NUS | Computing

National University of Singapore

CS5228: Knowledge Discovery and Data Mining

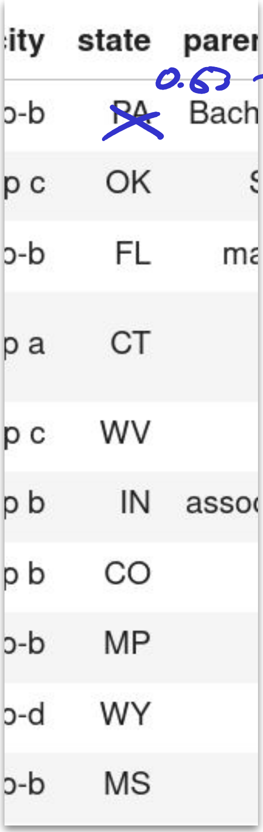
Lecture 3 — Clustering II

Course Logistics — Update

- Assignment 1
  - Topics: EDA, Data Preparation & Clustering
  - Submission deadline: Thu, Sep 18 (11.59 pm)

- Reminder: Honor Code
  - Cheat and plagiarism is serious academic offence
  - Do not read, copy, steal, etc. others code
  - Lecture slides & videos should easily suffice

Quick Recap — Tutorial

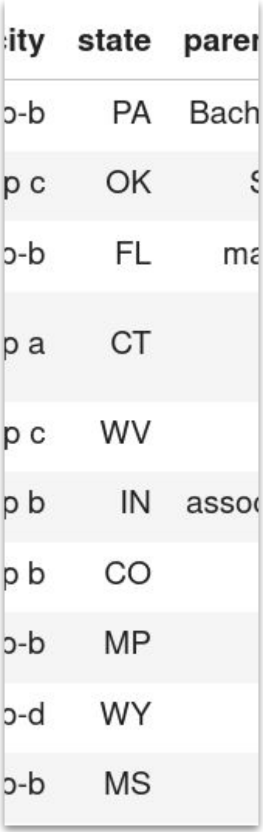


ity	state	parer
o-b	~~PA~~ 0.63	Bach
p c	OK	S
o-b	FL	ma
p a	CT	
p c	WV	
p b	IN	assoc
p b	CO	
o-b	MP	
o-d	WY	
o-b	MS	

- Alternative encoding of nominal attributes: *"proxy encoding"*
  - Replace nominal values with 1 or more numerical values
  - Numerical values should reflect underlying assumption of the impact of attribute
  - Example: *"What makes 'state' a potentially useful attribute?"*

Interpretation		Encoding through replacement
Average Political leaning	➜	Percentage of democrats/republicans
State education budget	➜	Dollar-per-student value
School system	➜	Rate of homeschooling
Urbanization	➜	#universities per capita
…	➜	…

Quick Recap — Tutorial



ity	state	parer
o-b	PA	Bach
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p a	CT	
p c	WV	
p b	IN	assoc
p b	CO	
o-b	MP	
o-d	WY	
o-b	MS	

- Important: "careless" encoding may imply questionable interpretation
  - Question: *"What is the interpretation of my encoding, and is it meaningful?"*
  - In practice, often very difficult to answer

Encoding		Interpretation
Ordinal values	➜	PA < OK < FL < CT < WV < …
Latitude/Longitude	➜	Geographic location of state matters
#KFC per capita	➜	Proliferation of fast food matters
…	➜	…

It **might** be correct, even if only incidentally!

Quick Recap — Lecture

- Clustering
  - Grouping data points based on their similarities
  - No single definition for cluster or clustering ➜ different meaningful intuitions
  - General-purpose data mining method ("only" distance/similarity measure required)

- Algorithms discussed so far:
  - K-Means (centroid-based, partitional, exclusive, complete)
  - DBSCAN (density-based, partitional, exclusive, partial)

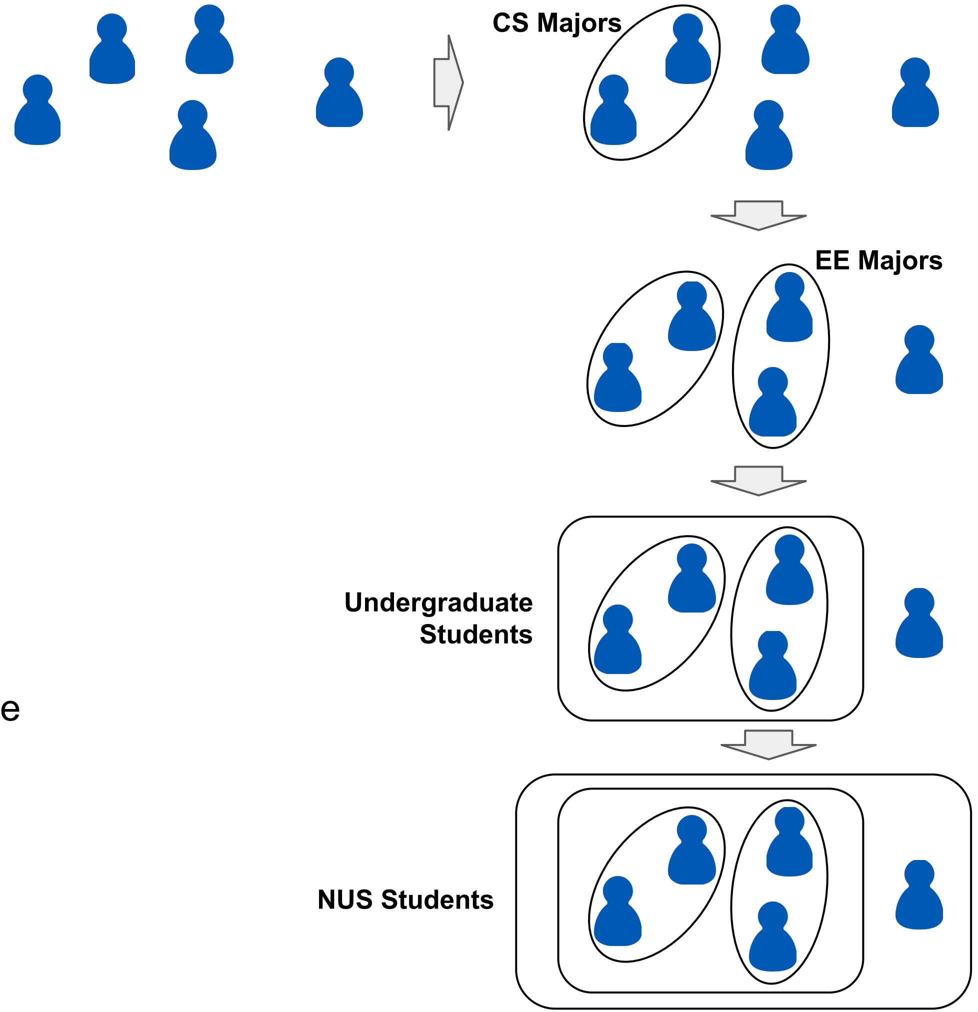
Outline

- Clustering
  - Overview
  - Concepts
  - Applications
- **Clustering algorithms**
  - K-Means
  - DBSCAN
  - **Hierarchical Clustering**
- Cluster Evaluation

Hierarchical Clustering

- Basic characteristics
  - Clusters: depends...
  - Clustering: hierarchical (duh!), complete, exclusive (at each level!)

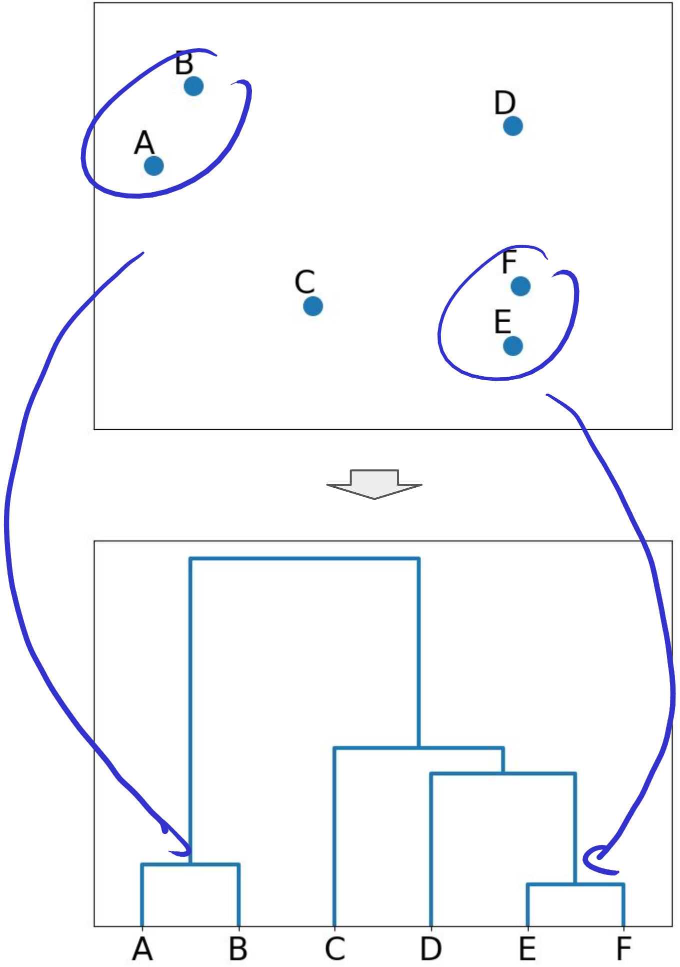
- No parameterization (in principle)
  - In practice, typically number of clusters is specified (similar to K-means)
  - Different choices of measures to calculate distances between clusters



Dendrograms

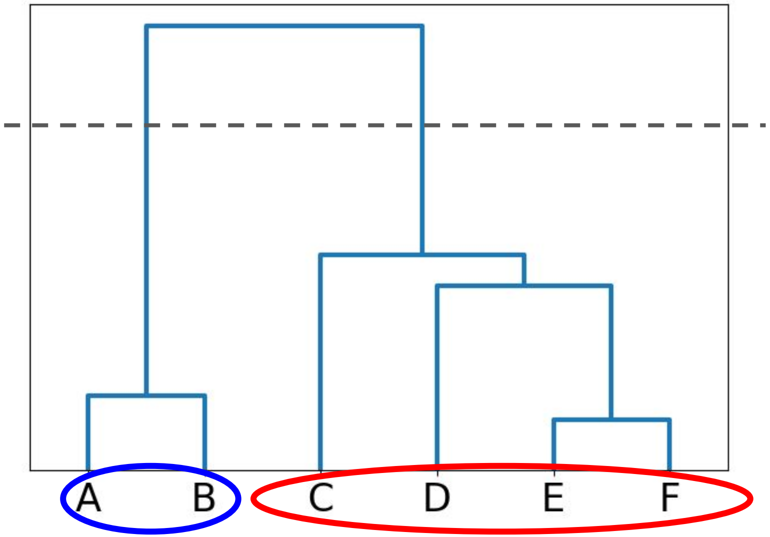
greek : tree

- Dendrogram: Visualization of hierarchical relationships
  - Binary tree showing how clusters are hierarchically merged/split
  - Each node is a cluster
  - Each leaf is a singleton cluster
  - Height reflects distance between clusters
    (e.g., large distance between A/B and C/D/E/F clusters)

