

CLUSTERING



FOR LATER
DOWNLOAD THE
“Exercise - Clustering.zip”
NOTEBOOK & DATASET
FROM THE COURSE SITE

CLUSTER ANALYSIS FINDS 'INTERESTING' GROUPS OF OBJECTS BASED ON SIMILARITY

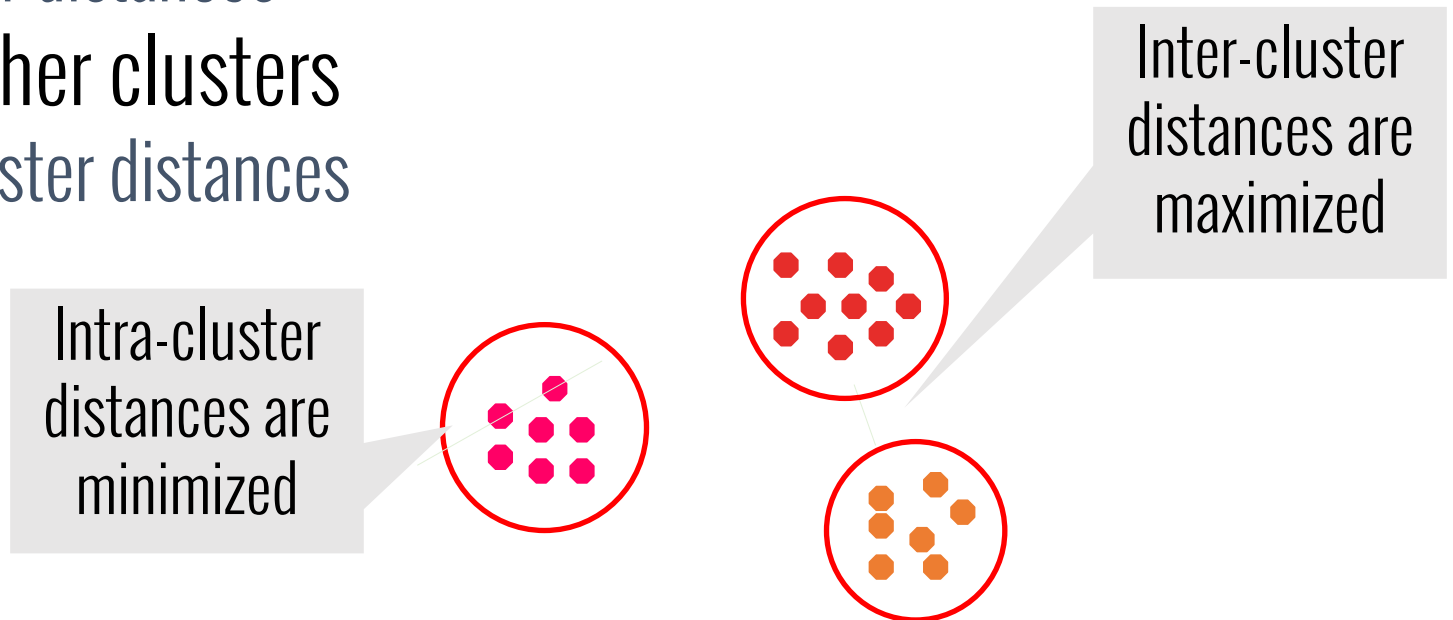
What typically makes a 'good' clustering?

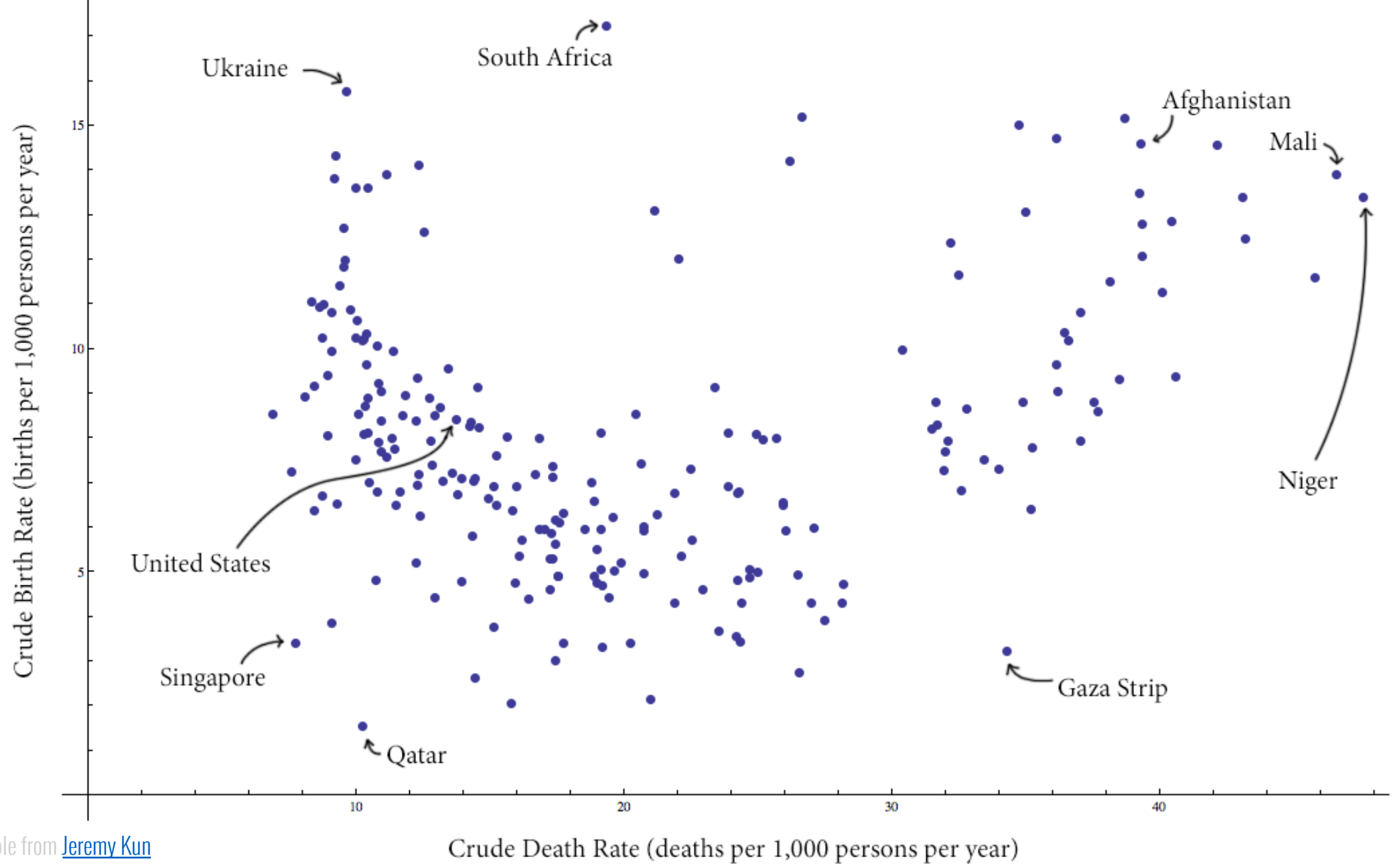
Members are highly similar to each other

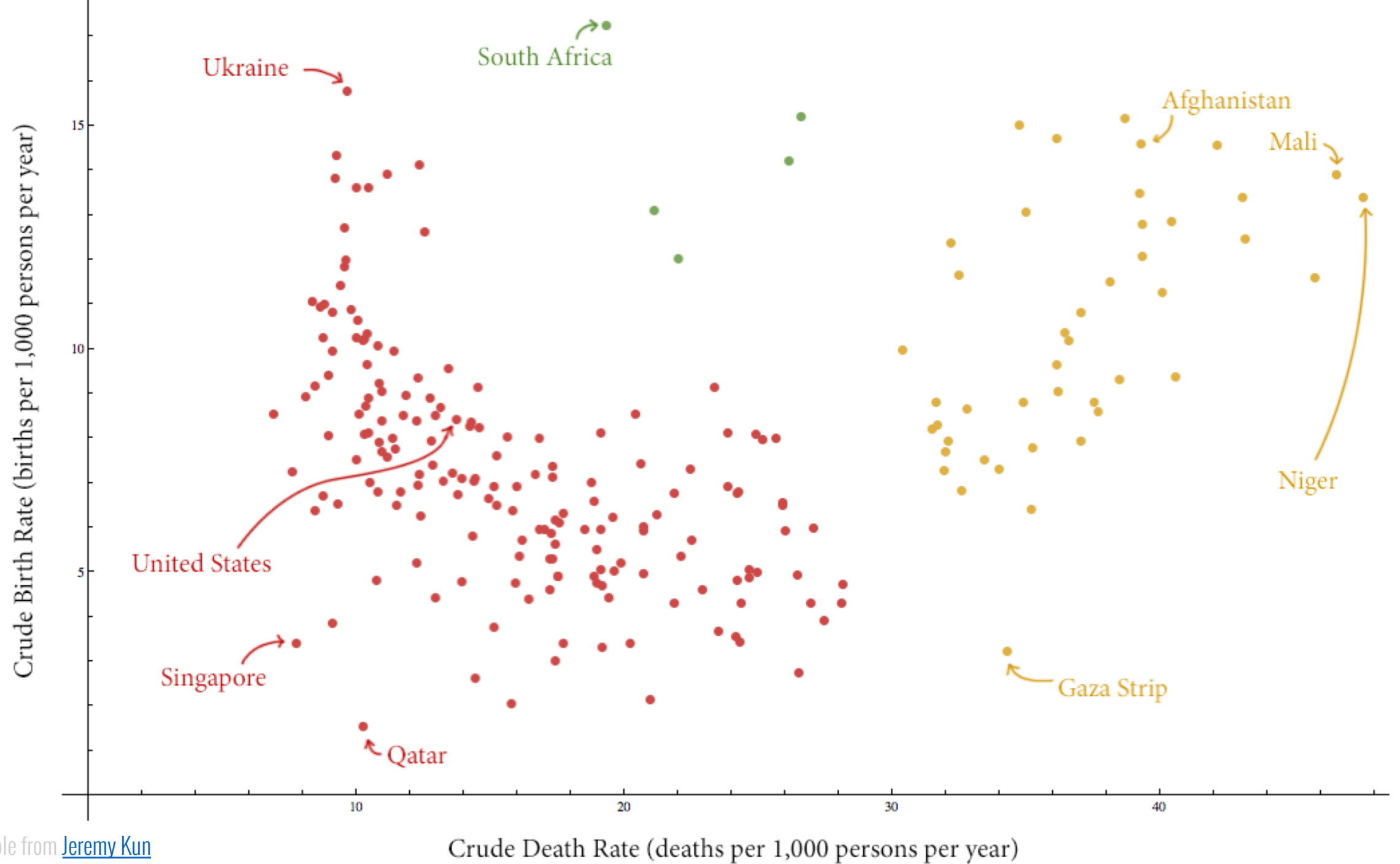
Minimize within-cluster distances

Well-separated from other clusters

Maximize between-cluster distances







APPLICATIONS OF CLUSTER ANALYSIS

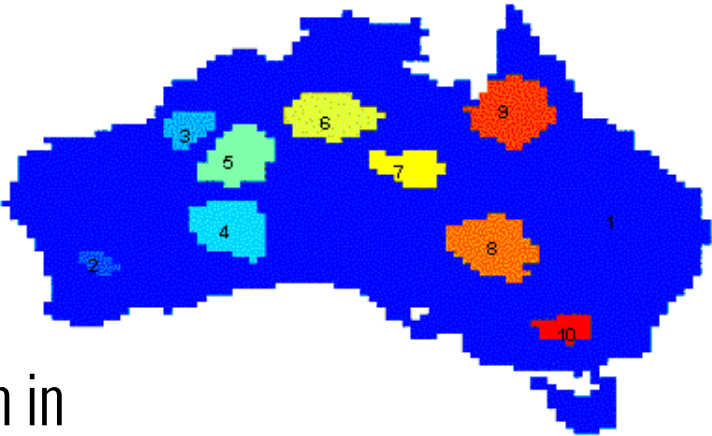
Understanding

- Group related documents for browsing
- Group genes and proteins with similar functionality
- Group stocks with similar price fluctuations
- ...

