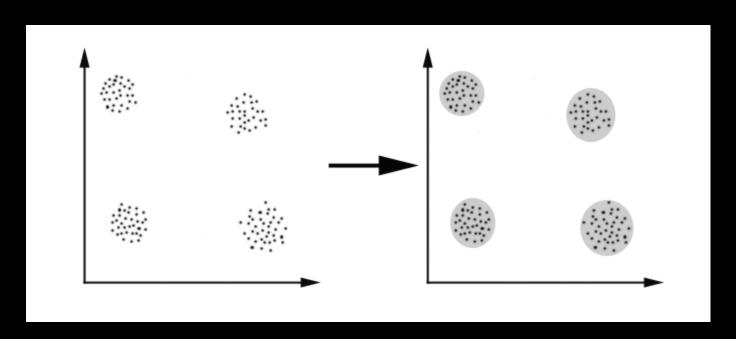
- 1. Definition and workflow
- 2. Clustering Methods
  - a. K-Means
  - b. Hierarchical
  - c. DBSCAN
  - d. Choosing clustering methods
- 3. Spatial Clustering
- Measuring Clustering Quality
  - a. SSE/Elbow Method
  - b. Silhouette Analysis
- 4. Next steps



# Clustering

## **Definition**

Type of analysis that divides data points into groups based on some similarity criteria





## Clustering

- Purpose of clustering
  - Discover groups of similar data points
  - Extract 'knowledge' from data
- What is a cluster?
  - A group of similar data points



## **Standardisation**

### Z score

(for not highly skewed data)

$$z = \frac{x - \mu}{\sigma}$$

## Min-Max Rescaling

(for highly skewed data)

$$x' = \frac{x - x_{min}}{x_{max} - x_{min}}$$

## **IDR Standardisation**

(Non-normal data with significant outliers)

$$x^{\text{IDR}} = \begin{cases} \frac{x - P_{50}}{P_{90} - P_{50}}, x \ge P_{50} \\ \frac{x - P_{50}}{P_{50} - P_{10}}, x < P_{50} \end{cases}$$

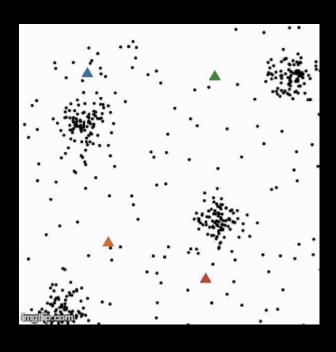
#### Criteria

- 1. Highly skewed distribution?
- 2. Significant outliers?



# Clustering K-Means Clustering

K-Means clustering **breaks down** a dataset into groups, based on proximity of points within a multidimensional space.



#### **Iterative Algorithm**

- 1 Place k centroids randomly within space
- 2 Assign points to nearest centroid
- 3 Recalculate centroids as the new mean of the cluster
- 4 Continue until centroid assignments no longer change

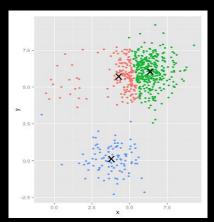
Interactive demo of kmeans: https://jeff3dx.github.io/kmc



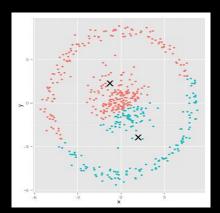
# Clustering Problems with K-Means Clustering

- Requires knowledge of the number of clusters, which you may not know in advance (solution: Elbow method);
- Sensitive to initialisation, which can lead to poor solutions (solution: try different random initialisation and pick up the best one);
- Sensitive to outliers, which can result in inaccurate clusters (solution: use another clustering method, or remove outliers);
- Incapable of handling clusters of a non-convex shape (no solution);
- Inapplicable to categorical data (solution: k-modes or k-prototypes).