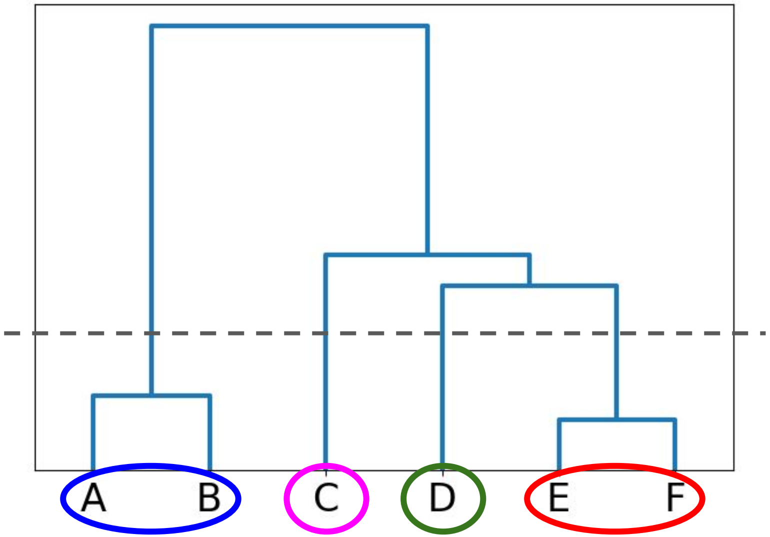


Hierarchical Clustering — Dendrograms

- A clustering can be obtained by cutting a dendrogram at the desired level



2 clusters



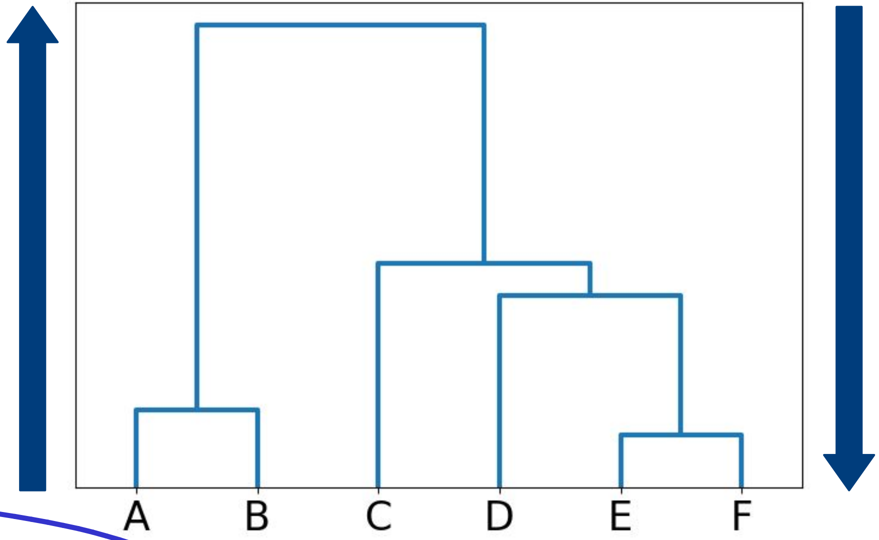
4 clusters

Hierarchical Clustering — 2 Main Types

Agglomerative

(bottom-up)

- Start with each point being its own cluster
- At each step, merge closest pair of clusters
- Stop when only one cluster is left



Divisive

(top-down)

- Start with one cluster containing all points
- At each step, split a cluster
- Stop when each cluster contains a single point

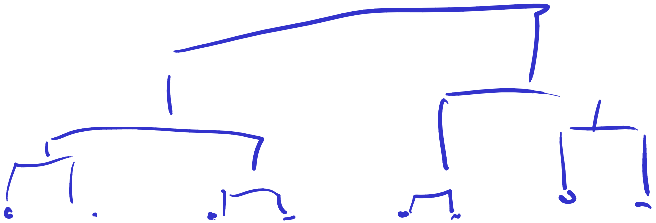
AGNES (AGglomerative NESting)

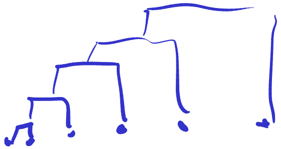
DIANA (DIvise ANAlysis)

Quick Quiz

What is the **minimum** and **maximum** possible **depth** of a dendrogram for a dataset with *N* data points?

(assume O-notation)





[✓ **A**] Min: $\log_2 N$ Max: N

B Min: $\sqrt{N}$ Max: $N \log_2 N$

C Min: $\sqrt{N}$ Max: N

D Min: $\log_2 N$ Max: $N \log_2 N$

AGNES — Basic Algorithm

1. Initialization: Each point forms its own cluster
2. Repeat

    2a) **Merge** the two closest clusters into one

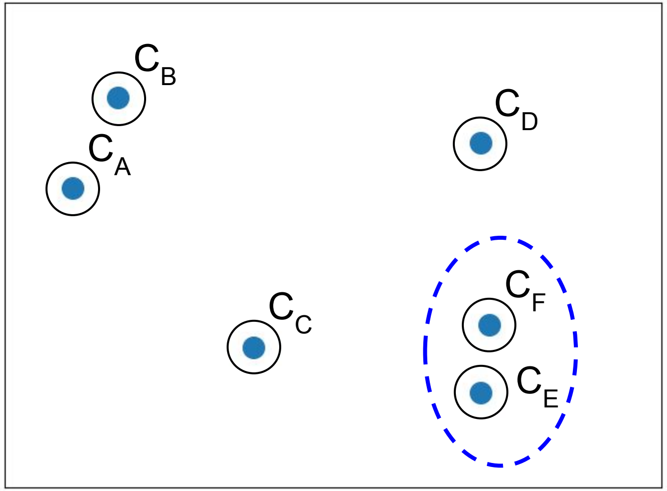
    Until only 1 cluster remains

Download at: https://www.comp.nus.edu.sg/~chris/downloads/images/agglomerative-clustering-visualized.gif

AGNES — Implementation

- Implementation using distance matrix

Initial clustering: each cluster, one point



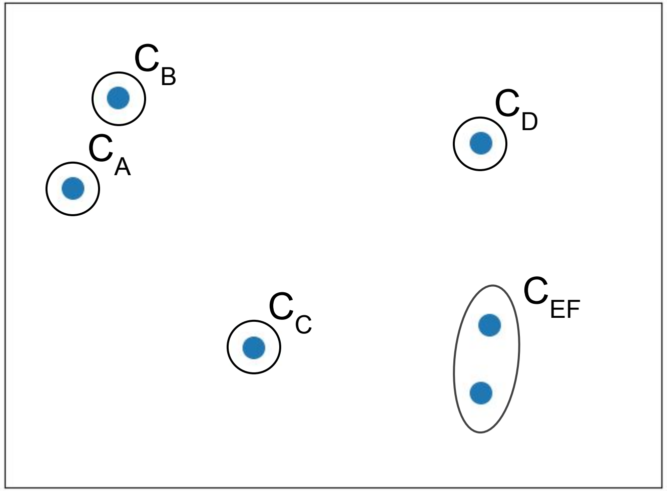
Distance between clusters = distance between points

	C_A	C_B	C_C	C_D	C_E	C_F
C_A	∞	2.25	5.32	9.06	9.79	9.49
C_B		∞	6.08	7.85	9.86	9.21
C_C			∞	6.73	4.81	5.02
C_D				∞	5.51	4.00
C_E					∞	**1.53**
C_F						∞

AGNES — Implementation

- What's the distance between clusters? (beyond containing single points)

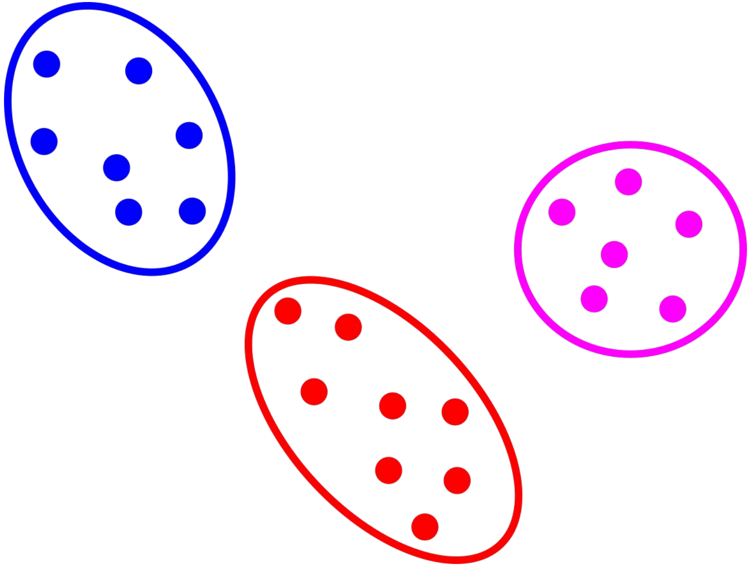
Clustering after merging C_E and C_F to C_{EF}



	C_A	C_B	C_C	C_D	C_{EF}
C_A	∞	2.25	5.32	9.06	???
C_B		∞	6.08	7.85	???
C_C			∞	6.73	???
C_D				∞	???
C_{EF}					∞

Quick "Quiz"

Which 2 clusters should get merged next?



[✓ A] Red & Blue

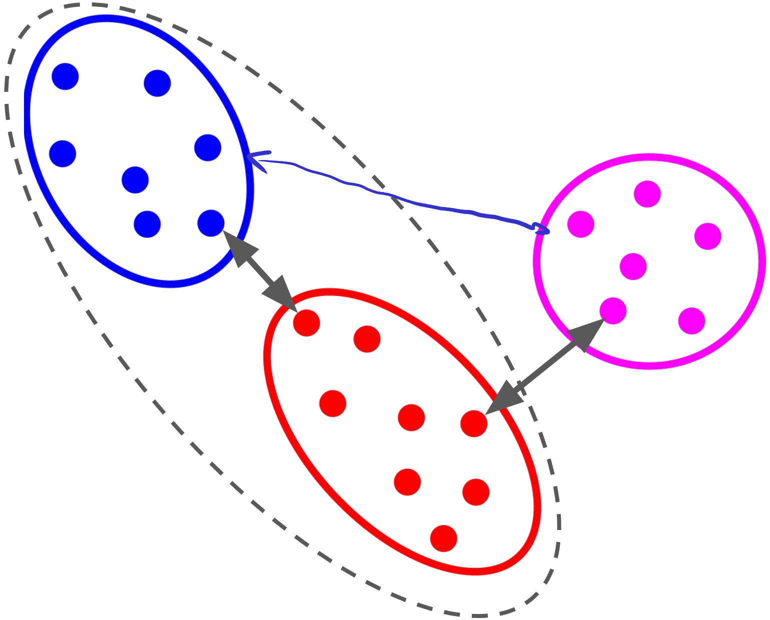
[✓ B] Red & Magenta

C Blue & Magenta

D It's Friday, I'm out...