	Discovered Clusters	Industry Group
1	Applied-Matl-DOWN,Bay-Network-Down,3-COM-DOWN, Cabletron-Sys-DOWN,CISCO-DOWN,HP-DOWN, DSC-Comm-DOWN,INTEL-DOWN,LSI-Logic-DOWN, Micron-Tech-DOWN,Texas-Inst-Down,Tellabs-Inc-Down, Natl-Semiconduct-DOWN,Oracl-DOWN,SGI-DOWN, Sun-DOWN	Technology1-DOWN
2	Apple-Comp-DOWN,Autodesk-DOWN,DEC-DOWN, ADV-Micro-Device-DOWN,Andrew-Corp-DOWN, Computer-Assoc-DOWN,Circuit-City-DOWN, Compaq-DOWN, EMC-Corp-DOWN, Gen-Inst-DOWN, Motorola-DOWN,Microsoft-DOWN,Scientific-Atl-DOWN	Technology2-DOWN
3	Fannie-Mae-DOWN,Fed-Home-Loan-DOWN, MBNA-Corp-DOWN,Morgan-Stanley-DOWN	Financial-DOWN
4	Baker-Hughes-UP,Dresser-Inds-UP,Halliburton-HLD-UP, Louisiana-Land-UP,Phillips-Petro-UP,Unocal-UP, Schlumberger-UP	Oil-UP

Summarization

- Reduce size of large data sets



Clustering
precipitation in
Australia

RELATIONSHIP TO OTHER APPROACHES

	continuous	categorical
supervised	regression	classification
unsupervised	dimension reduction	clustering

SUMMARY: CONDUCTING CLUSTER ANALYSIS

Formulate the Problem

Select a Distance/Similarity Measure

Select a Clustering Procedure

Decide on the Number of Clusters

Interpret and Profile Clusters

Assess the Validity of Clustering

CLUSTERING IS OFTEN USED AS AN EXPLORATORY DATA ANALYSIS TOOL

Data understanding

Finding underlying factors, groups, structure

Data navigation

Creating hierarchies to support browsing

Data reduction

Clustering creates a new nominal variable that can be used in any further analysis.

A good way to quantize variable measures into non-uniform buckets

Data cleaning / smoothing

Infer or interpolate missing attributes from cluster neighbors

CLUSTERING ARISES NATURALLY IN MANY FIELDS

Business

- Market segments
- Web site visitors

Social network analysis

- Find communities

Information Retrieval

- Search results clustered by similarity, event or topic
- Personalization for groups of similar users

Health

- DNA gene expression
 - Cluster cancer variants into treatment groups, based on immunomarkers of cell samples
- Medical imaging
 - Find likely tumors

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SELECTING A DISTANCE METRIC

Your distance function determines the clusters you'll get

Can be formulated in different (opposite) ways.

Maximize *intra-cluster similarity* while minimizing *inter-cluster similarity*
Minimize *intra-cluster distances* while maximizing *inter-cluster distances*

But what's the right metric for distance/similarity?

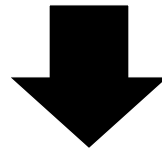
How do we calculate *genetic similarity*?

How do we calculate the *similarity in musical taste*?

How do we calculate the *similarity in car features*?

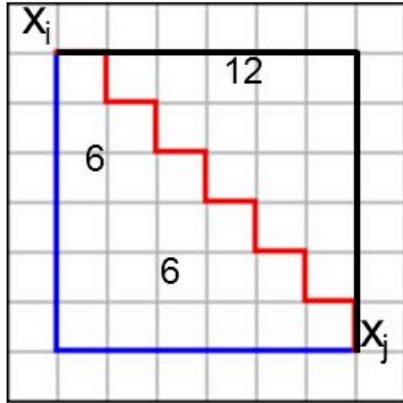
DATAFRAME TO SIMILARITY MATRIX

Entry #	Variable 1	Variable 2	Variable 3
1	A	True	3.5
2	A	False	4.5
3	B	True	5.6



	1	2	3
1	Sim(1,1)	Sim(1,2)	Sim(1,3)
2	Sim(2,1)	Sim(2,2)	Sim(2,3)
3	Sim(3,1)	Sim(3,2)	Sim(3,3)

SOME OPTIONS

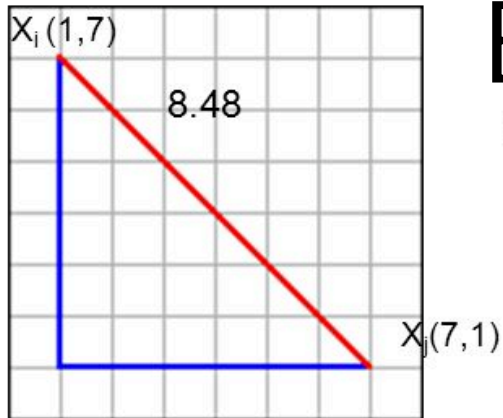


Manhattan distance

Think of it as city grids

$$|x_1 - x_2| + |y_1 - y_2| + |z_1 - z_2| + \dots$$

Not often used



Euclidean distance

$$\sqrt{(x_1 - x_2)^2 + (y_1 - y_2)^2 + (z_1 - z_2)^2 + \dots}$$

Good for: quantitative data, equal importance/weights, normalized data

SOME OPTIONS

Jaccard distance

When data is nominal (true/false, words, etc.)

$$\frac{|intersection(A, B)|}{|union(A, B)|}$$

“How many things we have in total versus how many things we have in common”

YOU HAVE LOTS OF CHOICES

For example, SciKit-Learn supports:

Metrics intended for real-valued vector spaces:

identifier	class name	args	distance function
"euclidean"	EuclideanDistance	•	$\sqrt{\sum (x - y)^2}$
"manhattan"	ManhattanDistance	•	$\sum x - y $
"chebyshev"	ChebyshevDistance	•	$\max(x - y)$
"minkowski"	MinkowskiDistance	p	$\sum x - y ^p^{(1/p)}$
"wminkowski"	WMinkowskiDistance	p, w	$\sum (w * x - y ^p)^{(1/p)}$
"seuclidean"	SEuclideanDistance	V	$\sqrt{\sum (x - y)^2 / V}$
"mahalanobis"	MahalanobisDistance	V or VI	$\sqrt{(x - y)' V^{-1} (x - y)}$

Metrics intended for two-dimensional vector spaces: Note that the haversine distance metric requires data in the form of [latitude, longitude] and both inputs and outputs are in units of radians.

identifier	class name	distance function
"haversine"	HaversineDistance	$2 \arcsin(\sqrt{\sin^2(0.5 * dx) + \cos(x1) \cos(x2) \sin^2(0.5 * dy)})$

Metrics intended for integer-valued vector spaces: Though intended for integer-valued vectors, these are also valid metrics in the case of real-valued vectors.

identifier	class name	distance function
"hamming"	HammingDistance	$N_{\text{unequal}}(x, y) / N_{\text{tot}}$
"canberra"	CanberraDistance	$\text{sum}(x - y / (x + y))$
"braycurtis"	BrayCurtisDistance	$\text{sum}(x - y) / (\text{sum}(x) + \text{sum}(y))$

Metrics intended for boolean-valued vector spaces: Any nonzero entry is evaluated to "True". In the listings below, the following abbreviations are used:

- N : number of dimensions
- NTT : number of dims in which both values are True
- NTF : number of dims in which the first value is True, second is False
- NFT : number of dims in which the first value is False, second is True
- NFF : number of dims in which both values are False
- NNEQ : number of non-equal dimensions, $NNEQ = NTF + NFT$
- NNZ : number of nonzero dimensions, $NNZ = NTF + NFT + NTT$

identifier	class name	distance function
"jaccard"	JaccardDistance	$NNEQ / NNZ$
"matching"	MatchingDistance	$NNEQ / N$
"dice"	DiceDistance	$NNEQ / (NTT + NNZ)$
"kulsinski"	KulsinskiDistance	$(NNEQ + N - NTT) / (NNEQ + N)$
"rogerstanimoto"	RogersTanimotoDistance	$2 * NNEQ / (N + NNEQ)$
"russellrao"	RussellRaoDistance	NNZ / N
"sokalmichener"	SokalMichenerDistance	$2 * NNEQ / (N + NNEQ)$
"sokalsneath"	SokalSneathDistance	$NNEQ / (NNEQ + 0.5 * NTT)$

SIMILARITY METRIC CHOICES

Try to use metrics that you **understand** and **can reason about!**

Very often you'll **roll your own** or use a known similarity/distance metric for your specific type of data.

Need to be careful if the metric is
or **similarity** (0 is far, 1 is close)
dissimilarity (1 is far, 0 is close)

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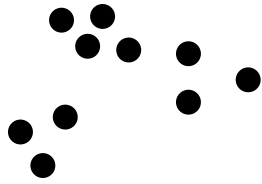
Decide on the Number of Clusters

Interpret and Profile Clusters

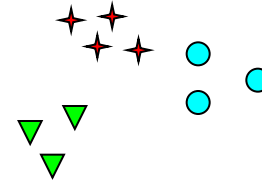
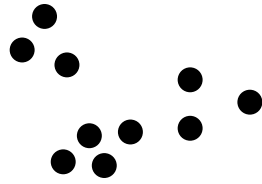
Assess the Validity of Clustering

SELECTING A CLUSTERING TECHNIQUE

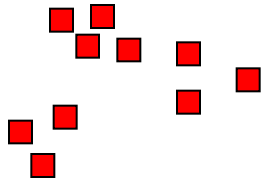
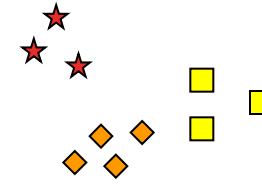
CLUSTERING CAN BE AMBIGUOUS: WHAT IS THE 'BEST' CLUSTERING HERE?



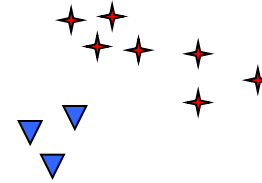
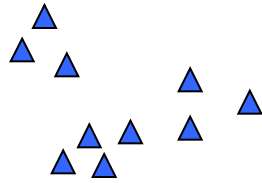
How many clusters?



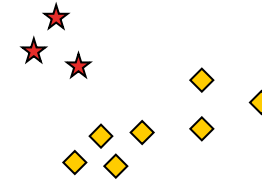
Six Clusters



Two Clusters



Four Clusters



THERE ARE LOTS OF CLUSTERING APPROACHES

Assigning objects to clusters

‘Hard’ (partitional) each object belongs to exactly 1 cluster

‘Soft’ : each object can belong to multiple clusters

Hierarchical vs non-hierarchical

A set of nested clusters organized as a tree

By far most widely-used fall into two types:

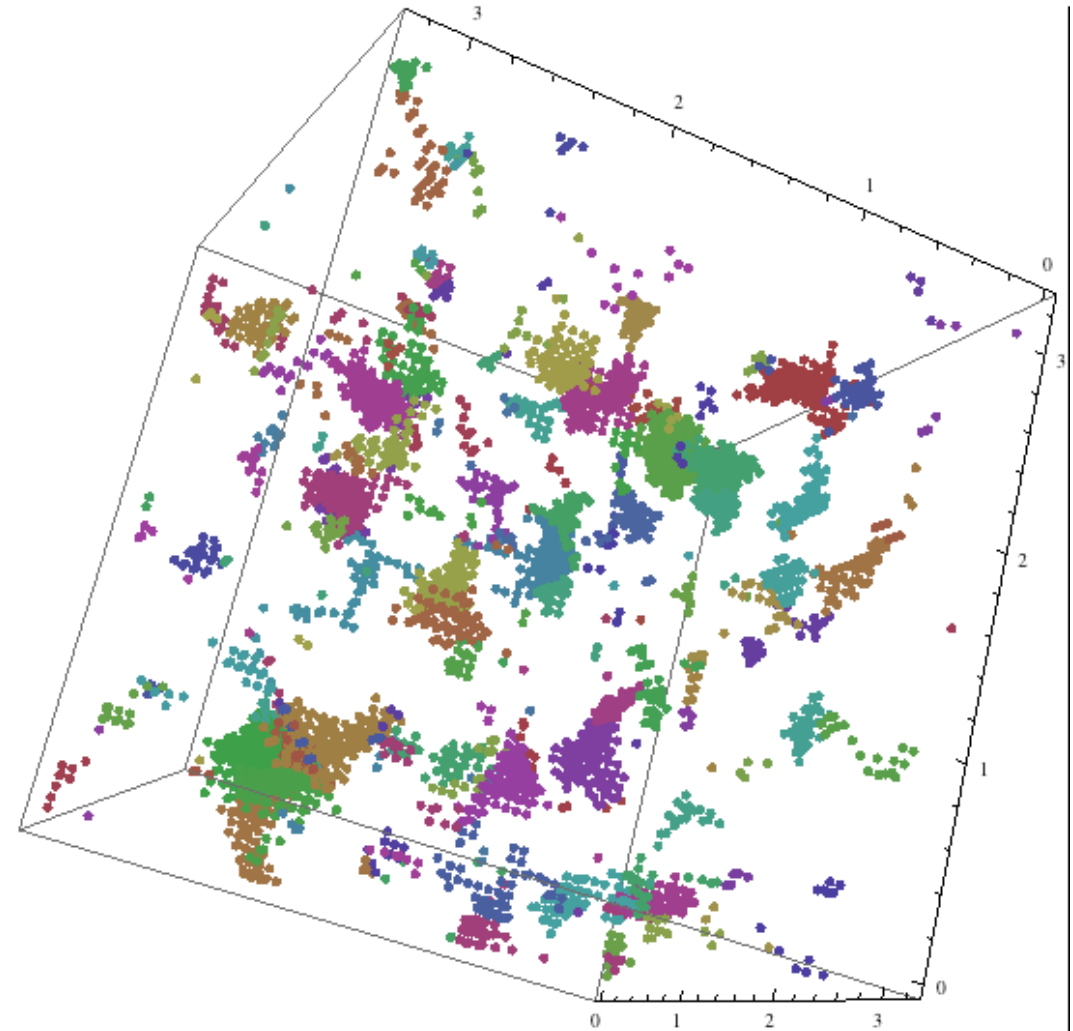
Hierarchical: agglomerative, single-link, etc.