### Choose k wisely



#### Non-convex shape



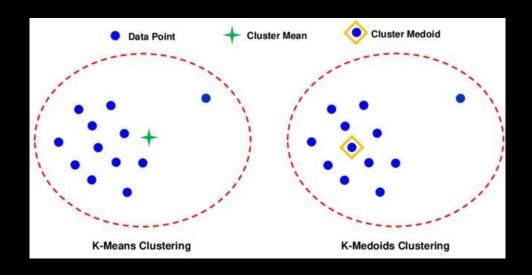


# **Extension of kmeans** K-modes and K-prototypes

method	Input variables
K-means	numerical
K-modes	categorical
K-prototypes	numerical and categorical



## K-medoids



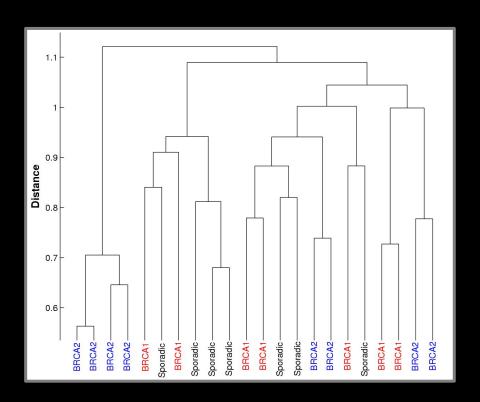
	Cluster 'centre'	Distance metric	Robustness to outlier	Computation cost
K-means	Mean of points in a cluster	Distance to the cluster mean	Not robust	Usually low
K-medoids	One of the points in a cluster	Any similarity measure	Robust	Much higher

https://scikit-learn-extra.readthedocs.io/en/stable/modules/cluster.html#k-medoids



# Hierarchical Clustering Agglomerative

Hierarchical clustering **builds up** clusters based on proximity of instances, ending on reaching predefined number of points



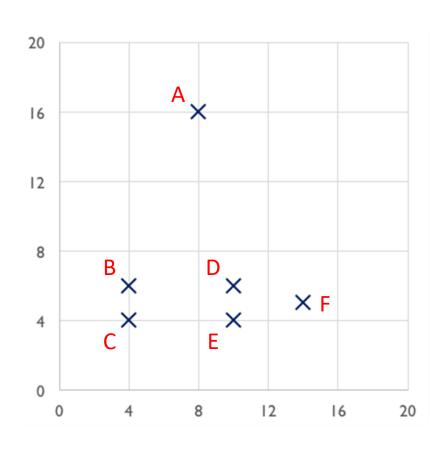
#### **Iterative Algorithm**

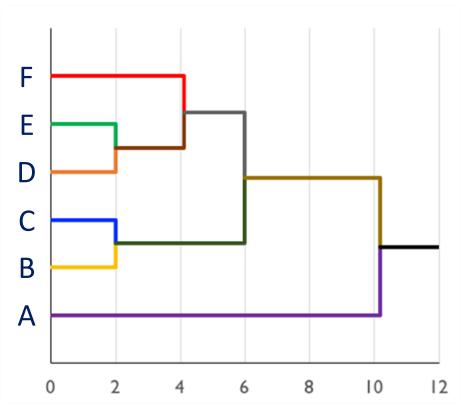
- 1 Start with every point in its own cluster
- 2 Merge points according to a *linkage* criterion (or distance)
- 3 Compute centroid of new clusters
- 4 Expand linkage threshold and continue until all points in one cluster