AGNES — Single Linkage

- Single Linkage Clustering
  - Distance between clusters = **minimum distance** between two points from each cluster



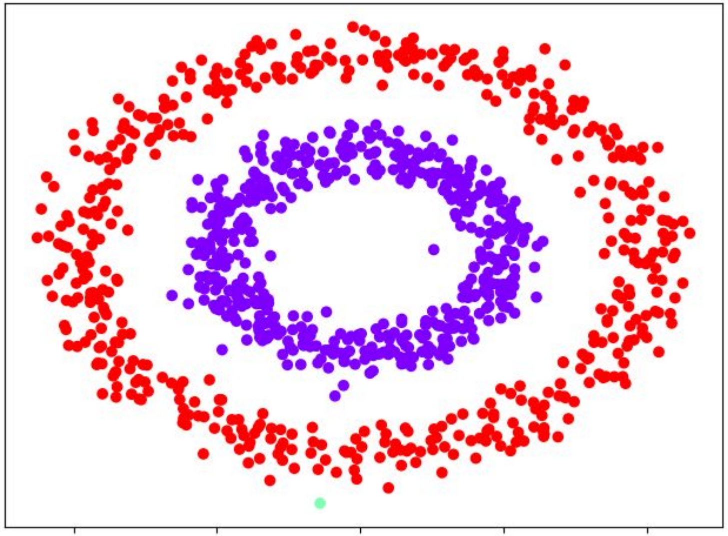
simple pointwise distance

$$d_{single}(C_i, C_j) = \min_{p \in C_i, q \in C_j} d(p, q)$$

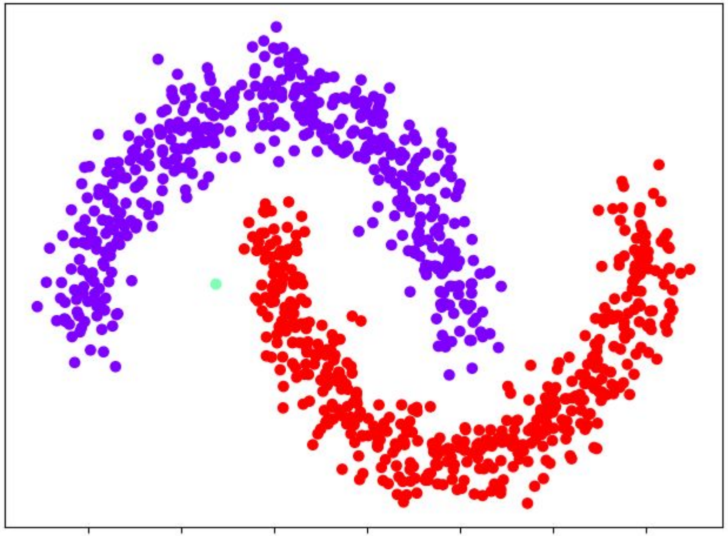
AGNES — Single Linkage

non-blob-like

- Strength: Can handle non-globular shapes



#cluster = 3

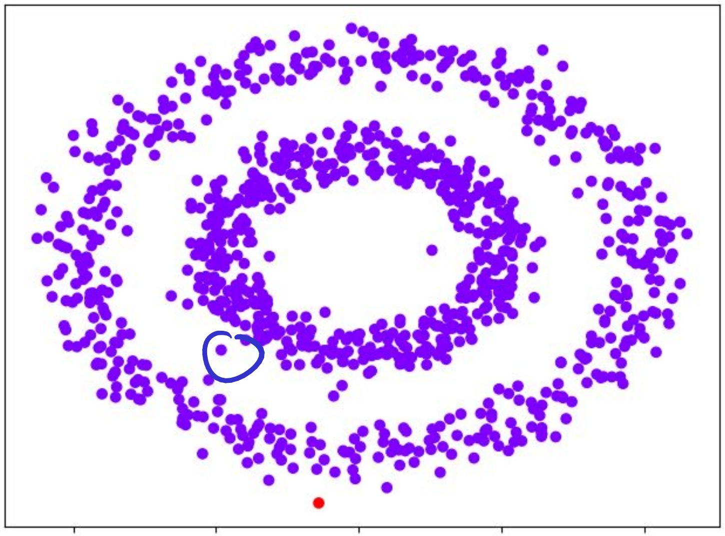


#cluster = 3

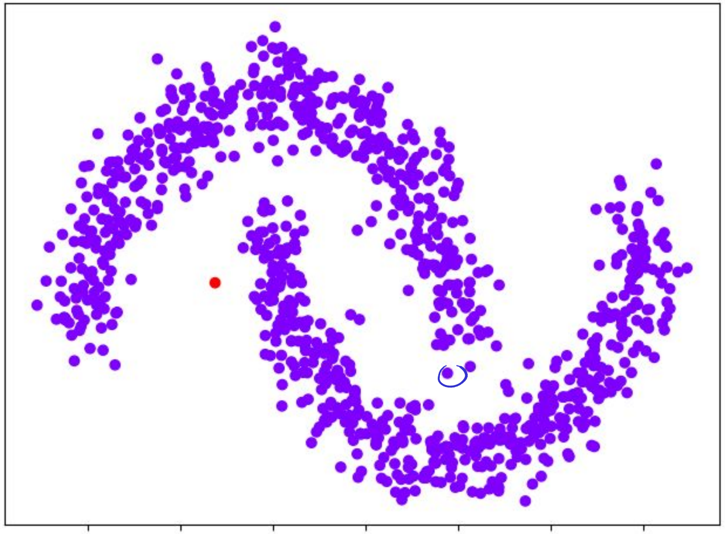
AGNES — Single Linkage

- Weakness: Very susceptible to noise ➜ "Chaining"
  - A single point may cause two clusters get merged

#cluster = 2

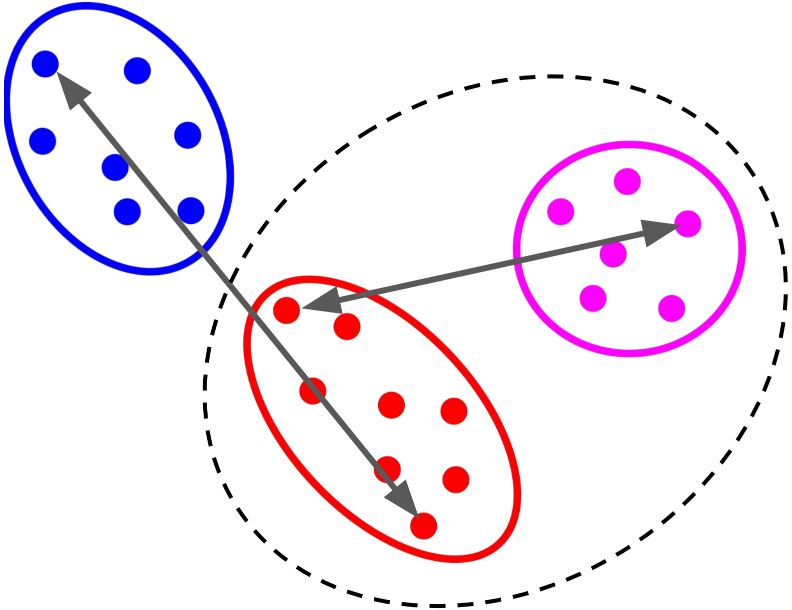


#cluster = 2



AGNES — Complete Linkage

- Complete Linkage Clustering
  - Distance between clusters = **maximum distance** between two points from each cluster

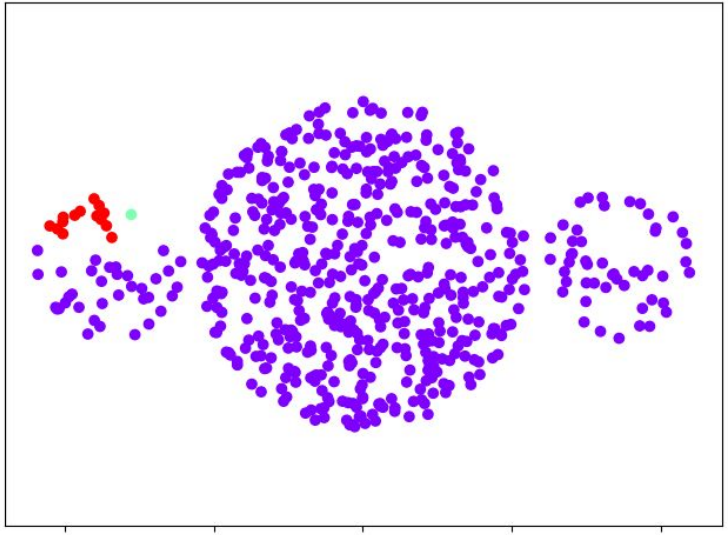


$$d_{complete}(C_i, C_j) = \underset{p\in C_i, q\in C_j}{\underline{max}} \; d(p, q)$$

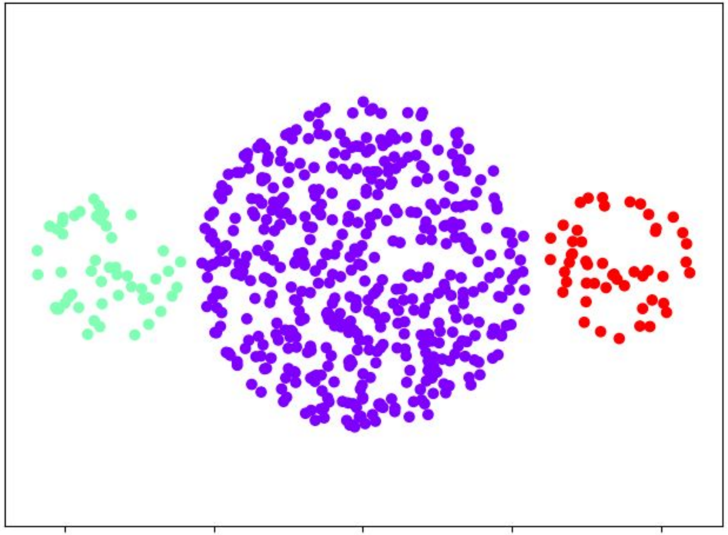
AGNES — Complete Linkage

- Strength: Less susceptible to noise or outliers

Single Linkage, #cluster = 3



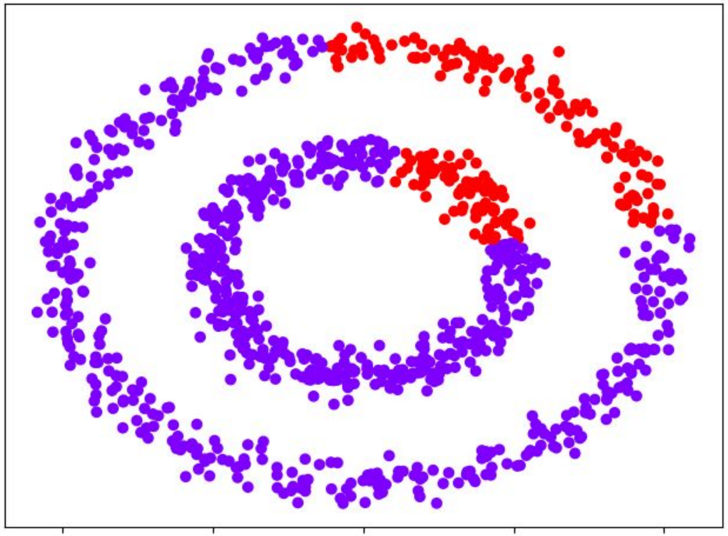
Complete Linkage, #cluster = 3



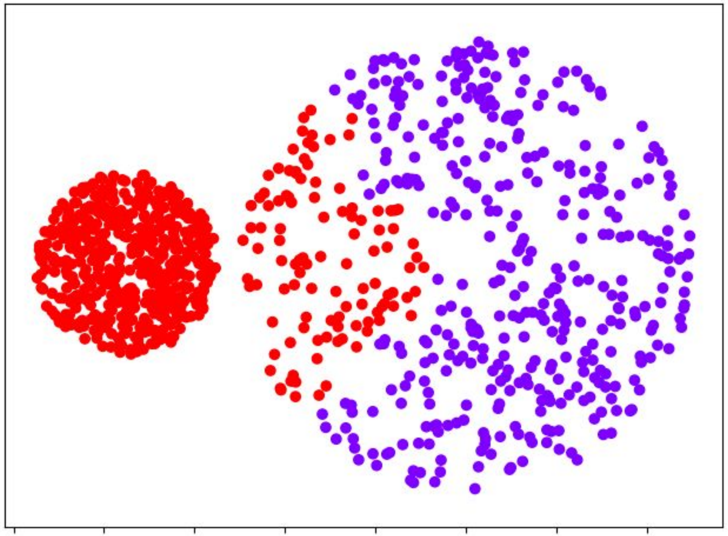
AGNES — Complete Linkage

- Weaknesses
  - Bias towards globular clusters
  - Tends to break large clusters