Partitional: k-means, k-median, etc.

CLUSTERING AND DIMENSIONALITY

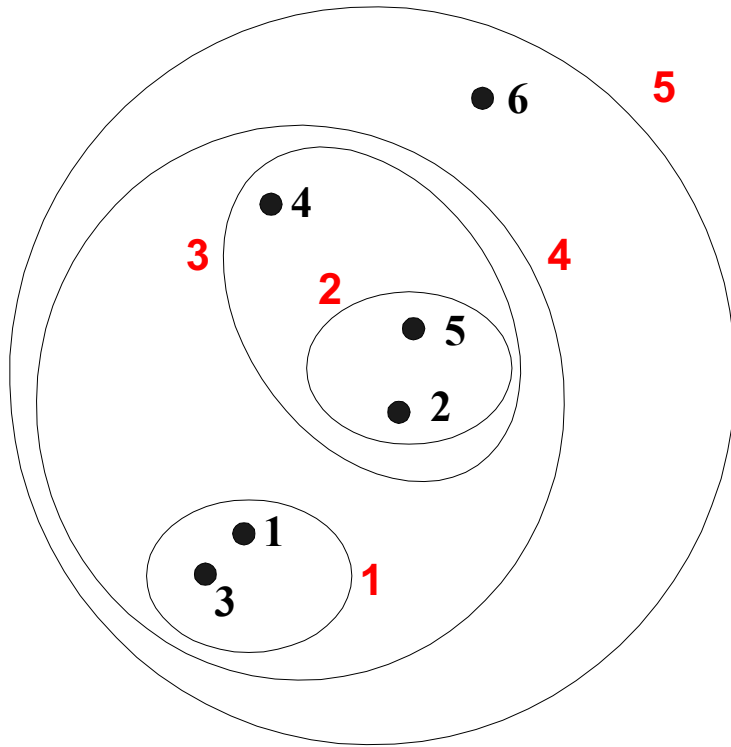
Most of our examples will be 2D
(its easy to illustrate).

But remember that your
distance/similarity metrics can
include **as many dimensions as
you want!**

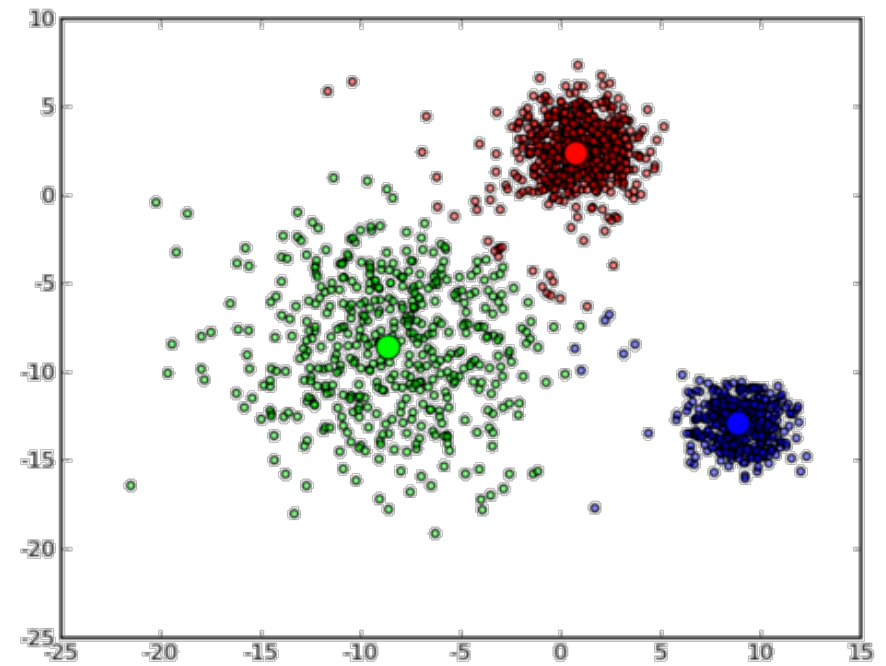


TODAY

HIERARCHICAL CLUSTERING



K-MEANS CLUSTERING

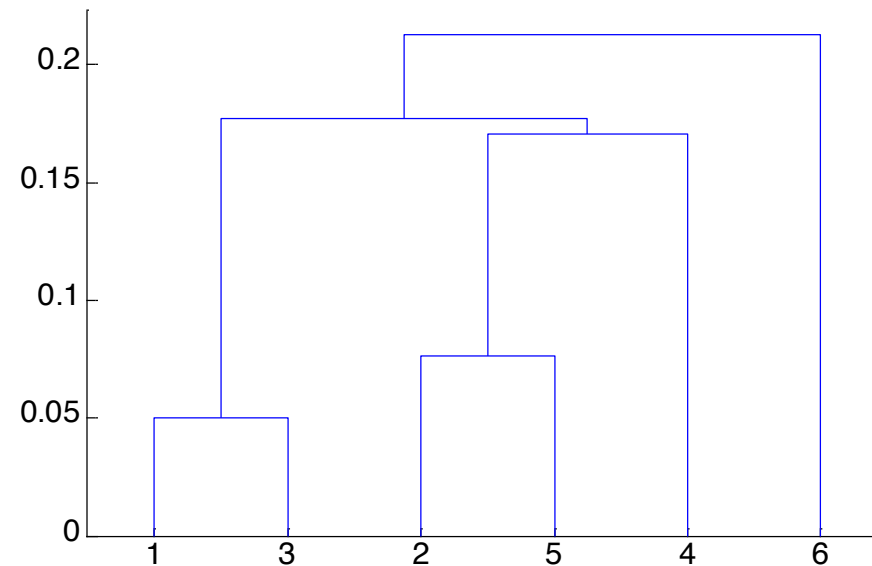
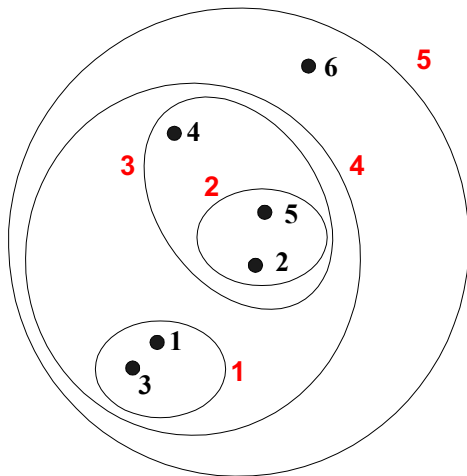


HIERARCHICAL CLUSTERING

Produces a set of nested clusters organized as a **hierarchical tree**

Can be visualized as a **dendrogram**

A tree like diagram that records the sequences of merges or splits



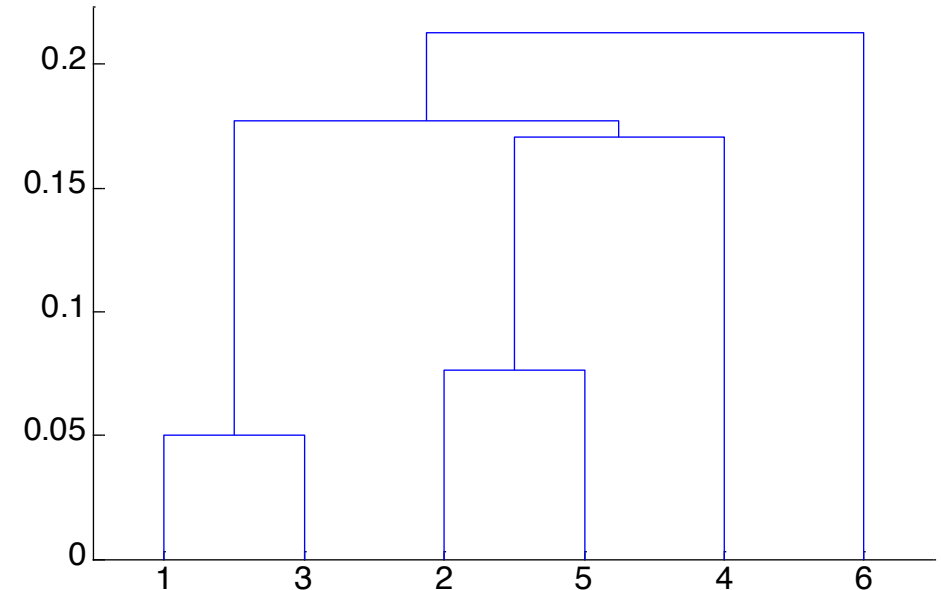
STRENGTHS OF HIERARCHICAL CLUSTERING

Do not have to assume any particular number of clusters

Any desired number of clusters can be obtained by 'cutting' the dendrogram at the proper level

They may correspond to meaningful taxonomies

Example in biological sciences (e.g., animal kingdom, phylogeny reconstruction, ...)



HIERARCHICAL CLUSTERING

Bottom-up ('Agglomerative')

- Start with each point being in its own cluster

- At each step

 - Merge the most similar pair of clusters based on a cost function

 - Continue until you have k clusters, or everything is in one big cluster

Top-down ('Divisive')

- Start with all points in a single big cluster

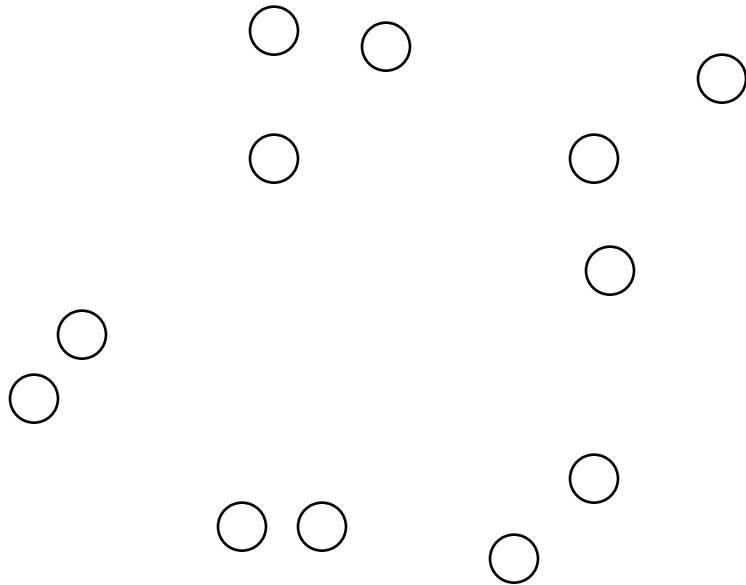
- At each step:

 - Split the cluster into two smaller clusters based on a cost function

 - Continue until you have k clusters, or each point is in its own cluster

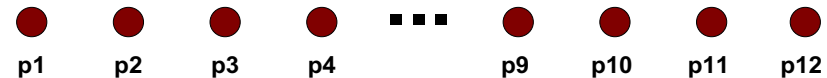
AGGLOMERATIVE (BOTTOM-UP) CLUSTERING: STARTING SITUATION

Start with clusters of individual points and a proximity matrix of object-to-object distances



	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						
.						