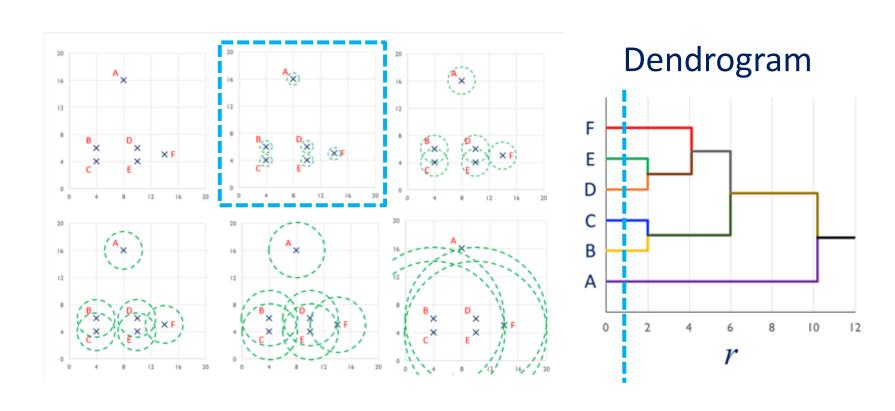
#### Pros

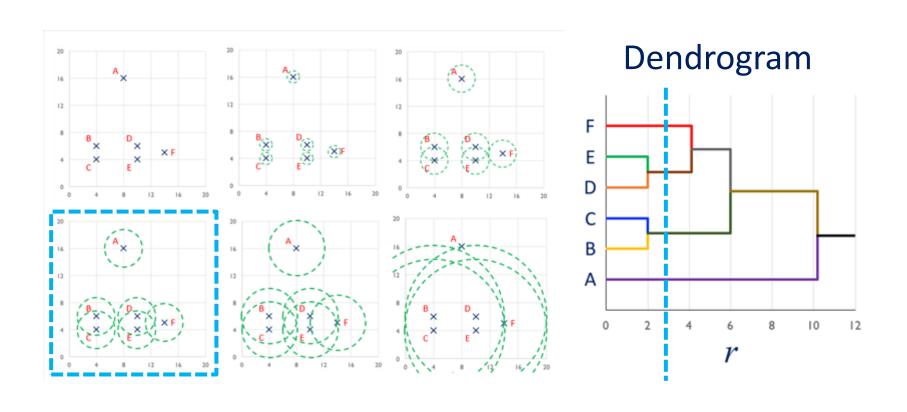
- Users can choose the level in a hierarchy structure;
- No prior knowledge of data required

## Dendrogram









## **Hierarchical Clustering**

Agglomerative

Bottom Up: Begins with one cluster per data point;

Gradually merge into larger clusters.

Divisive

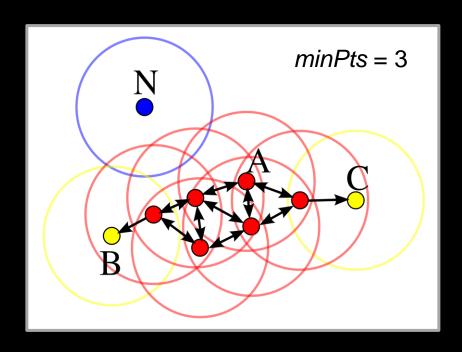
Top Down: Begins with one big cluster;

Gradually split into smaller clusters.



# Clustering Density-based – DBSCAN Clustering

DBSCAN builds clusters of points based on local proximity, considering neighbours within a maximum distance threshold

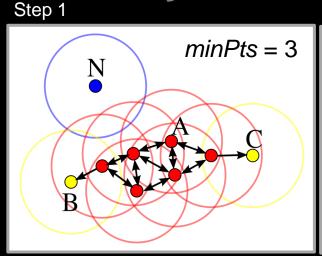


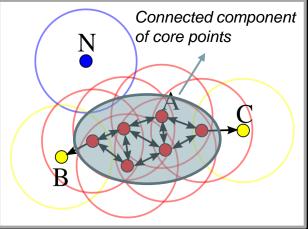
Given ε (search radius) and *minPts*, points are classified into three classes:

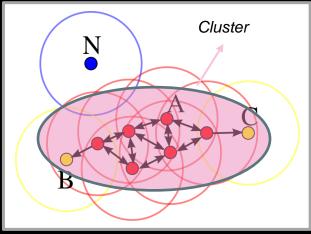
- 1. Point p is **core point**: if at least *minPt*s points are within distance ε of it (including p)
- Point p is edge point: if p is not a core point but it is reachable from a core point
- 3. Point p is **outlier**: all points not reachable from any core points



# Clustering Density-based – DBSCAN Clustering Step 2 Step 3







### Process (given ε and minPts)

- 1 Identify core points (with at least *minPt*s neighbours)
- 2 Connect core points while ignoring non-core points (forming connected components)
- 3 Assign each non-core point to a nearby cluster if it is within ε of a cluster, otherwise assign it to noise



# **Summary Three clustering methods**

method	required parameters	Extensions
kmeans	k (number of clusters)	K-modes, k-medoids, K-prototypes
hierarchical	No required parameter before clustering, but you should decide number of clusters afterwards	NA
DBSCAN	ε and <i>minPt</i> s	NA