#cluster = 2

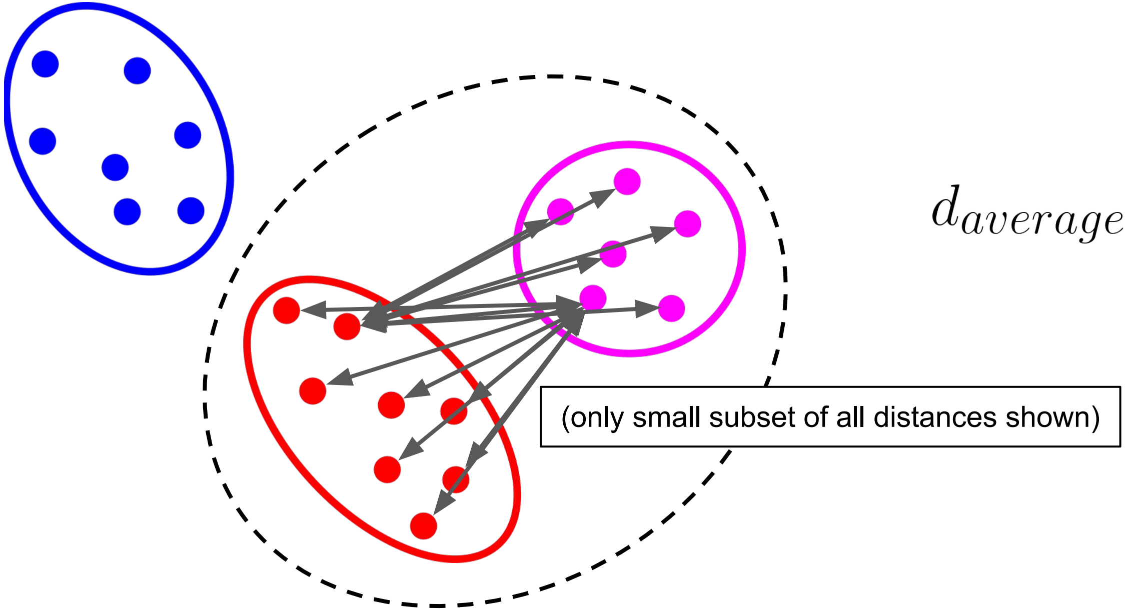


#cluster = 2



AGNES — Average Linkage

- Complete Linkage Clustering (compromise between single and complete linkage)
  - Distance between clusters = **average distance** between two points from each cluster



(only small subset of all distances shown)

$$d_{average}(C_i, C_j) = \underset{p \in C_i, q \in C_j}{\underline{avg}} \; d(p, q)$$

AGNES — Linkage Alternatives

- Centroid linkage
  - Distance between clusters = distance between the centroids of each cluster

$$d_{centroid}(C_i, C_j) = d(\underbrace{m_i, m_j}_{\text{centroid of cluster } i \text{ and } j \text{ (m for mean)}})$$

sklearn
✓

- Ward linkage

$$d_{Ward}(C_i, C_j) = \overbrace{\sum_{k \in C_i \cup C_j} ||x_k - m_{ij}||^2}^{\text{Variance of } C_{ij}} - \overbrace{\sum_{k \in C_i} ||x_k - m_i||^2}^{\text{Variance of } C_i} - \overbrace{\sum_{k \in C_j} ||x_k - m_j||^2}^{\text{Variance of } C_j}$$

$$= \frac{n_i n_j}{n_i + n_j} ||m_i - m_j||^2$$

n_i = #points in cluster C_i

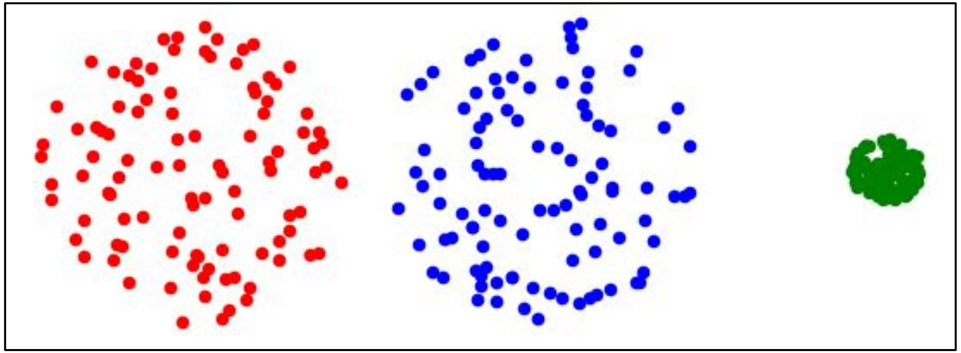
Ward Linkage — Intuition

$$d_{Ward}(C_i, C_j) = \overbrace{\sum_{k \in C_i \cup C_j} ||x_k - m_{ij}||^2}^{\text{Variance of } C_{ij}} - \overbrace{\sum_{k \in C_i} ||x_k - m_i||^2}^{\text{Variance of } C_i} - \overbrace{\sum_{k \in C_j} ||x_k - m_j||^2}^{\text{Variance of } C_j}$$

- Example for Ward Linkage
  - Each blob: 100 data points

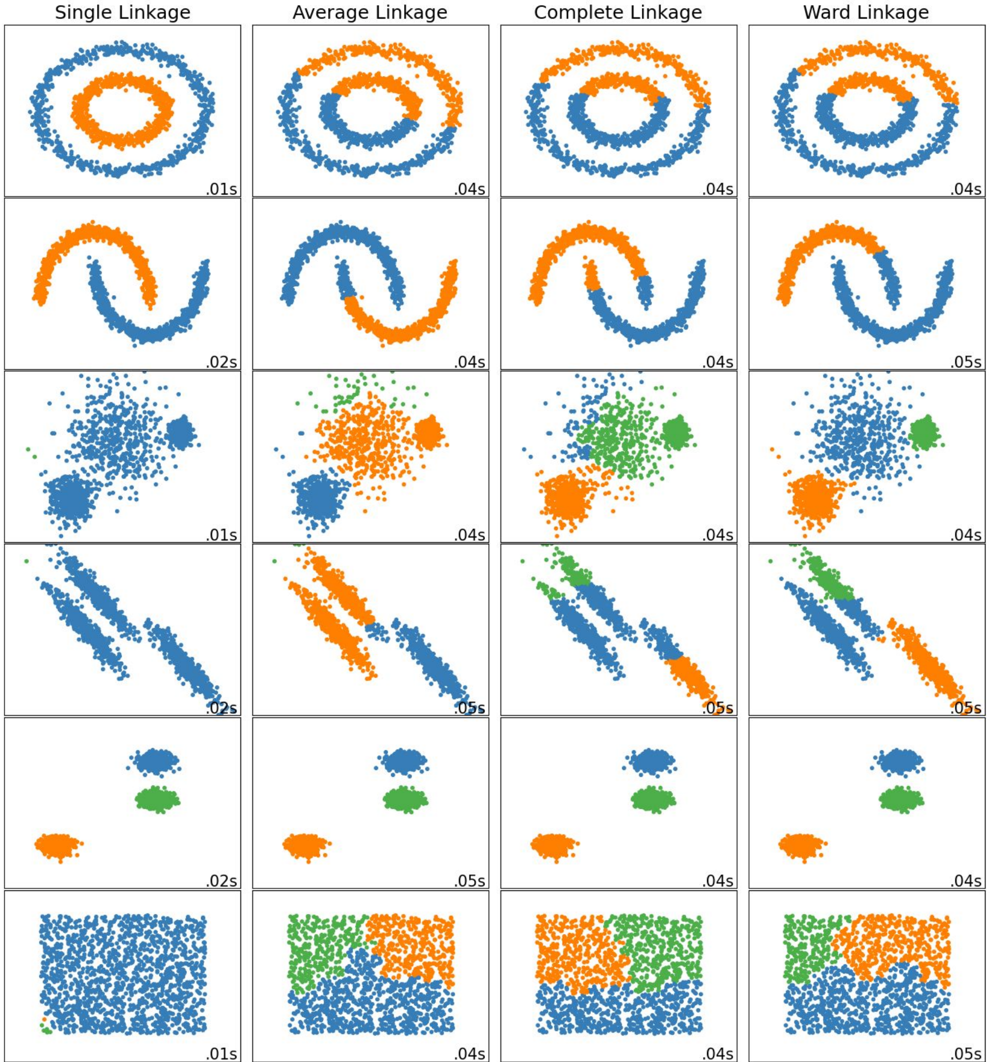
d_{Ward}(Red, Blue) = 1,635 - 195 - 200 = **1,240**

d_{Ward}(Blue, Green) = 1,450 - 200 - 10 = **1,240**



AGNES

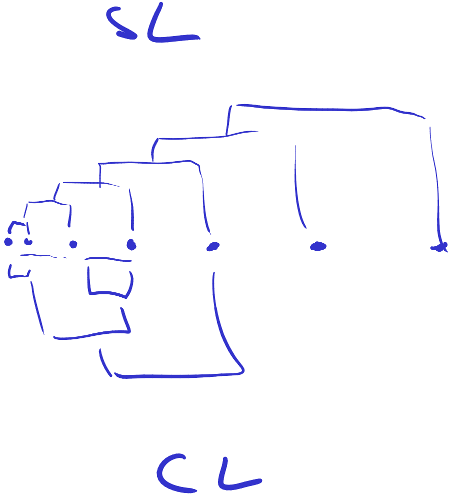
- Linkage comparison



Source:
https://scikit-learn.org/stable/auto_examples/cluster/plot_linkage_comparison.html

Quick Quiz

Which linkage method has intuitively the **highest chance** of returning a clustering where the dendrogram has a **depth of *N-1***?