Similarity Matrix



AGGLOMERATIVE CLUSTERING ALGORITHM

One popular hierarchical clustering technique

Basic algorithm is straightforward

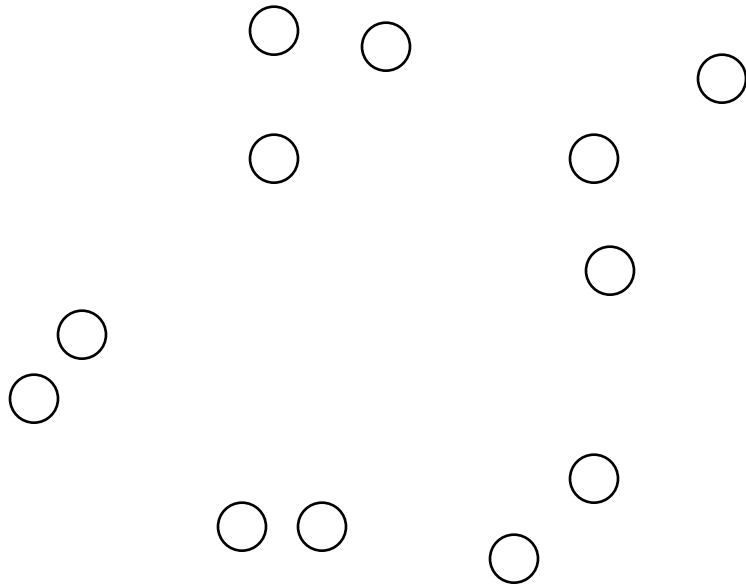
1. Compute the proximity matrix
2. Let each data point be a cluster
3. Repeat:
 4. Merge the two closest clusters
 5. Update the proximity matrix
 6. Stop if only a single cluster remains

For this discussion
“proximity” = “similarity” = “distance”

Key operation: computation of the proximity of two clusters. The cost function.
Different approaches to defining the distance between clusters distinguish the different algorithms

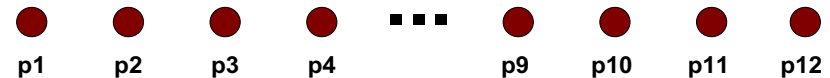
AGGLOMERATIVE (BOTTOM-UP) CLUSTERING

Start with clusters of individual points and a similarity matrix of object-to-object distances



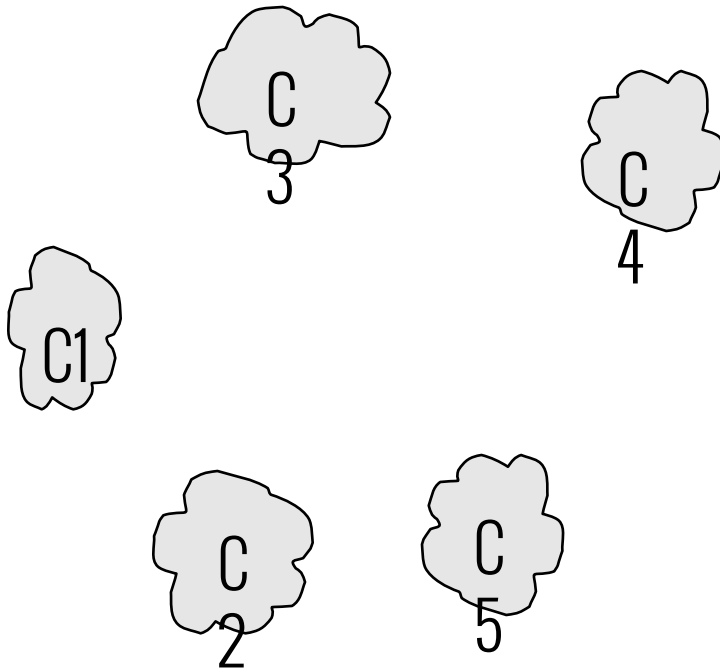
	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						
.						

Similarity Matrix



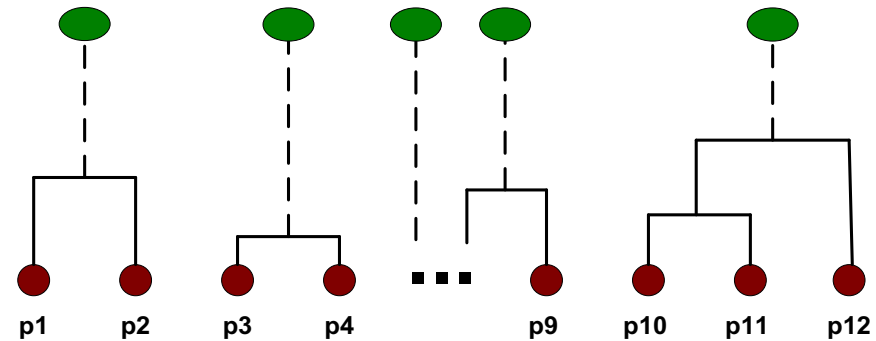
INTERMEDIATE SITUATION

After some merging steps, we have some clusters



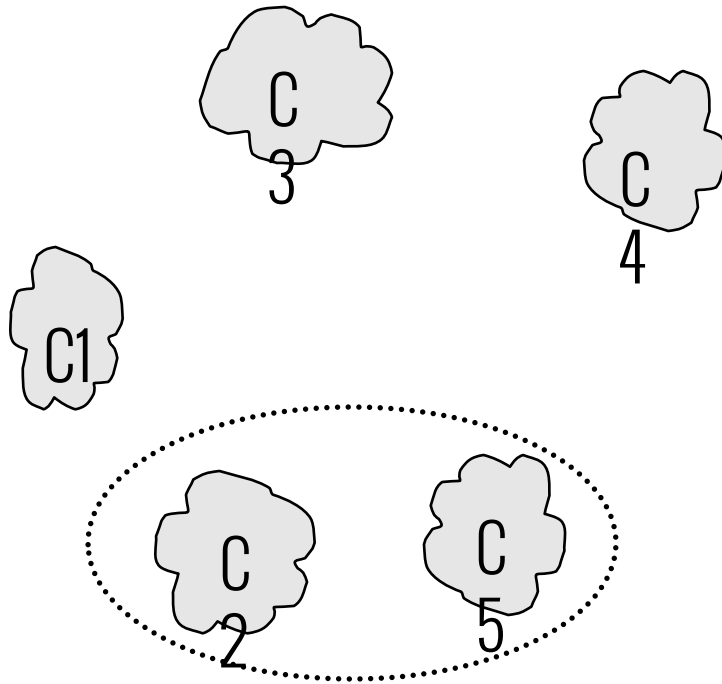
	C1	C2	C3	C4	C5
C1					
C2					
C3					
C4					
C5					

Similarity Matrix



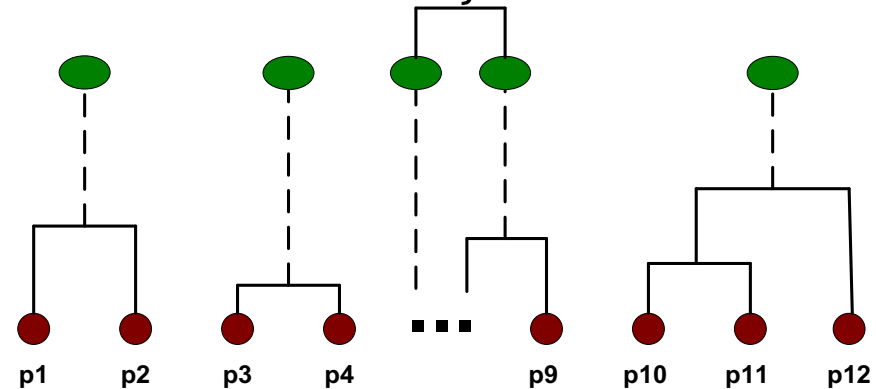
INTERMEDIATE SITUATION

We want to merge the two closest clusters (C2 and C5) and update the proximity matrix.



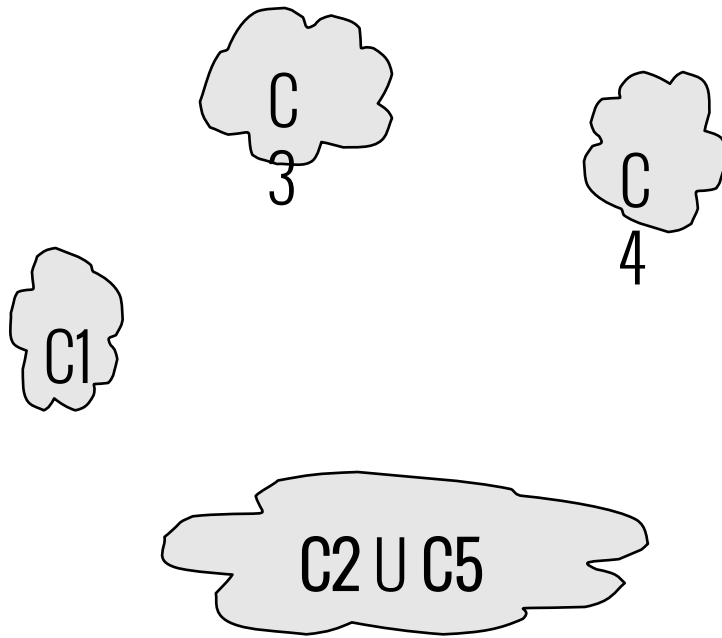
	C1	C2	C3	C4	C5
C1					
C2					
C3					
C4					
C5					

Similarity Matrix



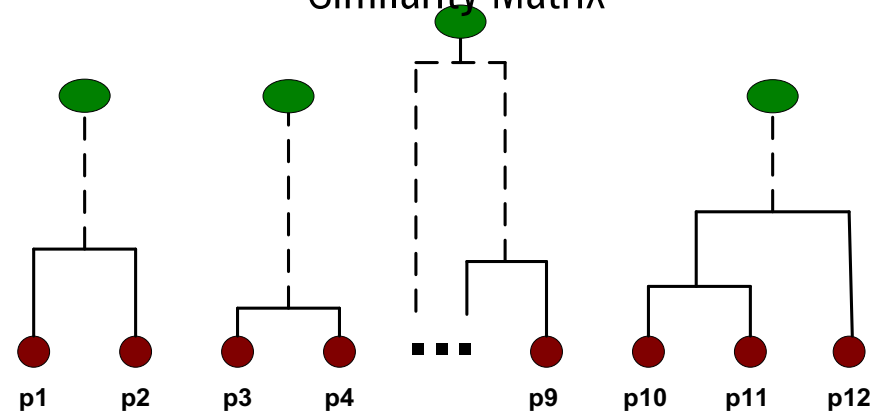
AFTER MERGING

The question is “How do we update the proximity matrix?”

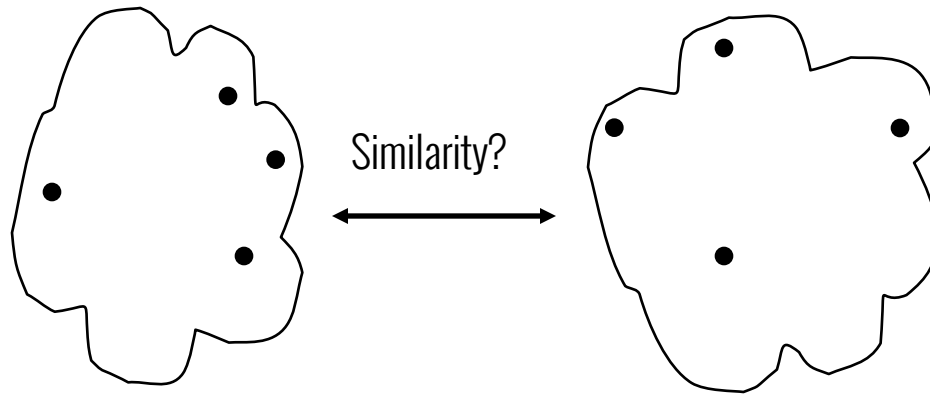


	C1	C2 + C5	C3	C4
C1				
C2 + C5				
C3				
C4				

Similarity Matrix



HOW TO DEFINE INTER-CLUSTER SIMILARITY



	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						

Similarity Matrix

MIN

MAX

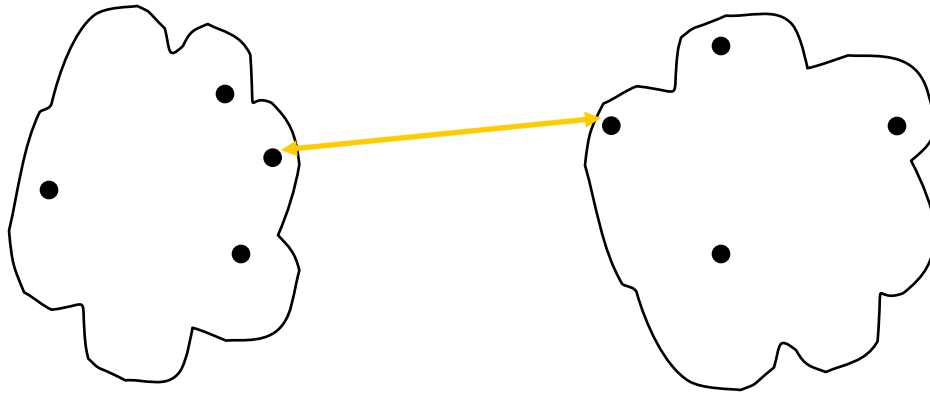
Group Average

Distance Between Centroids

Other methods driven by an objective function

Ward's Method uses squared error

HOW TO DEFINE INTER-CLUSTER SIMILARITY



	p1	p2	p3	p4	p5	...
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p2						
p3						
p4						
p5						
.						