## Choosing a cluster method ...

## There is no best way. Some issues are important:

- 1 Ability to cluster at speed for the given data size (the larger data, the fewer choices)
- 2 Accommodating the data types (numerical, categorical, or mixed)
- 3 Ability to cope with outliers (if the data have many outliers, then choose a method with robustness to outliers)
- 4 You can compare different methods and choose the best one



## Spatial clustering

- The methods above are 'general-purpose' clustering.
- They are applicable to non-spatial variables (e.g. house price, # bedrooms), or spatial variables (e.g. long-lat).
- Note the different implications
  - If you use non-spatial variables for clustering, a cluster represents a type of 'observation' with similar attributes ('<u>feature homogeneity</u>' or '<u>attribute similarity</u>')
  - If you explicitly consider spatial variables in clustering, a cluster represents a geographical 'place' or a region ('geographic cohesion')



## **Spatial clustering**

Given data points with both non-spatial and spatial variables, there are three approaches to cluster these points:

- 1. Clustering on only non-spatial variables, and then exploring the geography of clusters;
- Clustering on non-spatial variables but with constraints of 'geographic cohesion';
- 3. Clustering on both non-spatial and spatial variables\*
- Note the tension and trade-off between them.

<sup>\*</sup> See this paper: https://doi.org/10.1080/13658816.2021.1934475



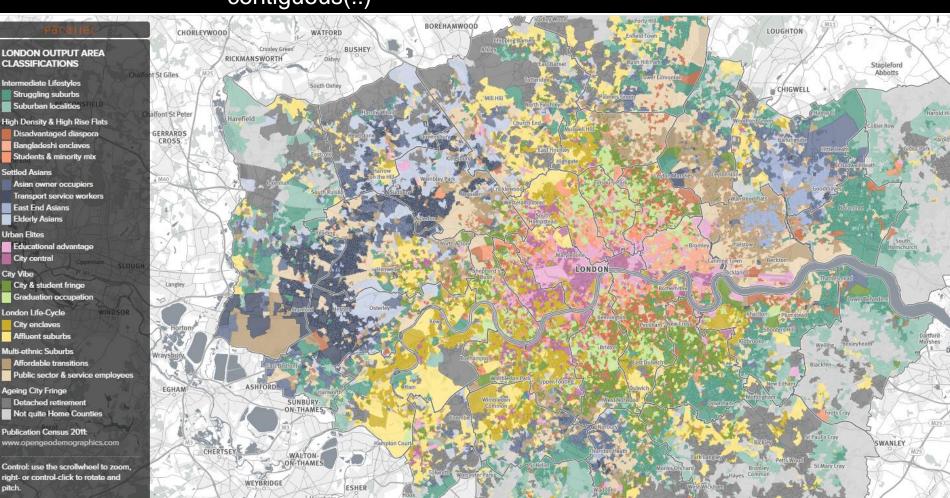
## **Approach One**

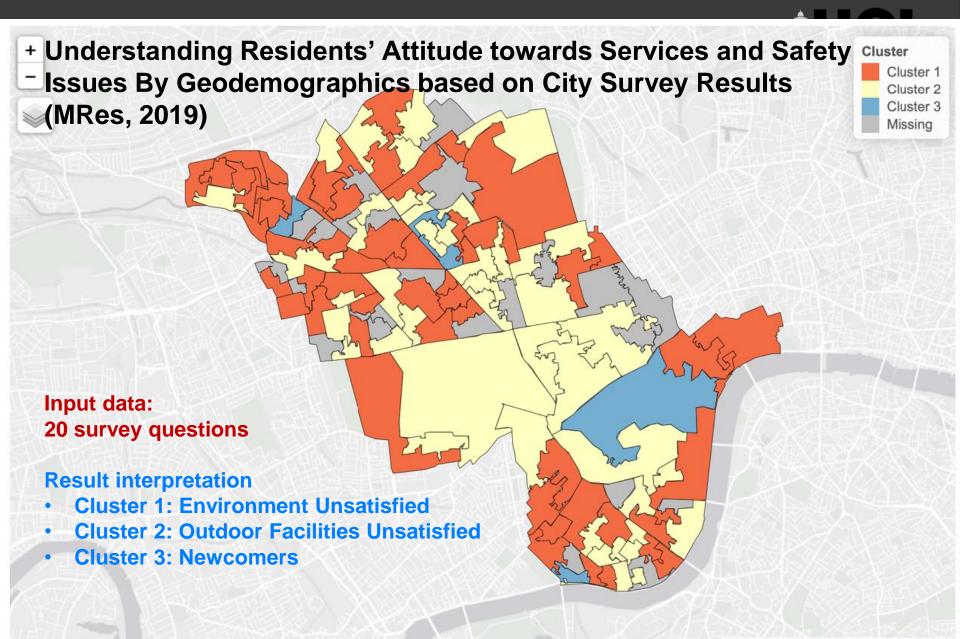
- Clustering on only non-spatial variables, and then exploring the geography of clusters;
- "usually ignores geographical coherence at the outset, but then explores the geography of uncovered solutions" (Wolf, 2021)
- Example: geodemographic analysis (London OA classification)
- Pros: it works well for geodemographics
- Cons: geographic cohesion is not sufficiently accounted for.