[✓ **A** Single]

B Complete

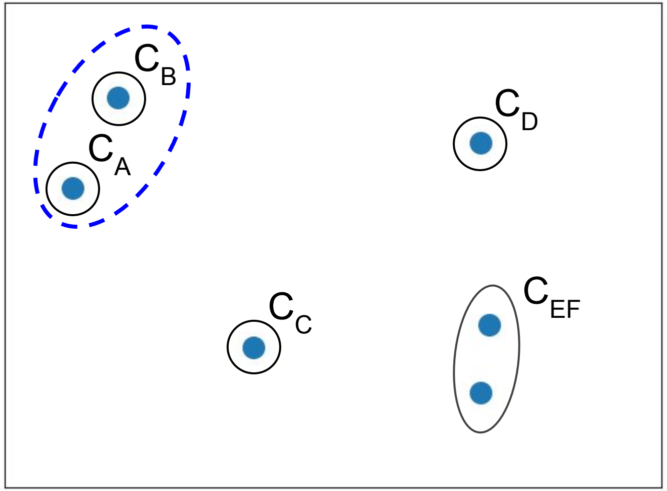
C Average

D No difference

AGNES — Implementation

- Distance matrix after merging Cluster C_E and C_F + Average Linkage

Clustering after merging C_E and C_F to C_{EF}

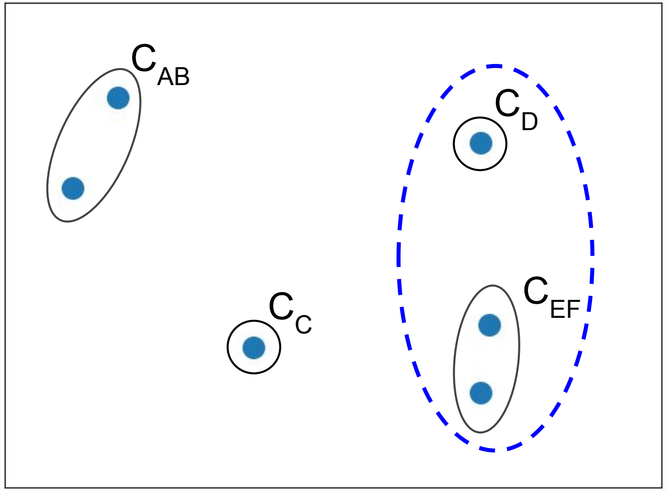


	C_A	C_B	C_C	C_D	C_{EF}
C_A	∞	**2.25**	5.32	9.06	9.64
C_B		∞	6.08	7.85	9.54
C_C			∞	6.73	4.92
C_D				∞	4.76
C_{EF}					∞

AGNES — Implementation

- Distance matrix after merging Cluster C_A and C_B + Average Linkage

Clustering after merging C_A and C_B to C_{AB}

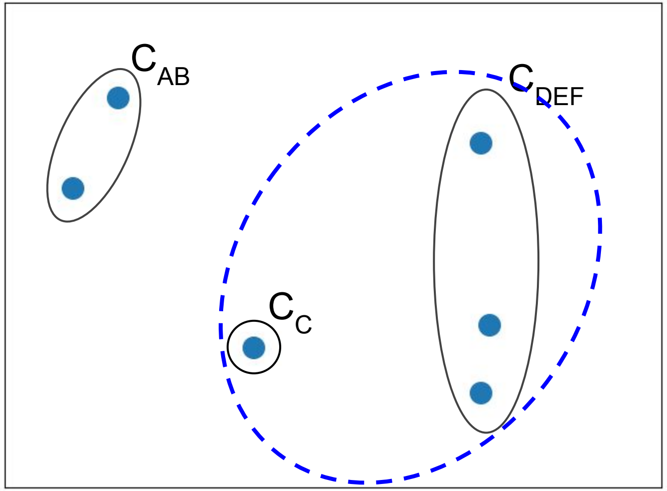


	C_{AB}	C_C	C_D	C_{EF}
C_{AB}	∞	5.70	8.45	9.59
C_C		∞	6.73	4.92
C_D			∞	**4.76**
C_{EF}				∞

AGNES — Implementation

- Distance matrix after merging Cluster C_C and C_{DEF} + Average Linkage

Clustering after merging C_D and C_{EF} to C_{DEF}

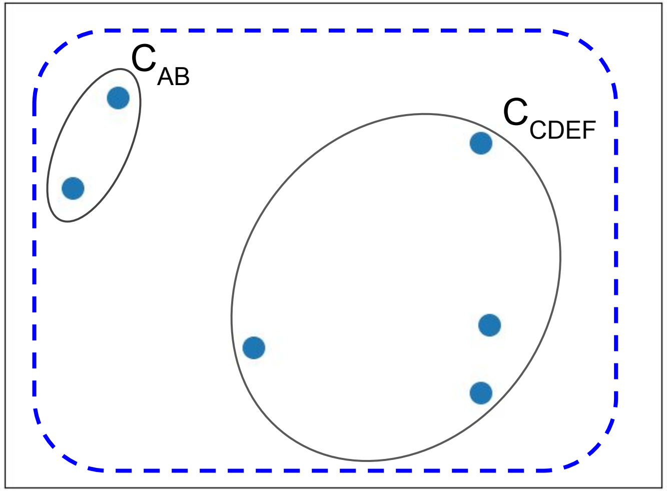


	C_{AB}	C_C	C_{DEF}
C_{AB}	∞	5.70	9.21
C_C		∞	**5.51**
C_{DEF}			∞

AGNES — Implementation

- Distance matrix after merging Cluster C_{AB} and C_{CDEF} + Average Linkage

Clustering after merging C_C and C_{DEF} to C_{CDEF}

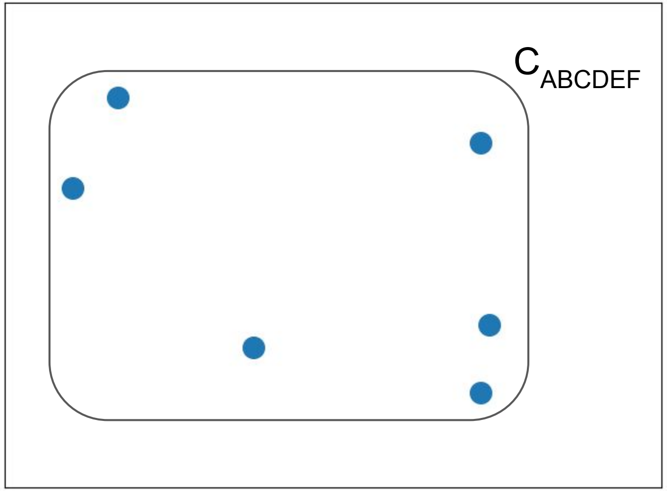


	C_{AB}	C_{CDEF}
C_{CDEF}	∞	**8.33**

AGNES — Implementation

- Distance matrix after merging Cluster C_{AB} and C_{CDEF} + Average Linkage

Clustering after merging C_{AB} and C_{CDEF} to C_{ABCDEF}



	C_{ABCDEF}
C_{ABCDEF}	∞

➜ Done!

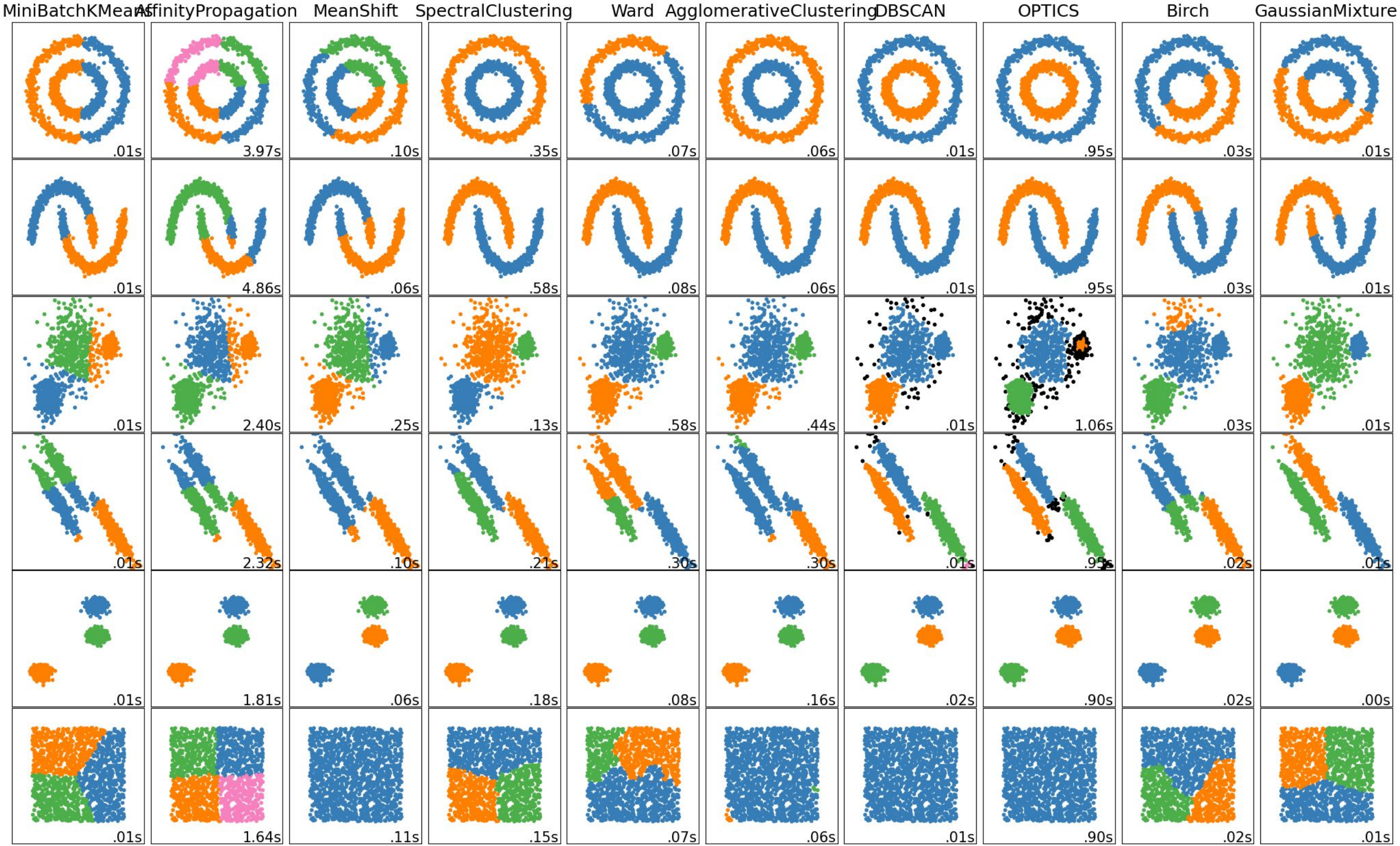
AGNES — Complexity Analysis

- Space Complexity: **$O(N^2)$**
  - Storing distance matrix

- Time Complexity
  - Baseline: **$O(N^3)$** — (N-1) steps, each step $O(N^2)$ to scan distance matrix
  - Using more sophisticated data structures, e.g, heap or priority queue: **$O(N^2 \log N)$**
  - Special optimization for Single Linkage Clustering: **$O(N^2)$**

DIANA — DIvisive ANAlysis

- Top-Down Hierarchical Clustering
  - Start with all points forming one cluster
  - Recursively split one cluster until all cluster have size 1
- Challenge: 2^n ways to split a cluster with *n* points
  - Heuristics needed to restrict search space
  - Generally slower and less common than AGNES
- Cases where DIANA can perform better
  - No complete clustering needed ➜ early stopping
  - Splitting can utilize global knowledge (merging based on local knowledge only)



Source: https://scikit-learn.org/stable/modules/clustering.html

Outline

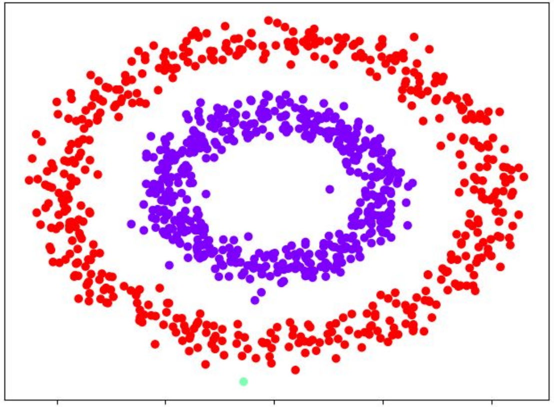
- Clustering
  - Overview
  - Concepts
  - Applications