Similarity Matrix

MIN

MAX

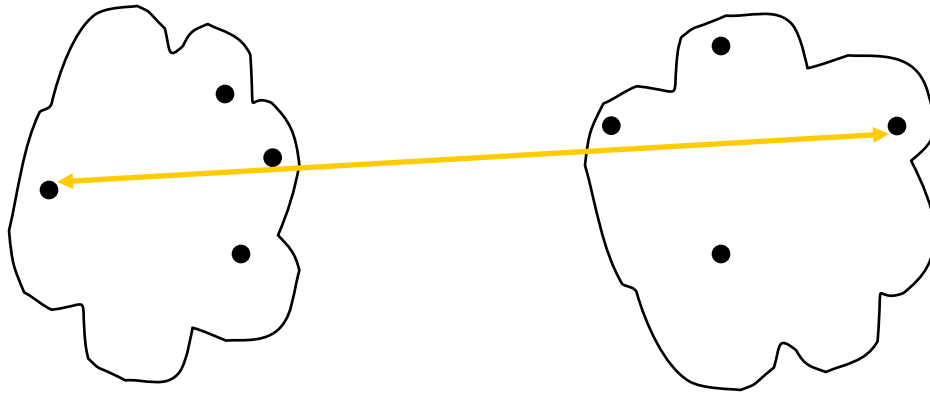
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HOW TO DEFINE INTER-CLUSTER SIMILARITY



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Similarity Matrix

MIN

MAX

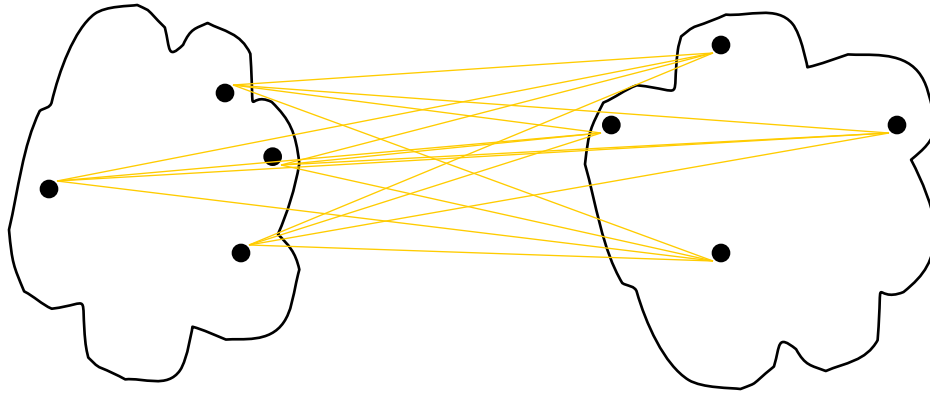
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HOW TO DEFINE INTER-CLUSTER SIMILARITY



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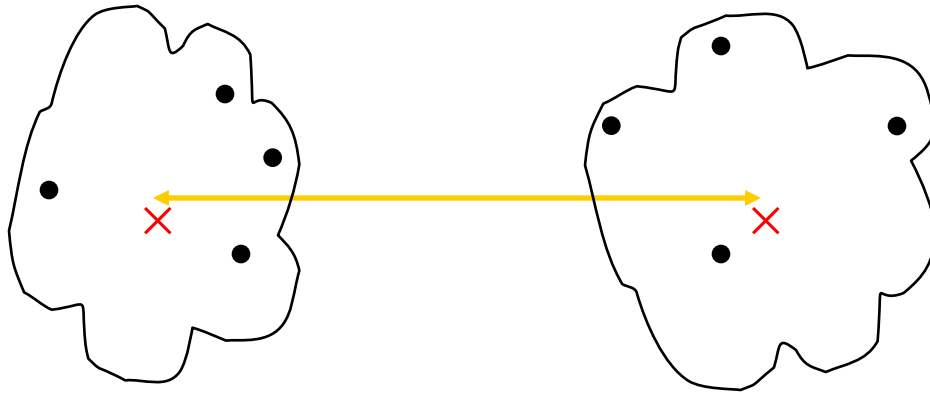
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HOW TO DEFINE INTER-CLUSTER SIMILARITY



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Similarity Matrix

MIN

MAX

Group Average

Distance Between Centroids

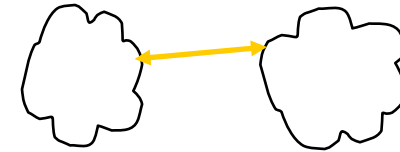
Other methods driven by an objective function

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COST FUNCTIONS FOR BOTTOM-UP (AGGLOMERATIVE) CLUSTERING

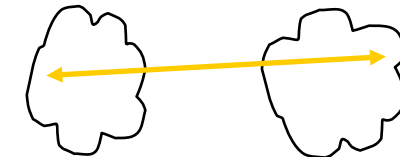
Single linkage

Minimum distance between clusters



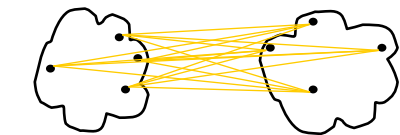
Complete linkage

Max distance between clusters



Average linkage

Average distance between clusters



WARD'S METHOD (1963)

Ward's distance between clusters C_i and C_j is the *difference* between the *total within cluster sum of squares for the two clusters separately*, and the *within cluster sum of squares resulting from merging the two clusters* in cluster C_{ij}

$$D_w(C_i, C_j) = \sum_{x \in C_i} (x - r_i)^2 + \sum_{x \in C_j} (x - r_j)^2 - \sum_{x \in C_{ij}} (x - r_{ij})^2$$

r_i : centroid of C_i

r_j : centroid of C_j

r_{ij} : centroid of C_{ij}

WARD'S DISTANCE FOR CLUSTERS

Similar to group average and centroid distance

Less susceptible to noise and outliers

Hierarchical analogue of k-means

Can be used to initialize k-means

WHICH TYPE OF HIERARCHICAL CLUSTERING TO USE?

Different methods have different strengths and weaknesses:

Ward's method tends to give equal sized clusters

Single linkage (nearest neighbor) tends to make long strings into a cluster.