## LOAC

- Clustering OAs on 70+ socio-economic variables (non-spatial);
   32000 OAs are clustered into 8 groups
- 2. Clusters have obvious spatial patterns but aren't spatially contiguous(!!)





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Figure 33. Cluster Map of HAC for Index of Service Usage Rate and Satisfaction

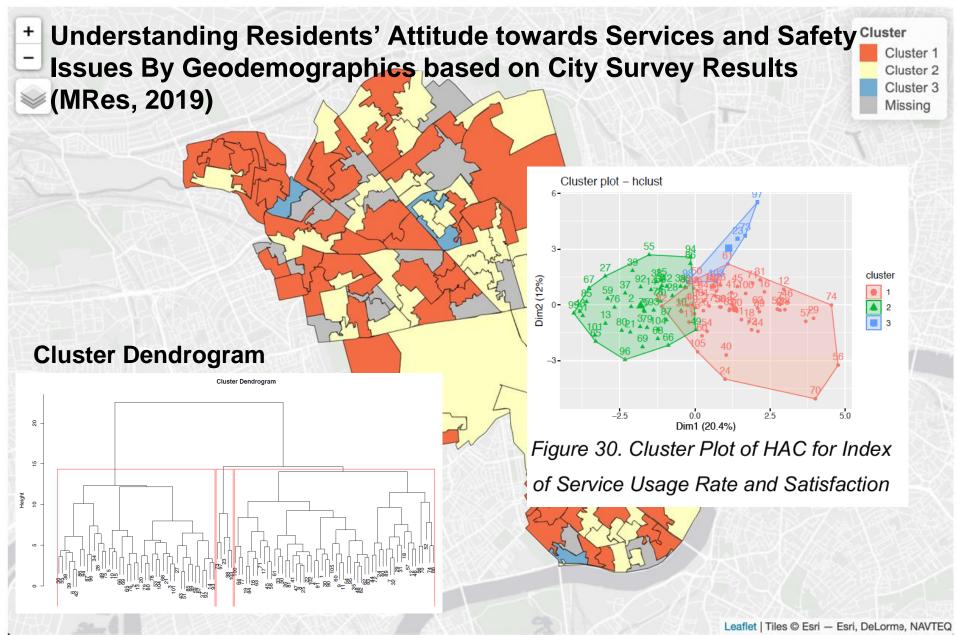


Figure 33. Cluster Map of HAC for Index of Service Usage Rate and Satisfaction



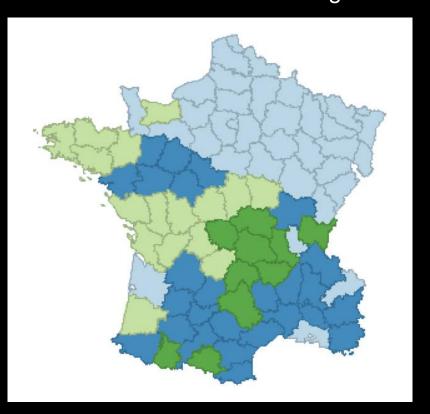
## **Approach Two**

- Clustering on non-spatial variables while adding constraints of 'geographic cohesion'
- Regions are 'coherent' if and only if they are geographically contiguous or connected
- Also called Regionalization, districting, spatially constrained clustering in literature.
- Pros: it simultaneously considers feature homogeneity and geographic cohesion
- Cons: computationally expensive