- Clustering algorithms
  - K-Means
  - DBSCAN
  - Hierarchical Clustering

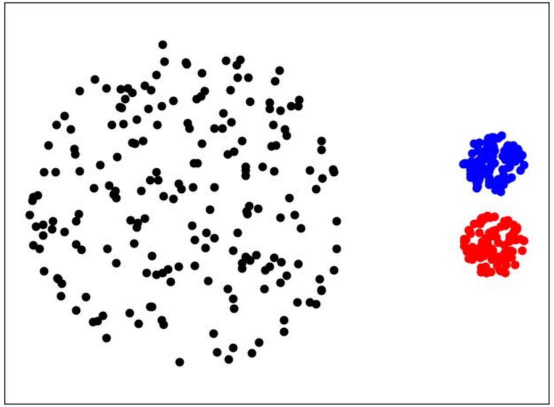
- **Cluster Evaluation**

Cluster Evaluation

- Problem 1: Just eyeballing the clustering is rarely possible
  - High-dimensional data (≥ 3 dimensions) difficult to impossible to visualize
  - Difficult to assess "nature" of clusters a-priori (e.g., variations in shape, size, density, etc)
  - Presence and distribution of noise or outliers



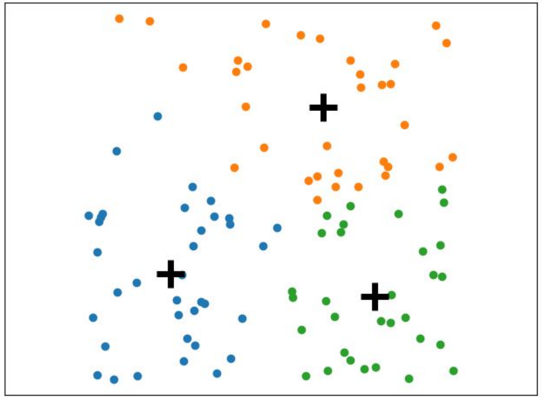
Your data usually does not look like this



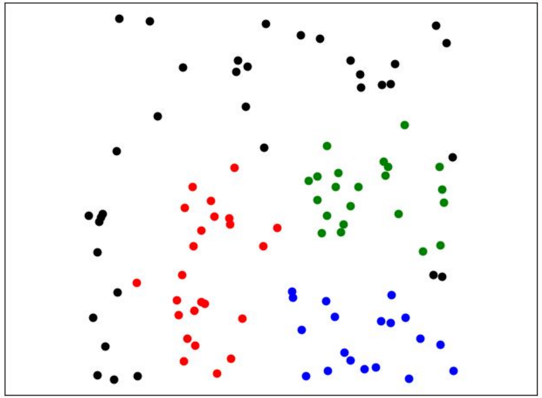
Cluster Evaluation

- Problem 2: Clustering algorithms will always find some clusters
  - Example: K-Means, DBSCAN and AGNES applied to random data

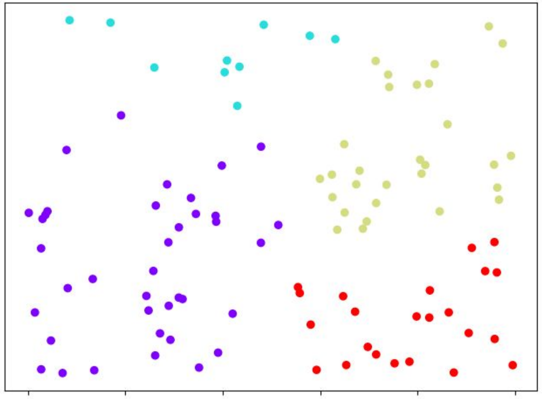
K-Means



DBSCAN



AGNES



Cluster Evaluation

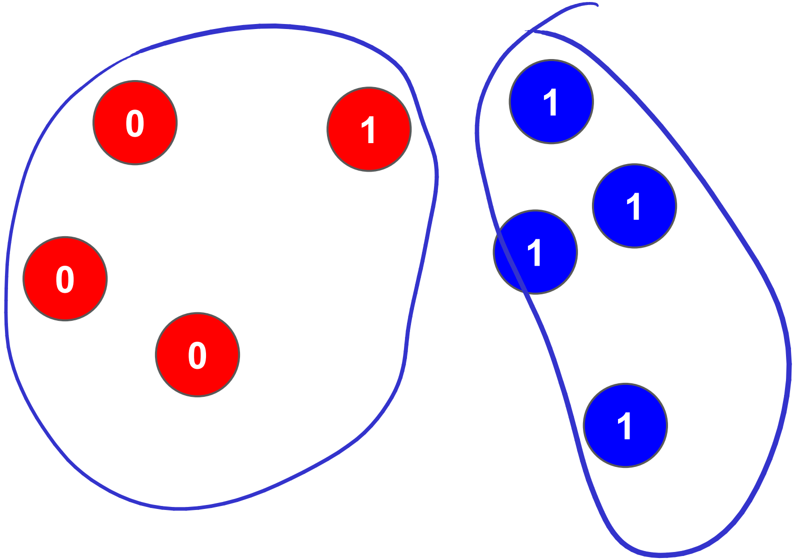
- Purpose of cluster evaluation
    - Comparing the results of different clustering algorithms
    - Comparing the results of a clustering algorithm with different parameters
    - Minimizing the effects of noise on the clustering

➜ Getting a sense of the "goodness" of a clustering

- Two main approaches
    - External quality measures: evaluate a clustering against a ground truth (if available)
    - **Internal quality measures**: evaluate clustering from the data itself

Cluster Evaluation — External Quality Measures

- Ground truth: Labeled data
  - Labels indicate that two points "belong together"
  - If cluster reflect this ➜ good clustering



truth

Cluster	Label
Red	0
Red	0
Red	0
Red	1
Blue	1
Blue	1
Blue	1
Blue	1

External Quality Measures — Cluster Purity

- Cluster purity P
  - N: #points, C: set of cluster, L: set of labels

$$P = \frac{1}{N}\sum_{c\in C}\overbrace{\underset{l\in L}{max}|c\cap l|}^{\text{\#points with most common label } l \text{ in cluster } c}$$

Purity for example:

$$P = \frac{1}{8}(3+4) = 0.875$$

Cluster	Label
Red	0
Red	0
Red	0
Red	1
Blue	1
Blue	1
Blue	1
Blue	1

- Limitations
  - Purity does not penalize having many cluster

    ➜ P=1 easy to achieve with all cluster containing single point

$$P = \frac{1}{8}(\underbrace{1+1+1 \ldots +1}_{8x})$$

External Quality Measures: Information Retrieval Metrics

- Established metrics from classification tasks
  - ■ **TP** — true positives
    same cluster, same label
    (A/B, A/C, B/C, E/F, …, G/H)
  - ■ **TN** — true negatives
    different clusters, different labels
    (A/E, A/F, A/G, A/H, B/E, …, C/H)
  - ■ **FP** — false positives
    same cluster, different labels
    (A/D, B/D, C/D)
  - ■ **FN** — false negatives
    different cluster, same label
    (D/E, D/F, D/G, D/H)

For the example:
- **TP** = 9
- **TN** = 12
- **FP** = 3
- **FN** = 4

ID	Custer	Label
A	Red	0
B	Red	0
C	Red	0
D	Red	1
E	Blue	1
F	Blue	1
G	Blue	1
H	Blue	1

External Quality Measures — Information Retrieval Metrics

- Rand Index RI
  - Reflects accuracy

$$RI = \frac{TP + TN}{TP + FP + FN + TN}$$

$$\boxed{RI_{example} = 0.75}$$

- Precision P, Recall R, F1-Score

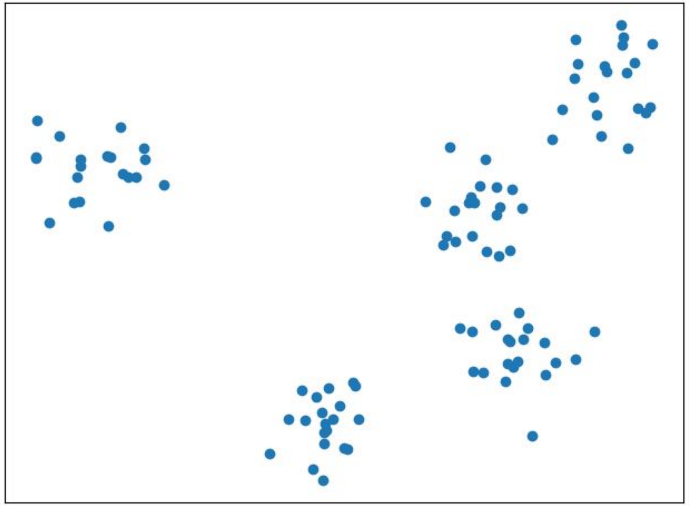
$$P = \frac{TP}{TP + FP} \qquad R = \frac{TP}{TP + FN} \qquad F1 = \frac{2 \cdot P \cdot R}{P + R}$$

$$\boxed{P_{example} = 0.75 \qquad R_{example} = 0.69 \qquad F1_{example} = 0.72}$$

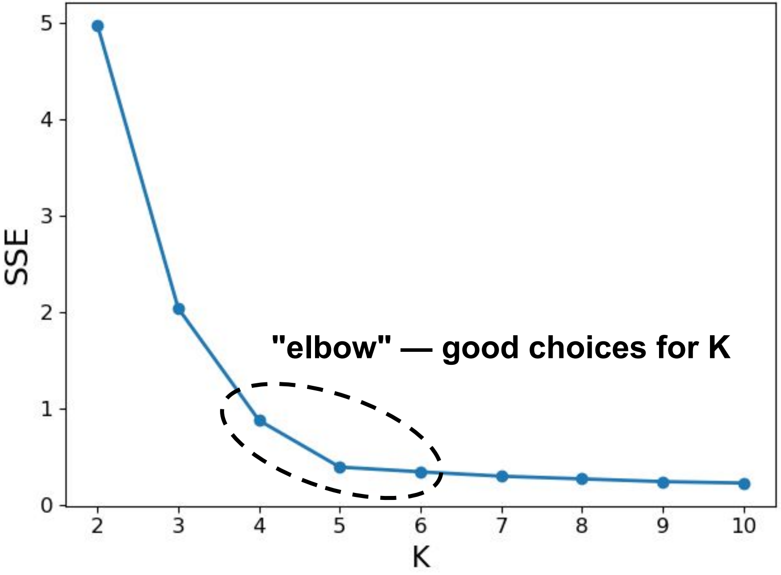
- ...and other metrics based on TP, TN, FP, FN