Top-down is sensitive to early errors

Bottom-up can't see the whole dataset

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Formulate the Problem

Select a Distance/Similarity Measure

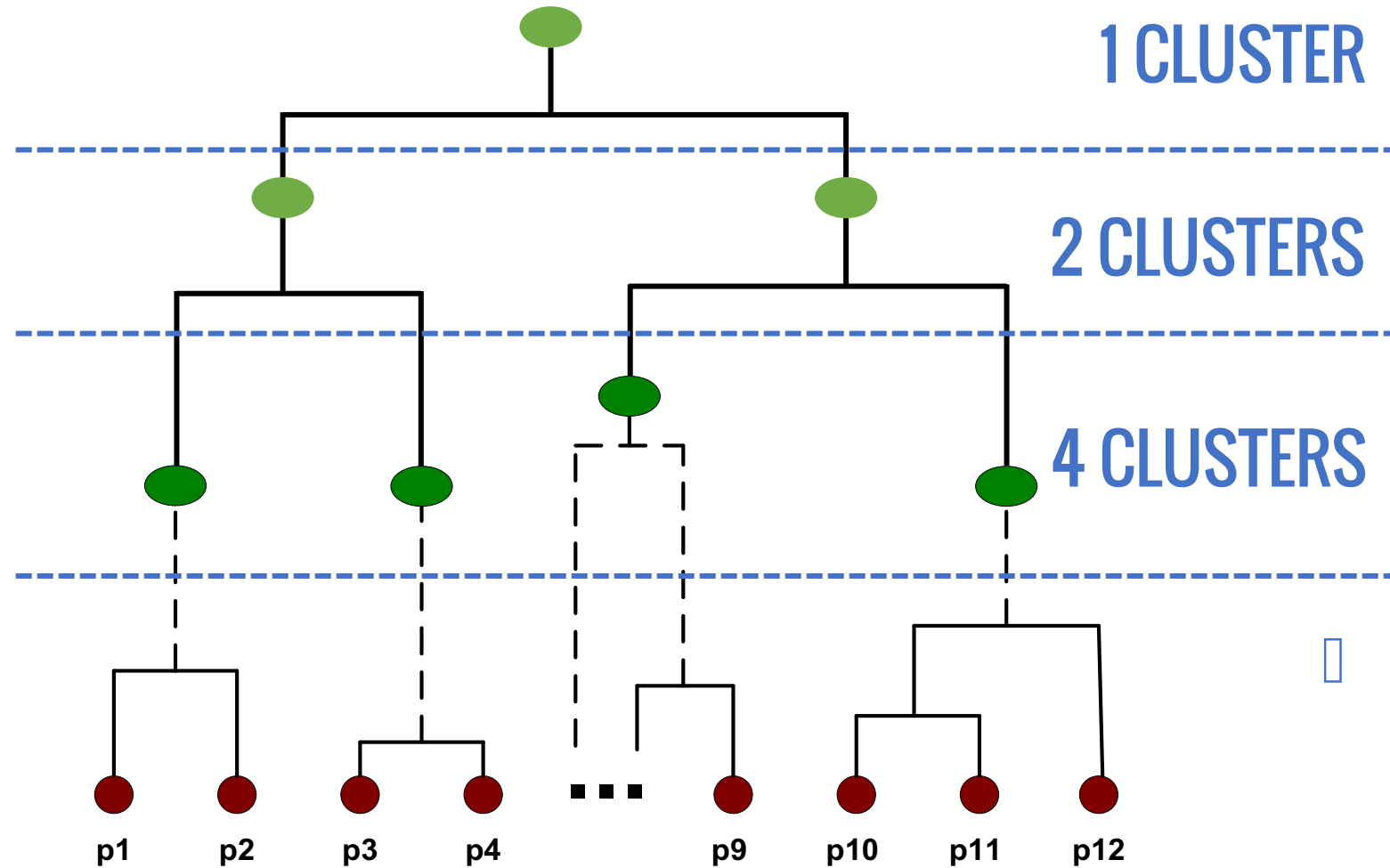
Select a Clustering Procedure

Decide on the Number of Clusters

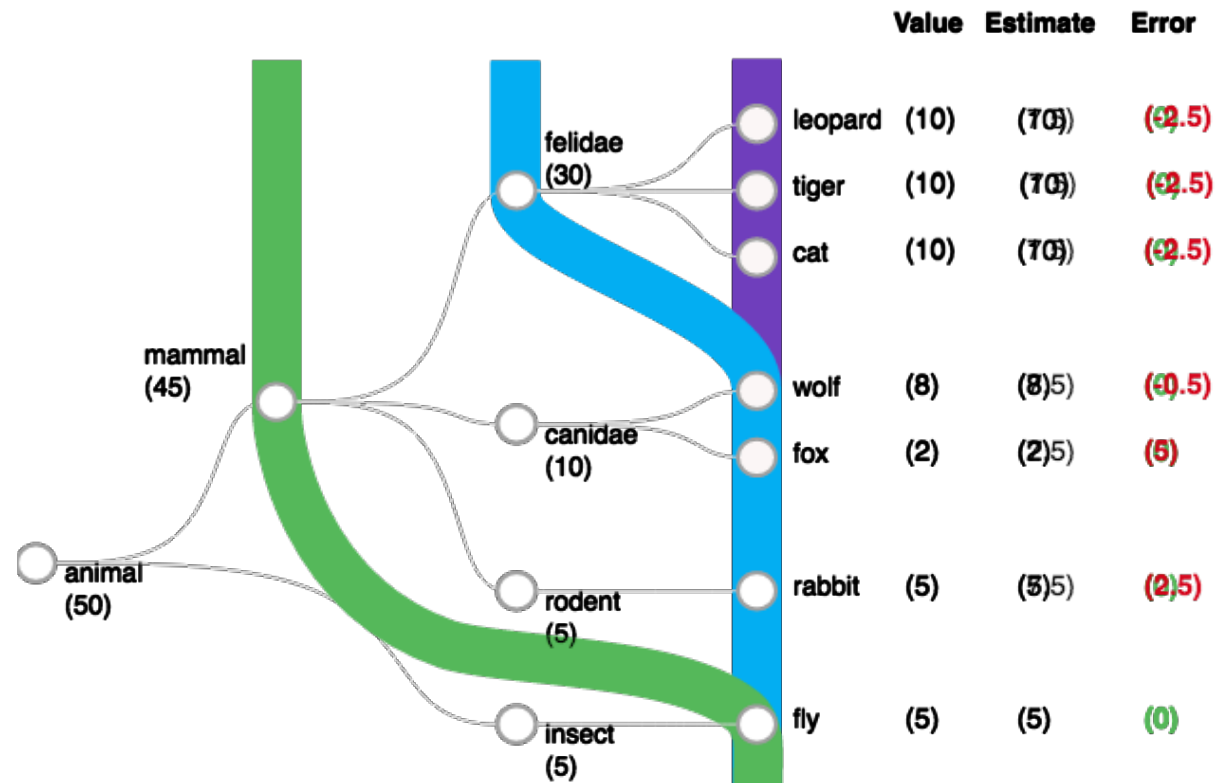
Interpret and Profile Clusters

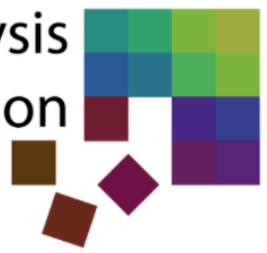
Assess the Validity of Clustering

CUTTING THE TREE INTO CLUSTERS



VARIABLE-DEPTH CUTS





k-means

Three Not Equal Circles

Custom Data

Parameters

k: 3

Distance
Measure:

Euclidian

$$d_2(x, y) = \sqrt{\sum_{i=1}^n |x_i - y_i|^2}$$

Navigation

Iteration #

8

Step #

15

Current Animation Speed

2



Current Step

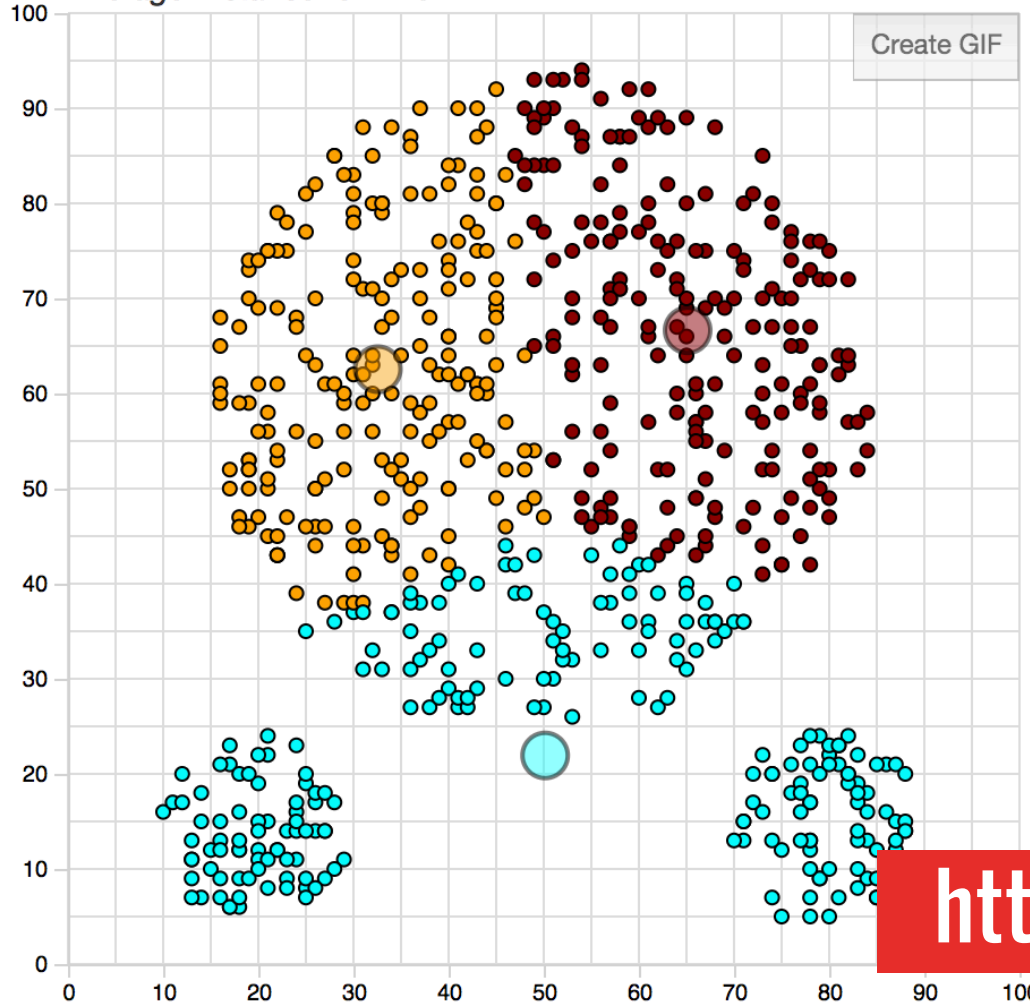
15

Visualization



Average Distance: 81.719

Create GIF



Algorithm: k-means

Complexity Range: $O(k \times n \times t)$

Input: k clusters

Output: k clusters

Pseudocode:

1. Choose k objects as initial **cluster centers**.
2. Assign each data point to the cluster which has the closest **mean point (centroid)** under chosen distance metric.
3. When all data points have been assigned, recalculate the positions of k **centroids (mean points)**.
4. Repeat steps 2 and 3 until the **centroids** do not change any more. All data points remain in their most recently assigned cluster.

<http://educlust.dbvis.de/>

K-MEANS

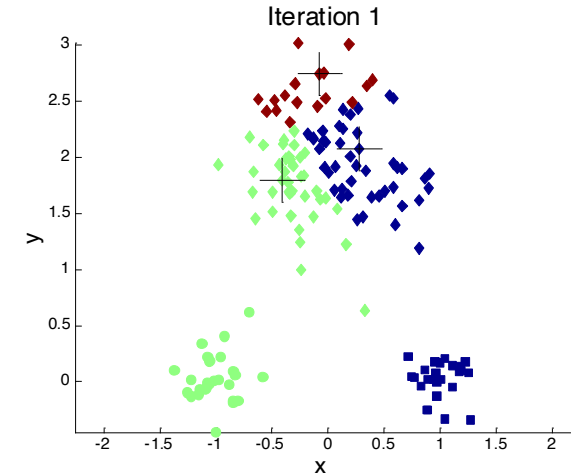
K-MEANS: THE OTHER MASSIVELY POPULAR CLUSTERING METHOD

Partitional clustering approach

Each cluster associated with a **centroid** (center point)

Each point is assigned to the cluster with the closest centroid

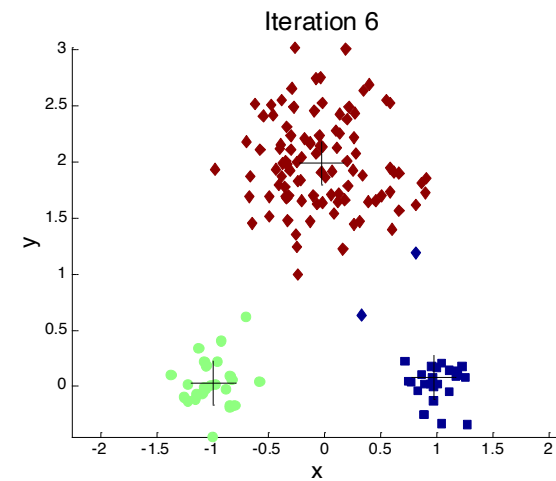
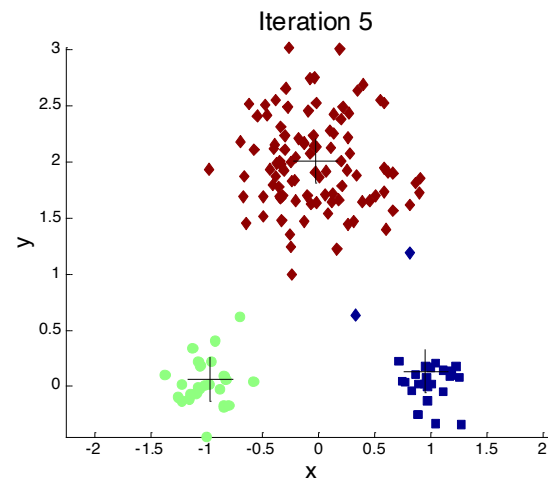
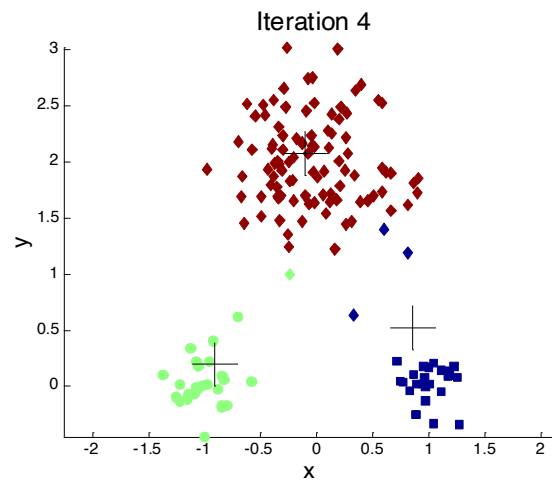
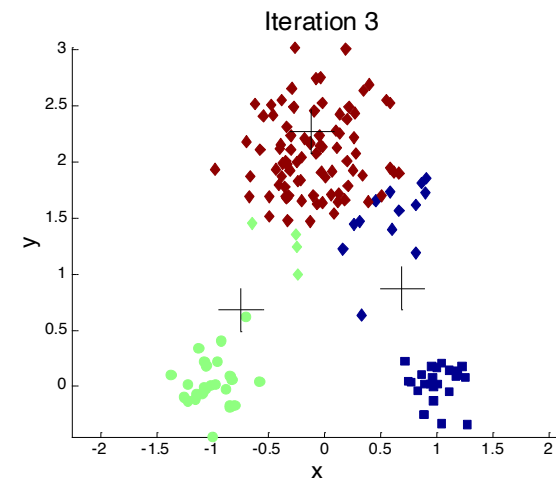
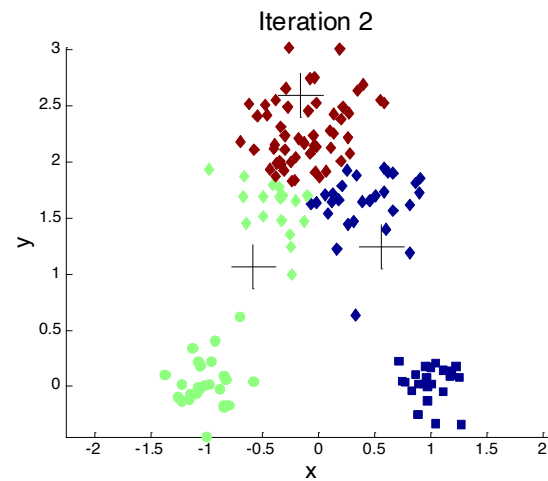
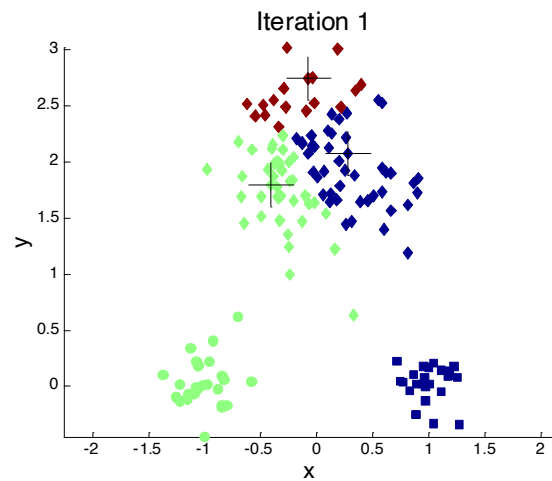
Number of clusters, **K**, must be specified in advance