# **Spatial contiguity**

None of the clusters are contiguous



Yes, all clusters are spatially contiguous





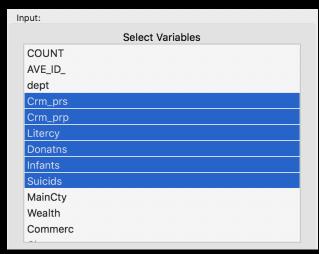
## The max-p method

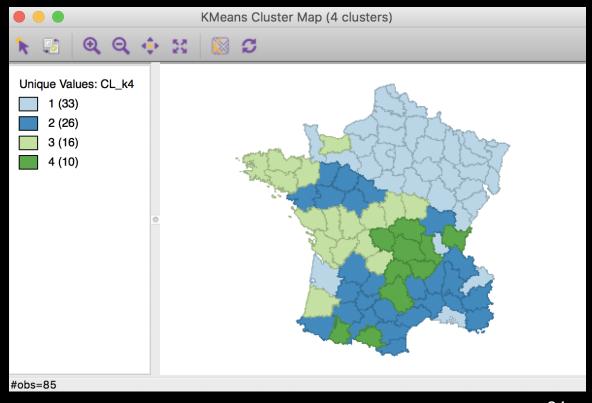
- Max-p: clustering of a set of geographic areas into the maximum number of regions such that the value of each region (e.g. population) is above a predefined threshold value
- What is a region? For each region, all parts are spatially connected to all other parts.
- Hyperparameter: the predefined threshold value
- The number of clusters or the maximum number of clusters is not predefined



## Comparing kmeans and max-p

- Case study: Guerry data set on moral statistics in 1830 France
- Method 1: generic kmeans (6 attributes, without considering geometric centroids), k=4. None of the clusters is spatially contiguous.

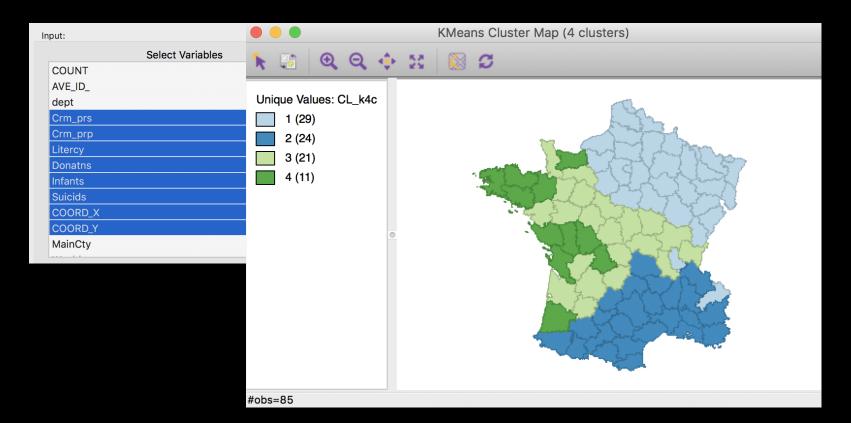






## Comparing kmeans and max-p

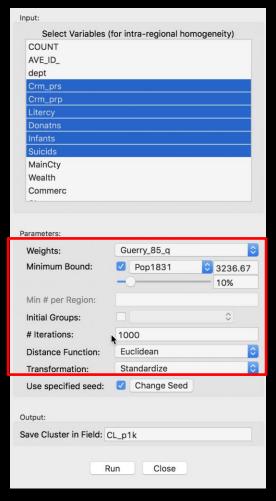
- Method 2: kmeans with centroids included as variables (8 attributes).
- Group 2 and 3 achieve contiguity. group 1 consists of three parts (including two singletons), and group 4 consists of four parts (including two singletons)