Internal Quality Measures — SSE

- Use SSE to select number of clusters



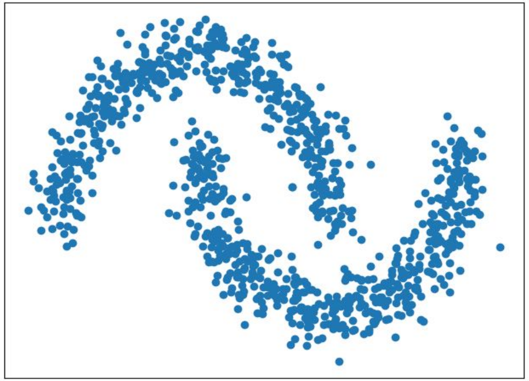
input data



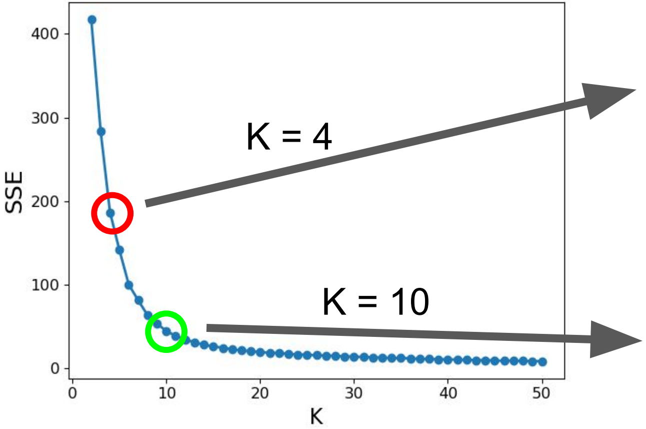
SSE for different K

Internal Quality Measures — SSE

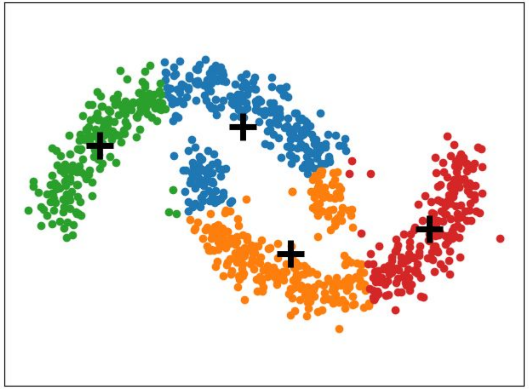
- Also applicable to more complicated data
  - But inherently "favors" globular clusters

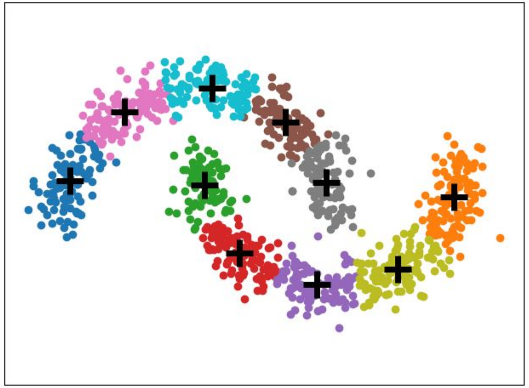


input data



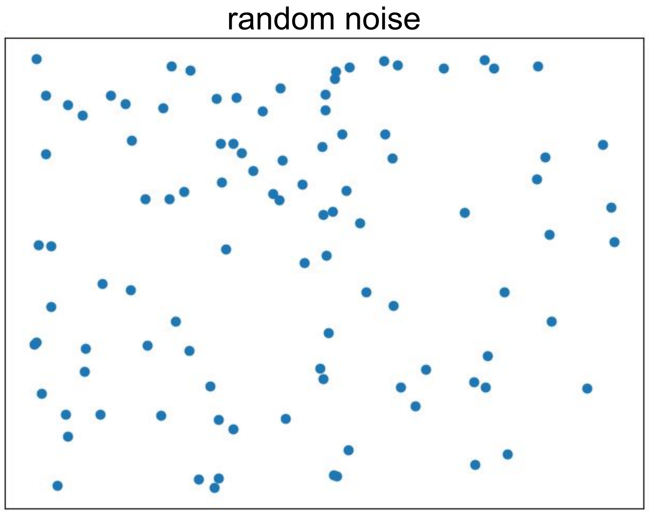
SEE for different K

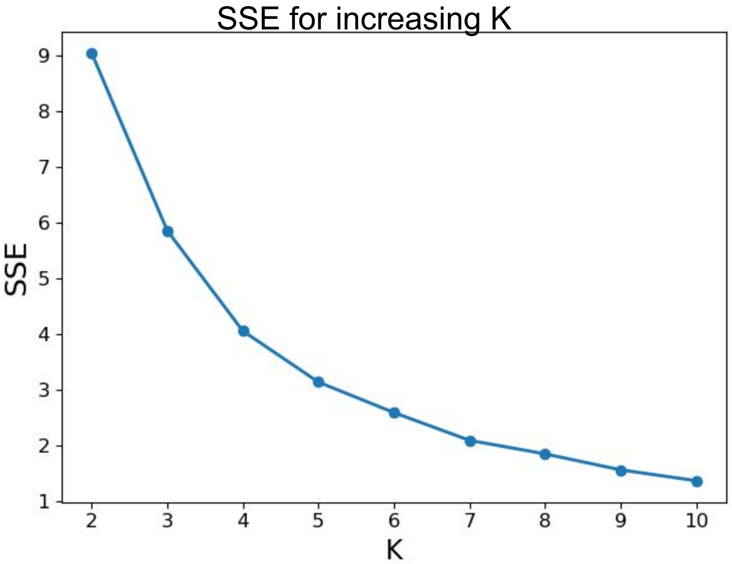




Internal Quality Measures — SSE

- Limitation of SSE as quality measure
  - SSE does not penalize large number of clusters
  - SSE decreases for increasing cluster counts
  - Applicable beyond K-Means, but less intuitive interpretation (in case of non-globular clusters)





Quick Quiz

If K << N, can SSE=0?

Why or **why not**?

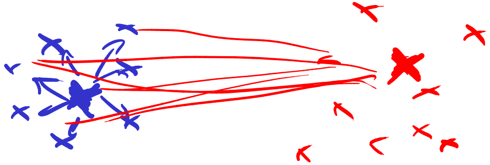
⇒ duplicates

[✓ **A** Yes]

B No

Internal Quality Measures — Silhouette Coefficient

- Intuition: A good clustering has
  - High inter-cluster distances
  - Low intra-cluster distances



- For each data point x, define
  - **Cohesion** $a(x)$: average distance to points in the same cluster
  - **Separation** $b(x)$: minimum average distance to points in a different cluster
  - **Silhouette**:

$x \in C_X$

$$a(x) = \frac{1}{|C_X - 1|} \sum_{p \in C_X, p \neq x} d(x,p)$$

the smaller, the better

$$b(x) = \min_{X \neq K} \frac{1}{|C_K|} \sum_{p \in C_K} d(x,p)$$

the larger, the better

$$s(x) = \frac{b(x) - a(x)}{max\{a(x), b(x)\}}, \text{ if } |C_x| > 1$$

$$s(x) = 0, \text{ if } |C_x| = 1$$

Internal Quality Measures — Silhouette Coefficient

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

- Interpretation

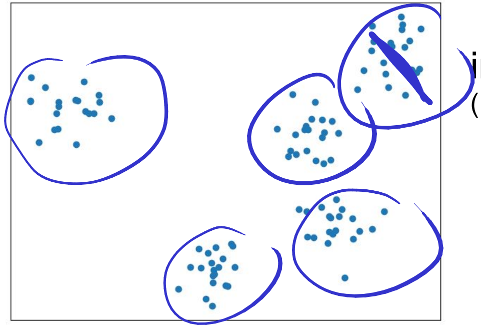
$$\underset{\textbf{BAD}}{\underline{-1}} \leq s(\underline{x}) \leq \underset{\textbf{GOOD}}{\underline{+1}}$$

- Silhouette Coefficient SC:

$$SC = \frac{1}{N} \sum_{i=1}^{N} s(x_i)$$

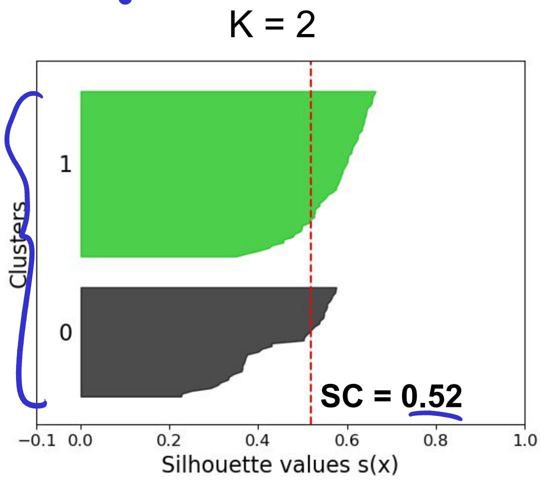
Internal Quality Measures — Silhouette Coefficient

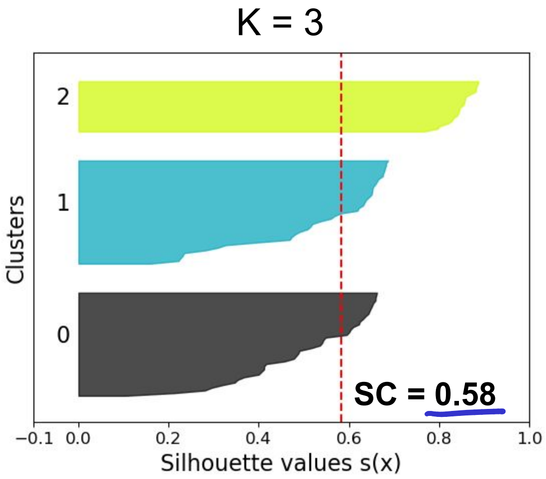
- Example: K-Means

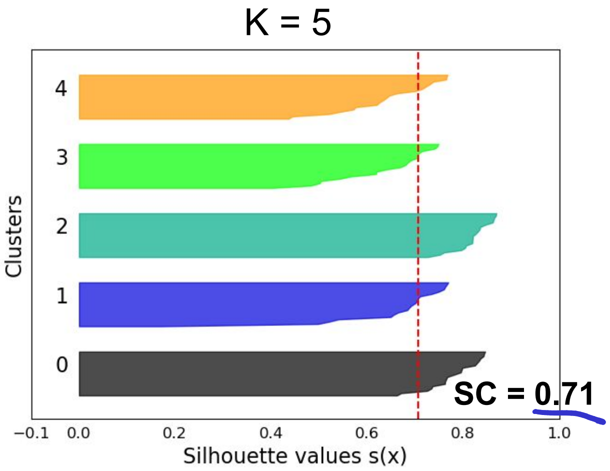


input data
(100 data points)

100 bars



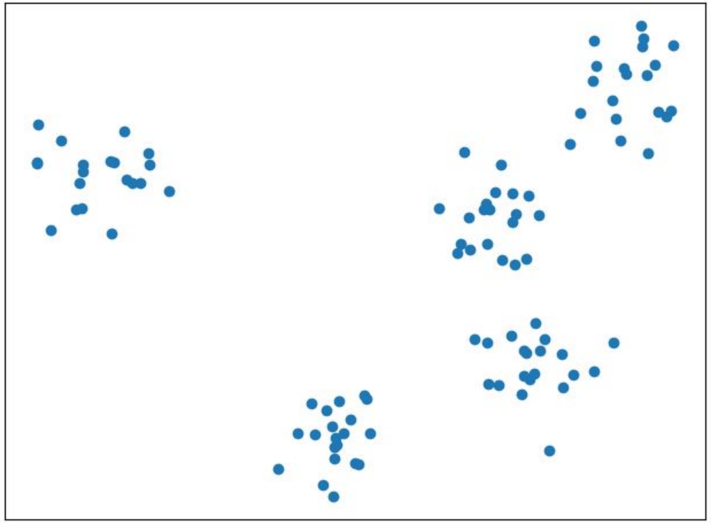




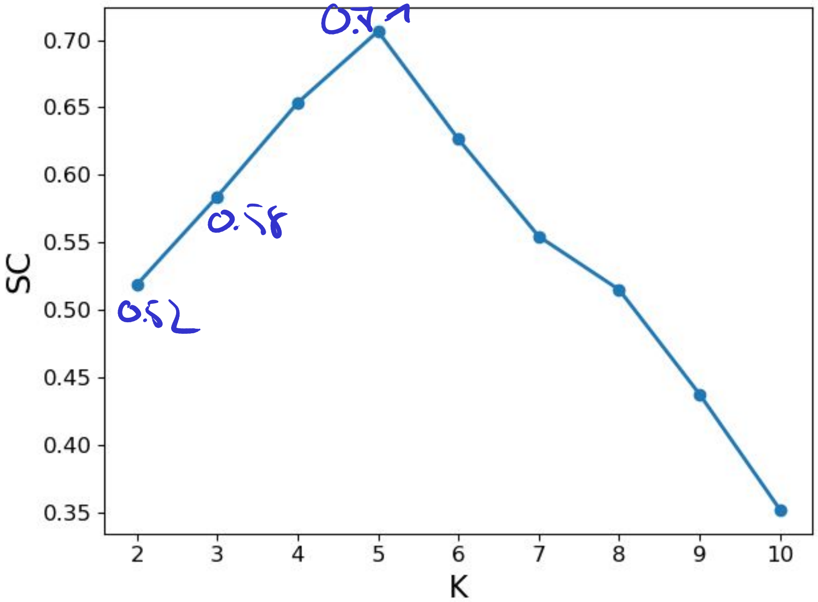
Internal Quality Measures — Silhouette Coefficient