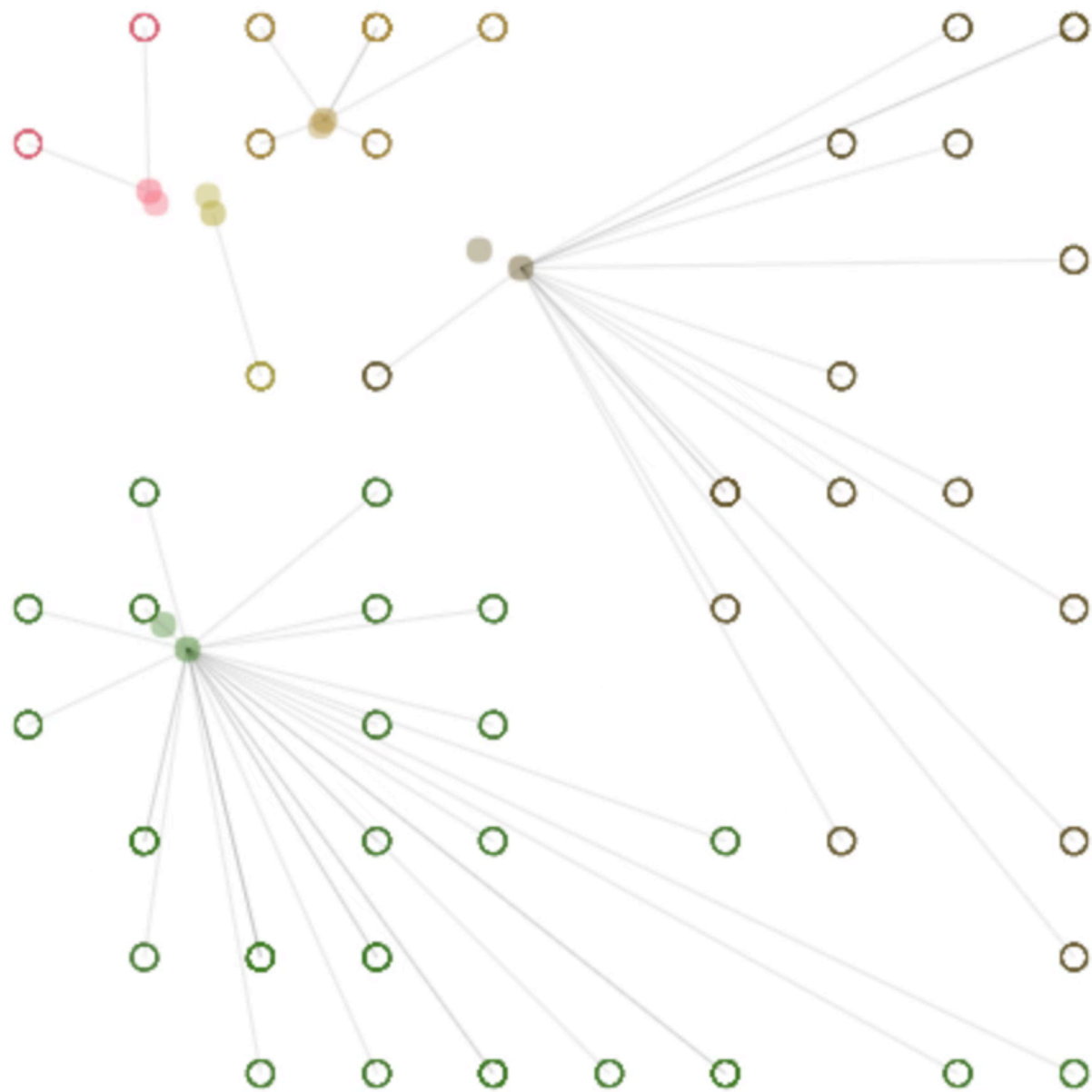


The basic algorithm is very simple

- 1: Select K points as the initial centroids.
- 2: **repeat**
- 3: Form K clusters by assigning all points to the closest centroid.
- 4: Recompute the centroid of each cluster.
- 5: **until** The centroids don't change

THE K-MEANS ALGORITHM (K = 3)





Miguel Mota
<https://goo.gl/LqhNUz>

K-MEANS CLUSTERING - DETAILS

Different initializations can result in different solutions

- Initial centroids are often chosen randomly.

- Clusters produced vary from one run to another.

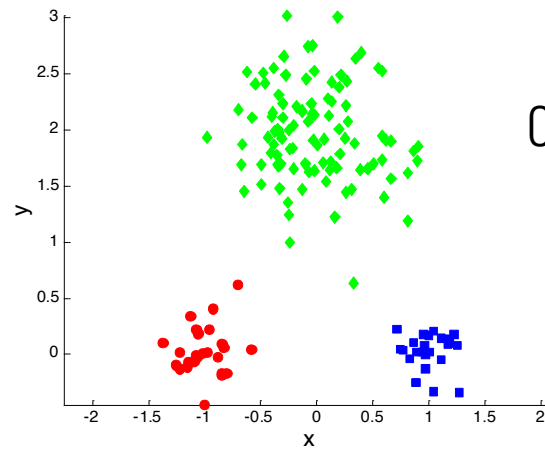
- So multiple runs are sometimes done

Centroid is typically the mean of the points in the cluster.

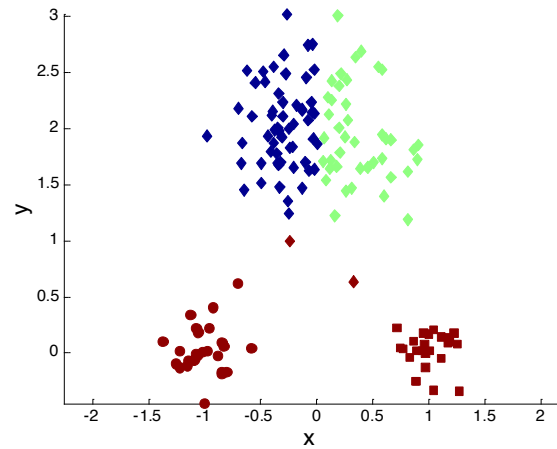
- “K-mediod” – center must be an actual datapoint.

- Useful when mean of a feature is not defined or available.

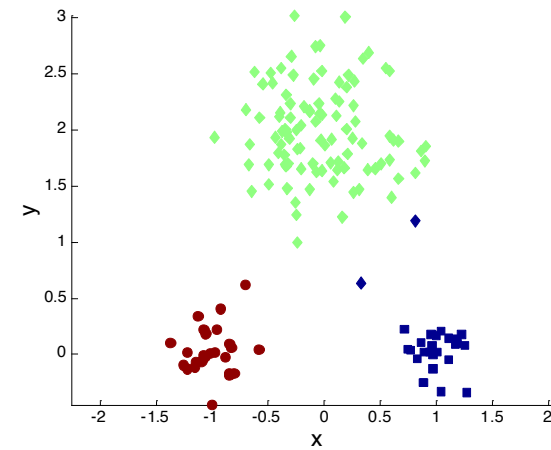
TWO DIFFERENT K-MEANS CLUSTERINGS



Original Points



Optimal Clustering? No.



Optimal Clustering

APPROACHES FOR IMPROVING K-MEANS

Idea 1: Be careful about where you start

Place first center on randomly chosen datapoint

Place second centroid on datapoint **as far as possible** from the first

Place n -th center on datapoint as far as possible from centers 1 thru $n - 1$

Idea 2: Do many runs of k-means

Each from a different random start configuration.

Use a heuristics to pick the best one.

LIMITATIONS OF K-MEANS

K-means has problems when clusters are of differing
Sizes
Densities

K-means has problems when the data contains **outliers**.

You **have** to pick the number of clusters (k) in advance.

WHEN TO USE K-MEANS VS HIERARCHICAL?

Do you need to easily interpret the clusters?

Do you know the right K ?

Does the data have a natural "tree" structure (living things, etc.)

How computationally expensive will each approach be for your data?

WHEN TO USE K-MEANS VS HIERARCHICAL?

k-means prefers solutions with similar sized clusters

very different cluster sizes, shapes, densities can confuse it
complex cluster geometry, or outliers
need to specify and test for good k choice

Can combine the two approaches – for example:

1. Try several hierarchical methods and see which gives the most interpretable clusters.
2. Use k-means (with the hierarchical cluster centroids as starting points) to clean up the hierarchical cluster.

SUMMARY: CONDUCTING CLUSTER ANALYSIS

Formulate the Problem

Select a Distance/Similarity Measure

Select a Clustering Procedure

Decide on the Number of Clusters

Interpret and Profile Clusters

Assess the Validity of Clustering

HOW MANY CLUSTERS?

Theoretical, conceptual or practical issues may suggest a number.

Hierarchical clustering:

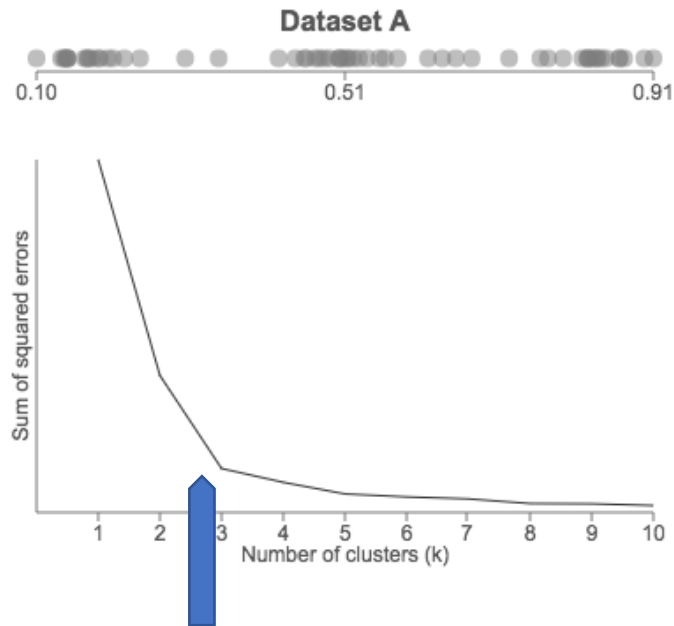
Distance threshold at which clusters are combined

K-means (and other non-hierarchical)

Compare ratio within-groups/between-group variance
against the number of clusters

THE “ELBOW” METHOD

Compare within-groups sum of squares vs # of clusters



Example from [Robert Gove](#)

The “**elbow**” shows the point at which adding more clusters helps reduce distortion measure less and less.

SET RULES FOR SPLITTING / MERGING

ISODATA Algorithm

A **K-means** variant that can add or remove centroids at each step based on user-defined thresholds like:

- Cluster size
- Standard deviation within cluster
- Distance between clusters
- Etc.

SUMMARY: CONDUCTING CLUSTER ANALYSIS

Formulate the Problem

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Interpret and Profile Clusters

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HOW TO TELL IF YOU'VE FOUND GOOD QUALITY CLUSTERS?