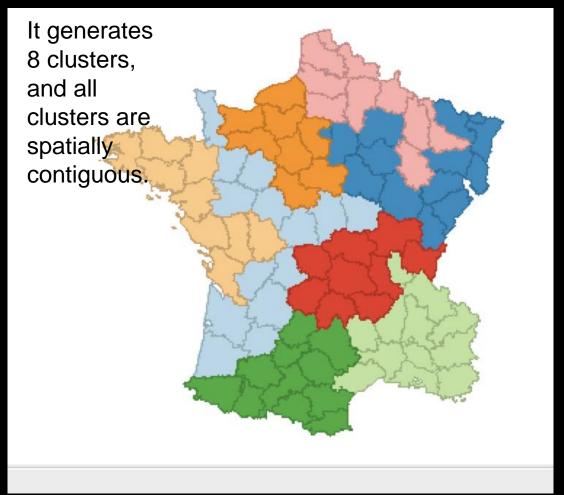




## Comparing kmeans and max-p

Method 3: max-p method (each region has at least 10% of total pop)







## **Measuring Clustering Quality**

## Necessary when...

- Comparing different implementations of a clustering method with randomness (e.g., k-means)
- Comparing clustering with different parameters (e.g., numbers of clusters)
- Comparing different clustering methods



## **Method 1: SEE / Elbow Method**

SSE: Sum of Squared Errors

$$SSE = \sum_{i=1}^{n} \sum_{j=1}^{k} w^{(i,j)} dist(x^{(i)}, \mu^{(j)})$$

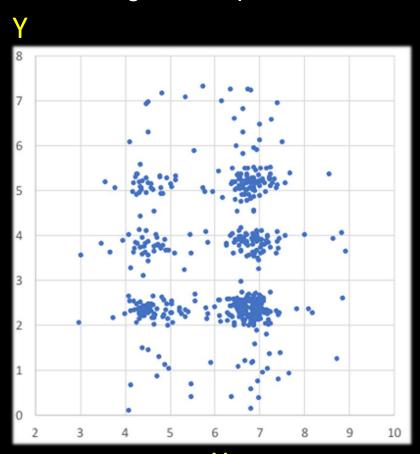
Where: i is a data point, j is a cluster, and  $\mu^{(j)}$  is the centre of a cluster. w(i,j)=1 when i is in cluster j, otherwise 0.

- The range of SSE? [0, infinity)
- A small SSE means that the data points are close to cluster centre and the clustering has good performance.



## **Method 1: SEE / Elbow Method**

Elbow diagram: help choose k for k-means







## Method 2: Silhouette Analysis

Silhouette of a point

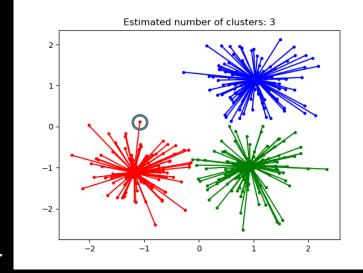
"Is this point closer to points of the same cluster, or any other cluster?"