- Example: K-Means

input data



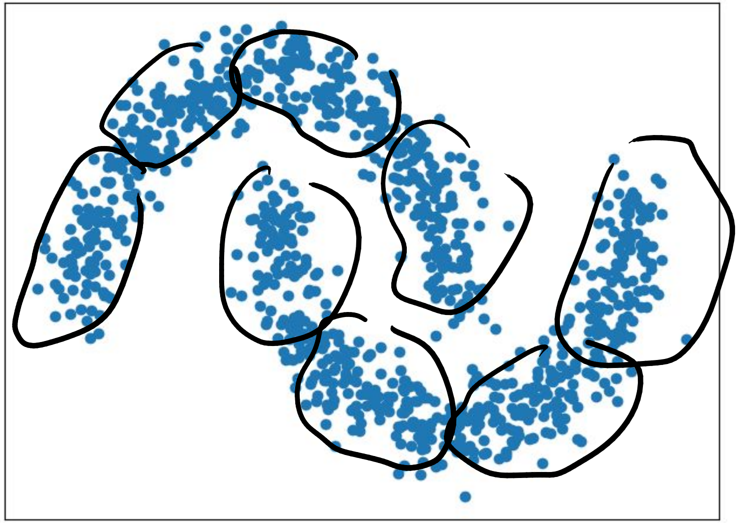
Silhouette Coefficients



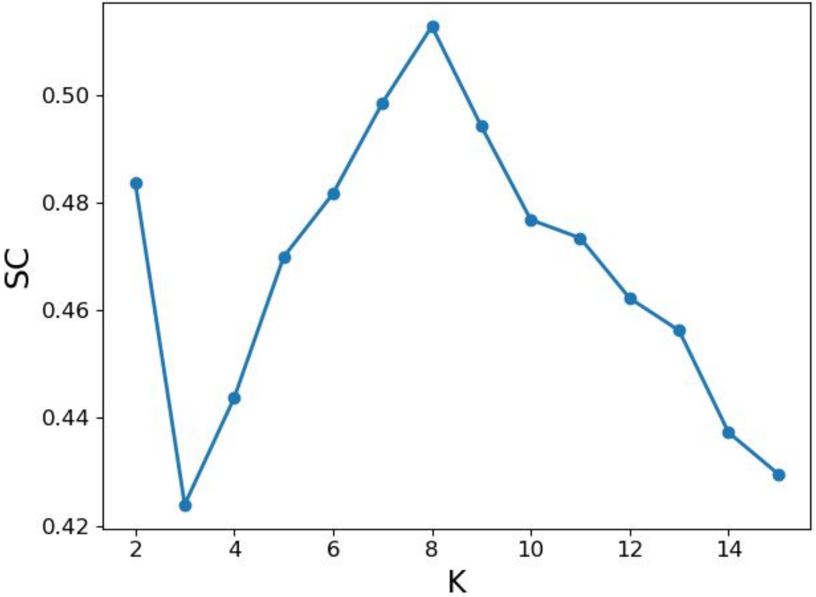
Internal Quality Measures — Silhouette Coefficient

- Example: K-Means

input data



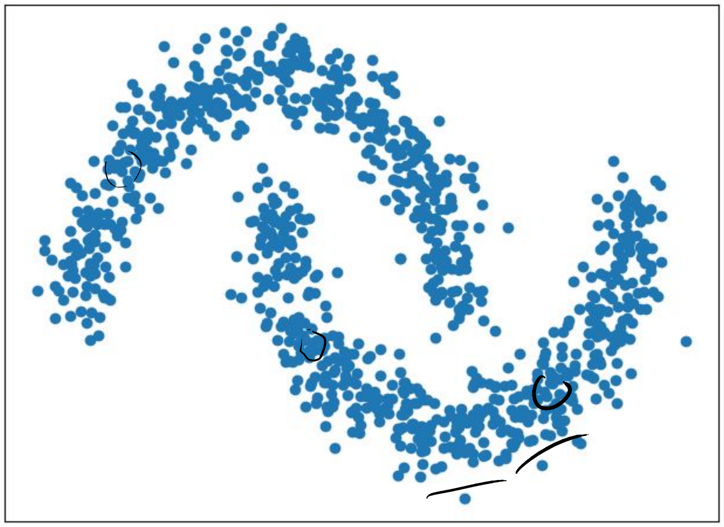
Silhouette Coefficients



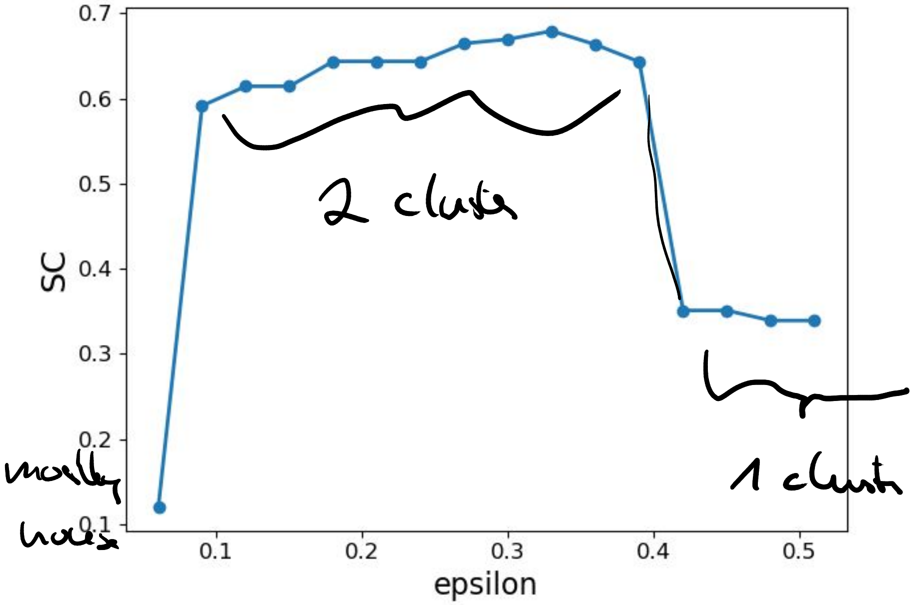
Internal Quality Measures — Silhouette Coefficient

- Example: DBSCAN

input data



Silhouette Coefficients



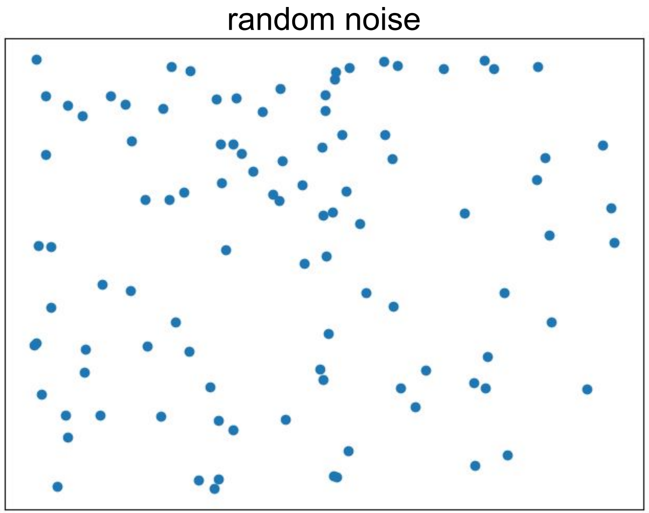
2 cluster

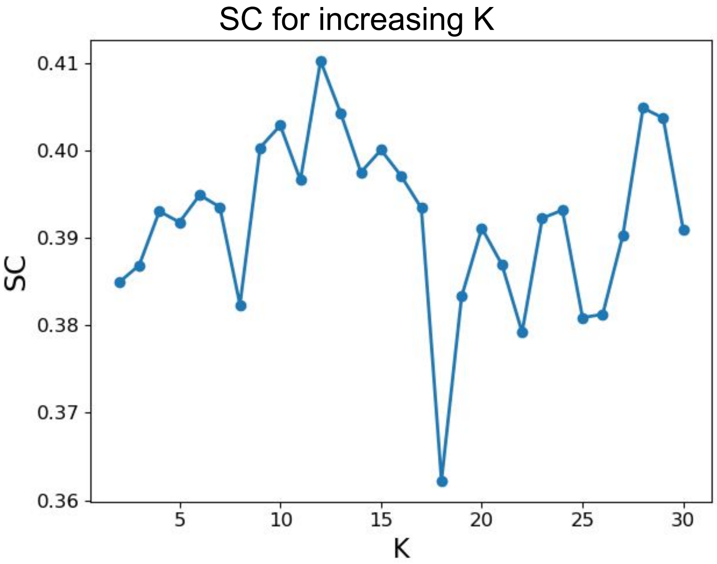
1 cluster

mostly noise

Internal Quality Measures — Silhouette Coefficient

- SC for random data (K-Means)





Note: DBSCAN on random data quickly results in 0 or 1 cluster, for which SC is not defined

Cluster Evaluation — Comments

- In practice, choice of "best" clustering often more pragmatic:
  - Fixed number of clusters (problematic for DBSCAN)
  - Parameters defined by tasks
    (e.g., "areas with more than 5 McDonalds within a radius of 500m")
  - Maximum, minimum, or average size of clusters
  - Focus in individual clusters instead of whole clustering
    (e.g., biggest/smallest cluster, cluster that contains certain points)
  - Set *K* "too high" and merge later if needed
  - ...

Outline

- Clustering
    - Overview
    - Concepts
    - Applications
- Clustering algorithms
    - K-Means
    - DBSCAN
    - Hierarchical Clustering
- Cluster Evaluation

Summary — Clustering

- Clustering: Finding patterns (here: cluster/groups) in unlabeled data
  - Very important concept in data mining
  - Wide range of clustering algorithms with varying characteristics (pros & cons)

➜ **No "one-size-fits-all" algorithm**

- Discussed algorithms: K-Means, DBSCAN, AGNES
  - Focus on the — arguably intuitive — conceptual inner workings
  - Emphasis on algorithms' strength and weaknesses
  - (Many tweaks and optimizations to improve performance)
- Major challenge: cluster evaluation
  - No fool-proof method to find the best algorithm or parameters (at least for unlabeled data)

Solutions to Quick Quizzes

- Slide 11: A
- Slide 15: A and/or B
- Slide 26: A
- Slide 46: A (in case of duplicates and K < #unique points)