Compare cluster stability across:

- Different **distance** measures

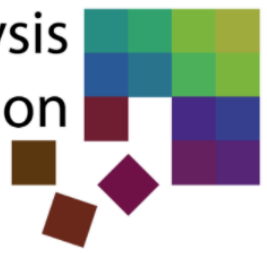
- Different **clustering** methods

- Different 50/50 random data splits (a bit like cross-validation)

- Different **variable/features** deletions

- Different **data orderings** (non-hierarchical)

“Good” clusterings (if they exist) are generally **stable** and **robust** to perturbations in methods or data.



k-means

Three Not Equal Circles

Custom Data

Parameters

k: 3

Distance
Measure:

Euclidian

$$d_2(x, y) = \sqrt{\sum_{i=1}^n |x_i - y_i|^2}$$

Navigation

Iteration #

8

Step #

15

Current Animation Speed

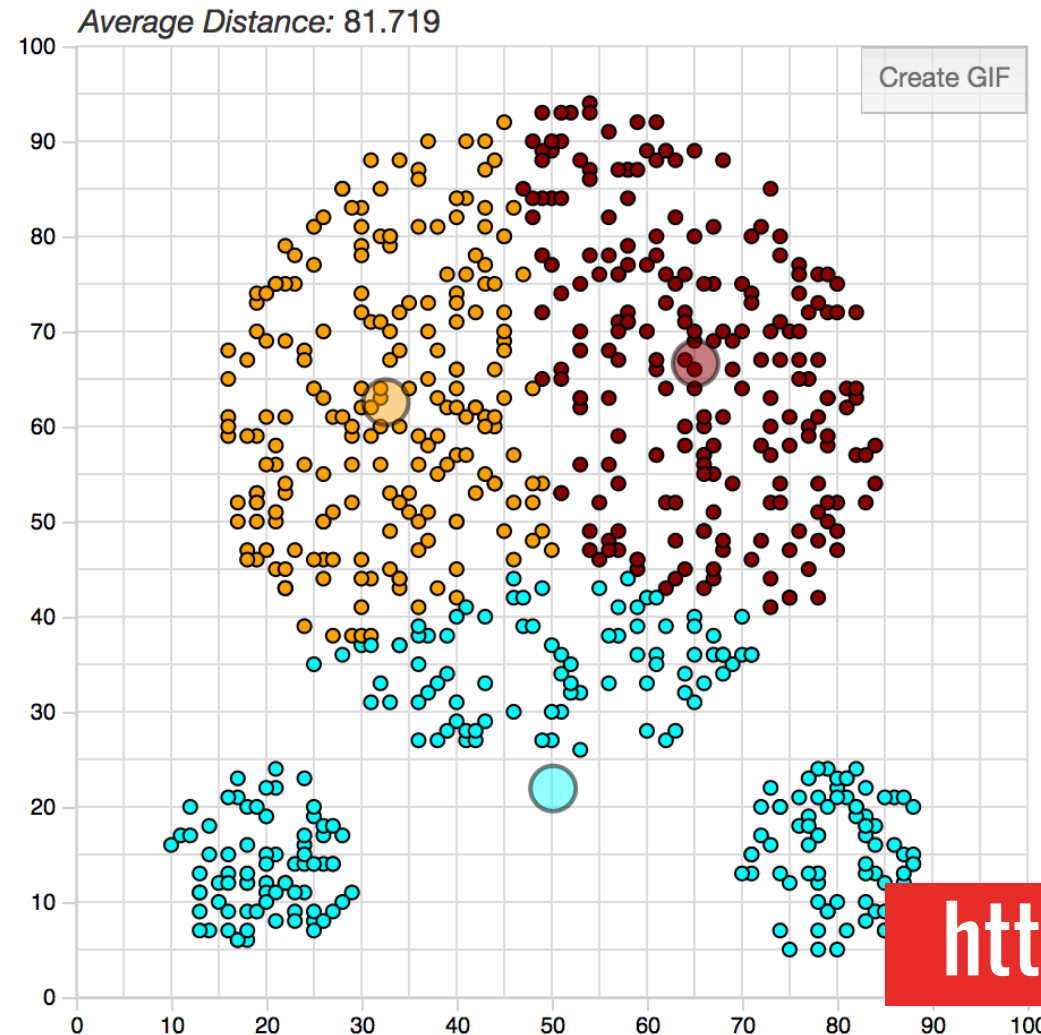
2



Current Step

15

Visualization



Algorithm: k-means

Complexity Range: $O(k \times n \times t)$

Input: k clusters

Output: k clusters

Pseudocode:

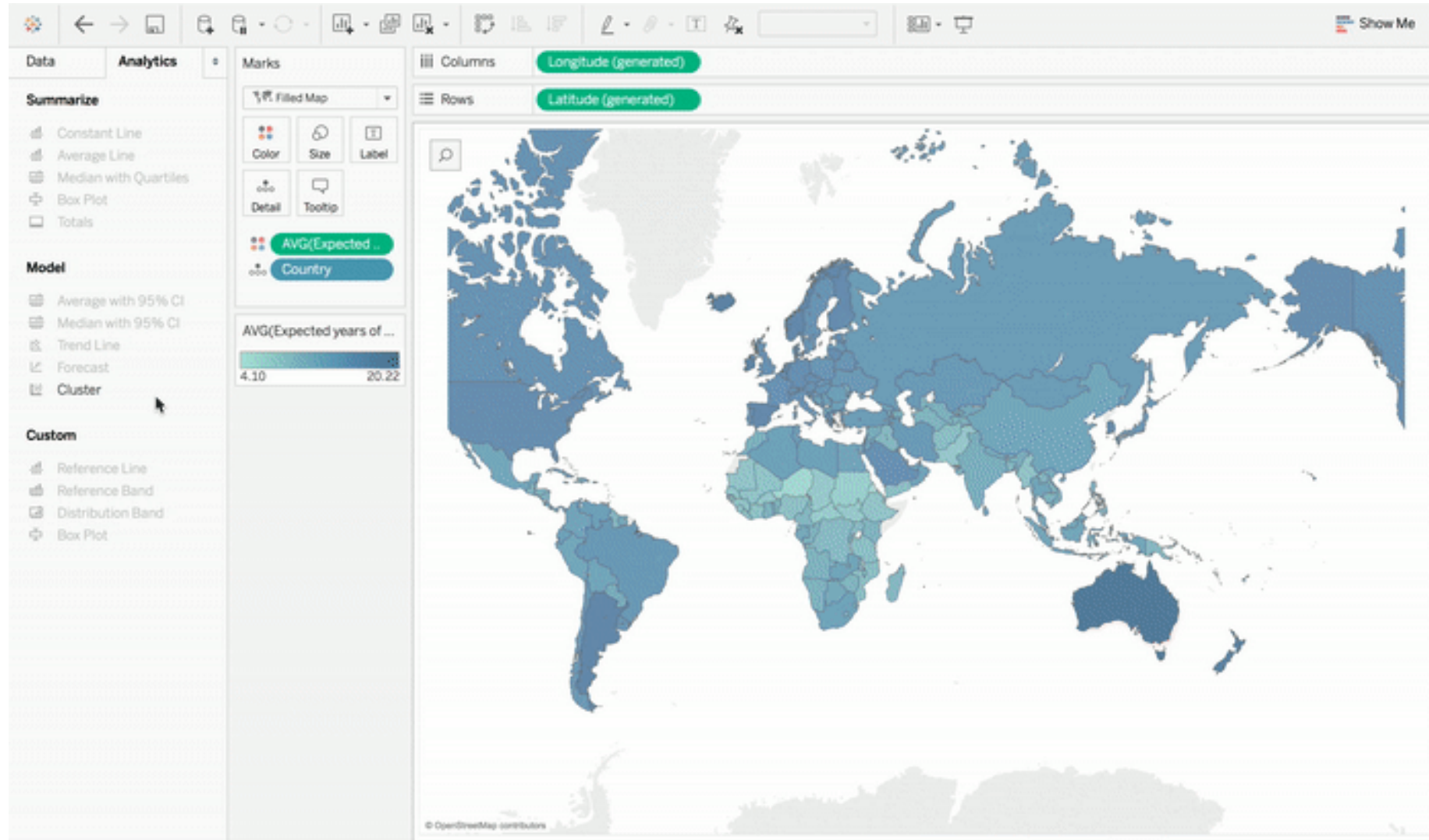
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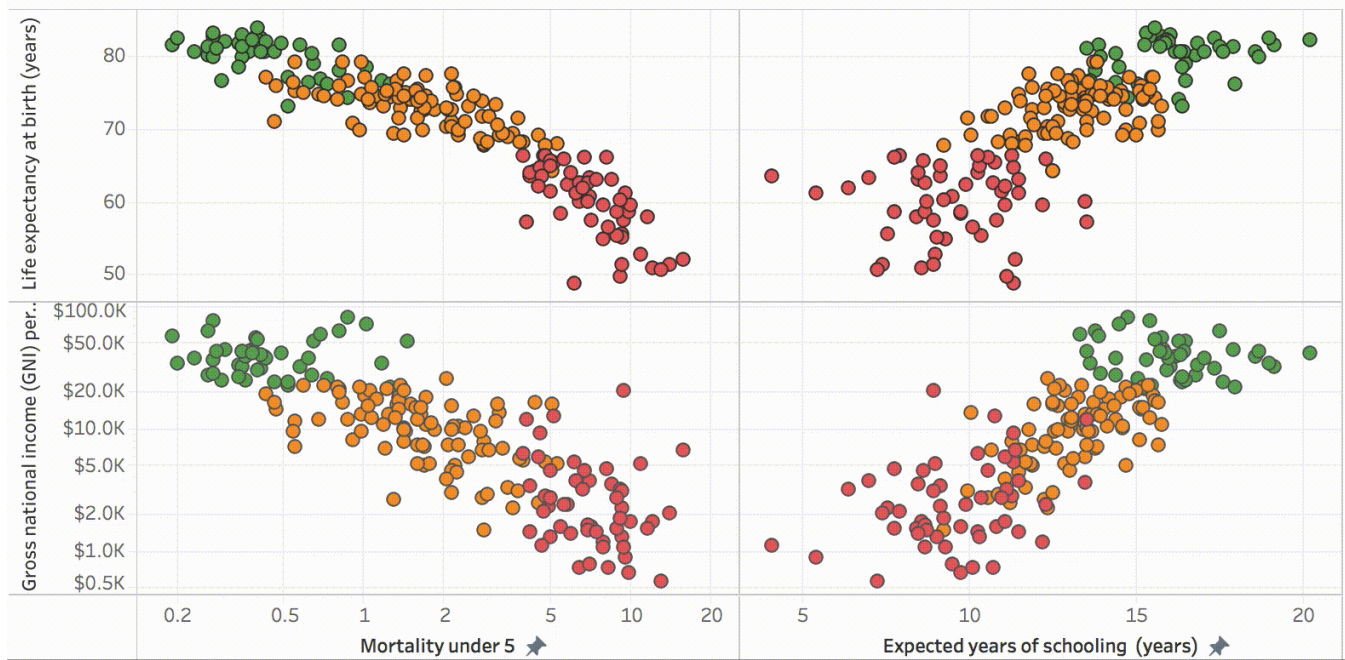
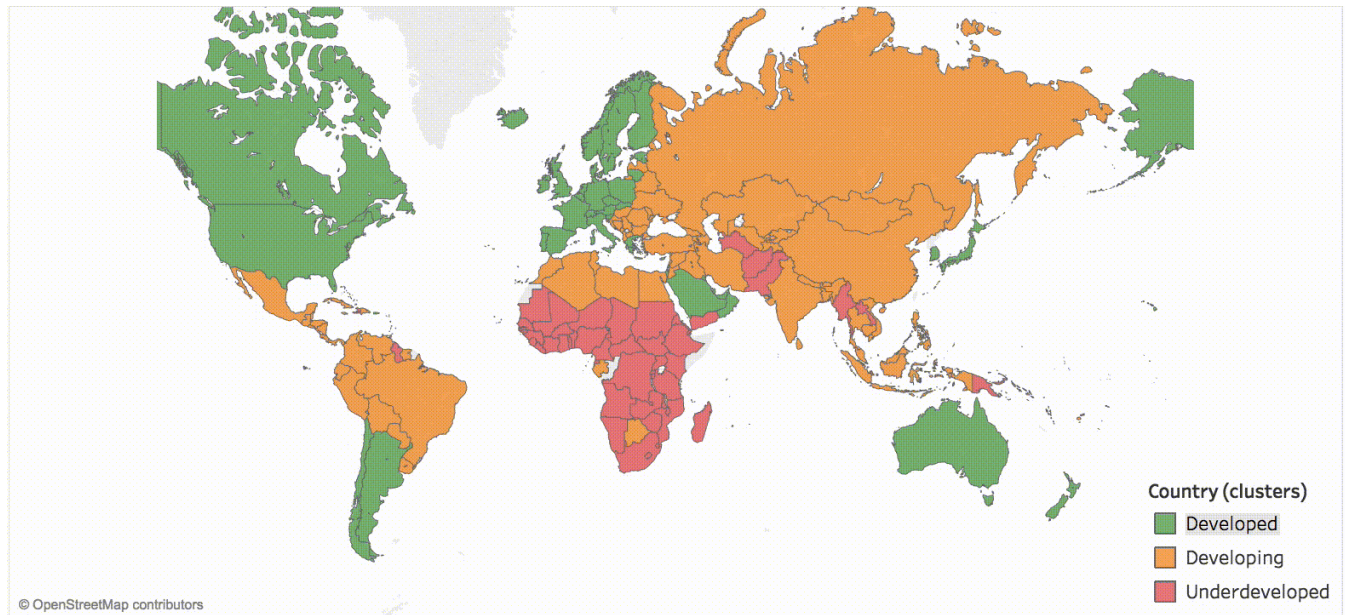
<http://educlust.dbvis.de/>

CLUSTERING IN PRACTICE

Lots of good Python and R libraries.
Demos coming up!

K-MEANS IN TABLEAU





SUMMARY

Clustering is a powerful and broad technique

Many, **many options** for each step

- Similarity metrics

- Type of clustering

 - Hierarchical, k-means, etc.

 - Decisions within each type

- Number of clusters

Many approaches **specifically** for text (coming next week)