$$s(i) = \frac{b(i) - a(i)}{\max\{a(i), b(i)\}}$$



b(i): minimum mean distance to points of another cluster

$$-1 \le s(i) \le 1$$
; the larger  $s(i)$ , the higher clustering quality



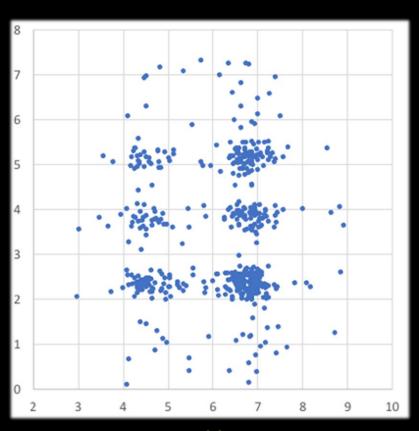
Silhouette Score for a Clustering

Average of s(i) for all points i

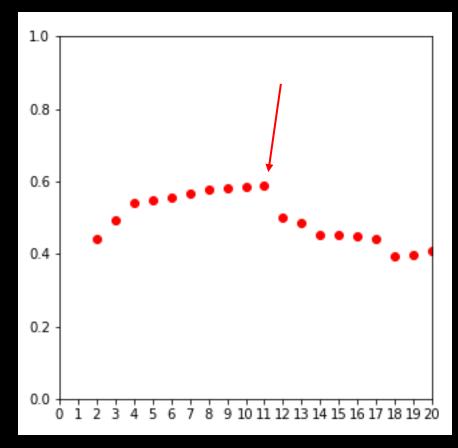


## Method 2: Silhouette Analysis

Choose k for k-means

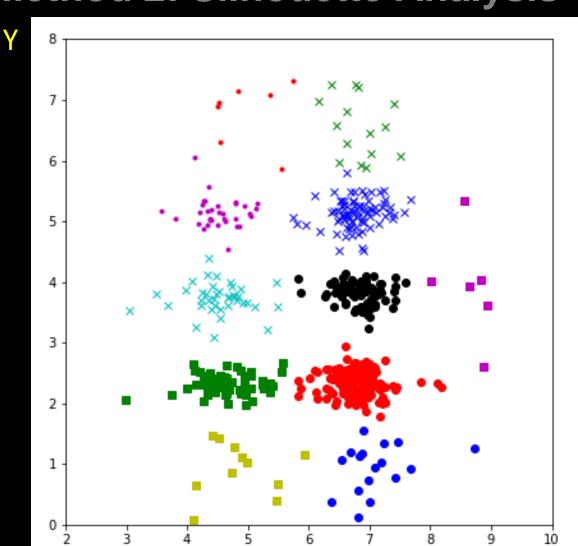


## Silhouette Score





## Method 2: Silhouette Analysis



'Optimal' k-Means

$$k = 11$$
  
S. Score = 0.59



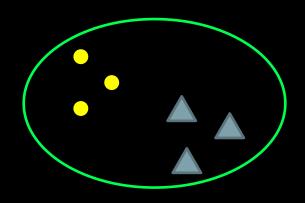
Homogeneity

All clusters contain only points from a single observed class – expressed as a proportion of clusters for which this is true

Completeness

All members of given class are within the same cluster – expressed as a proportion of classes for which this is true

#### Scenario 1



	C1	C2	Average
Homogeneity	1	1	1
	OC1		Average
Completeness	0		0

Clustering

Observed class (ground truth) OC1



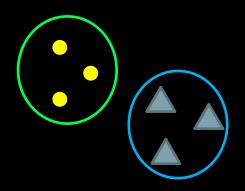
Homogeneity

All clusters contain only points from a single observed class – expressed as a proportion of clusters for which this is true

**Completeness** 

All members of given class are within the same cluster – expressed as a proportion of classes for which this is true

### Scenario 2



	C1	C2	Average
Homogeneity	1	1	1
			_
	OC1	OC2	Average

Clustering

Observed class (ground truth)

C

OC1

 $\triangle$  C2

OC2



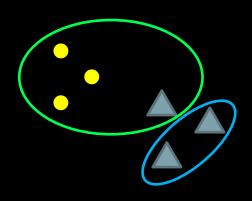
Homogeneity

All clusters contain only points from a single observed class – expressed as a proportion of clusters for which this is true

Completeness

All members of given class are within the same cluster – expressed as a proportion of classes for which this is true

#### **Scenario 3**



	C1	C2	Average
Homogeneity	1	0	0.5
	OC1	OC2	Average
Completeness	0	1	0.5

Clustering

Observed class (ground truth)

C1

C2

OC1

OC2



- Where is the 'ground truth' from?
  - You have some ground truth available;
  - The 'ground truth' can come from a different but relevant task. Should prove these tasks are relevant.
  - You can ask some domain experts for their opinions.
     This is very common and useful.



## **Measuring Clustering Quality**

 If you use the methods above to choose the hyperparameters (e.g. k of kmeans), the result of these methods might be different. You can simply use one method to determine the k value.



## **Next steps of clustering**

- Visualisation (often combined with dimension reduction, e.g. PCA)
- Qualitatively describe cluster characteristics
- Mapping the clusters (do these clusters cluster in space?)
- Compare against expert knowledge





## Workshop

#### **Dimension reduction**

- Weekly quiz on Moodle: please finish them before the workshop and we will discuss the quiz in the workshop
- Python notebooks for workshop: will be ready by 5pm Thursday.
- See you in the workshop on Friday 1-3pm