Boston University CS 506 - Lance Galletti

K-means - Lloyd's Algorithm

Q1: Will this algorithm always converge?

Proof (by contradiction): Suppose it does not converge. Then, either:

- 1. The minimum of the cost function is only reached in the limit (i.e. after an infinite number of iterations).
 - **Impossible** because we are iterating over a finite set of partitions
- The algorithm gets stuck in a cycle / loop
 Impossible since this would require having a clustering that has a lower cost than itself and we know:
 - If old ≠ new clustering then the cost has improved
 - If old = new clustering then the cost is unchanged

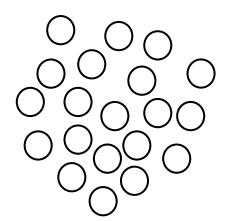
Conclusion: Lloyd's Algorithm always converges!

K-means - Lloyd's Algorithm

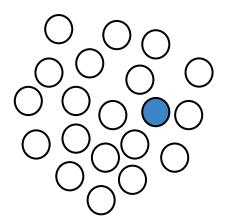
Q2: Will this always converge to the optimal solution?

NO

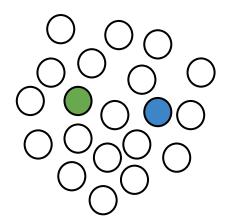
If the centers are too closed together, they would split up a naturally occuring cluster into 2 unnatural clusters



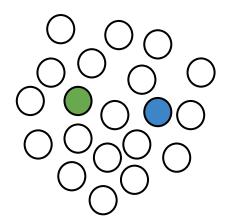


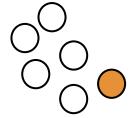


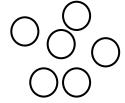


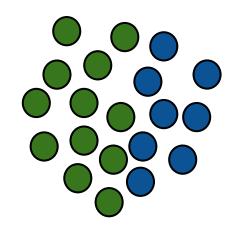


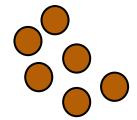


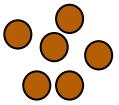




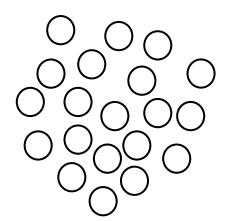




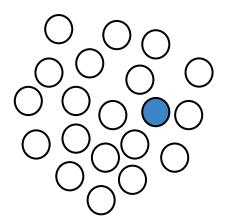




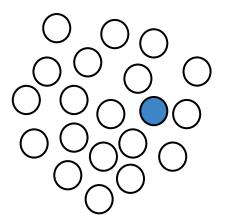
What's the problem?

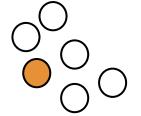


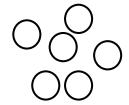


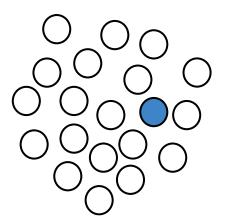


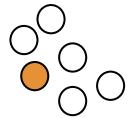


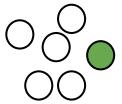


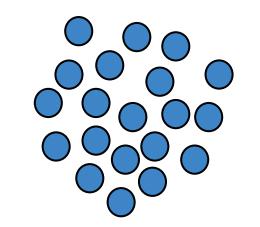


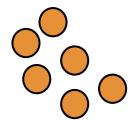


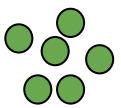




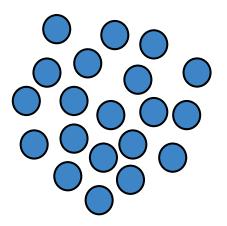


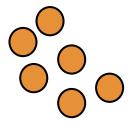


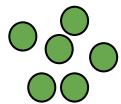




Farthest First Traversal

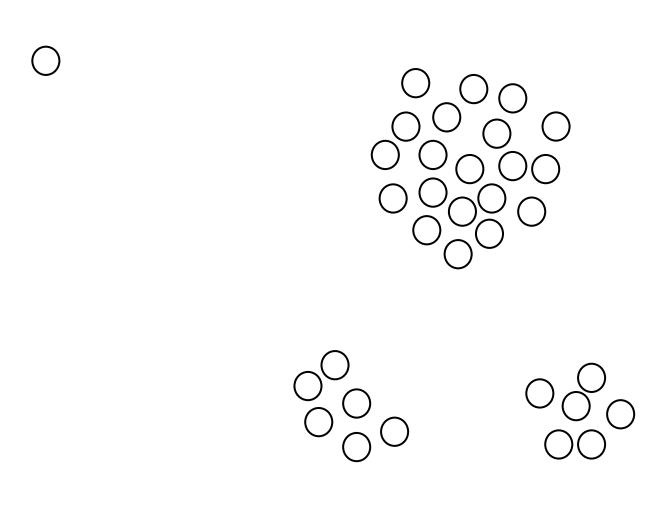




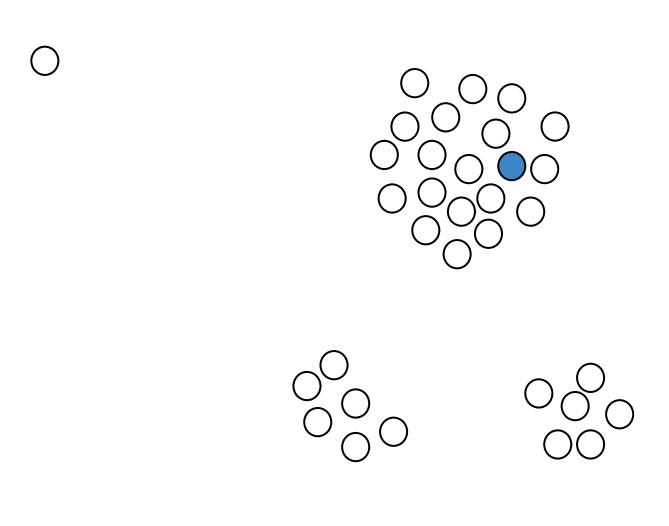


But...

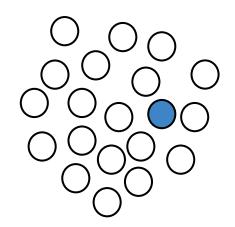
The problem is we could pick outliers as our centers

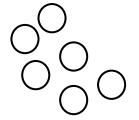


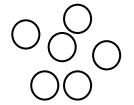


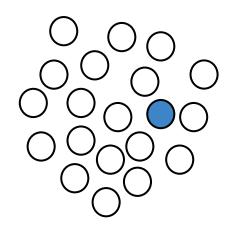


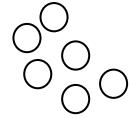


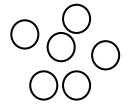


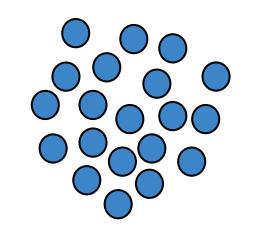


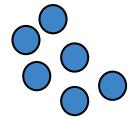


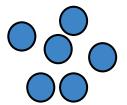




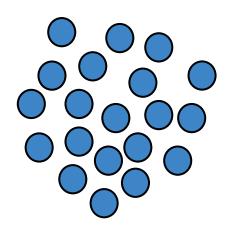


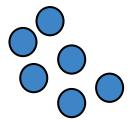


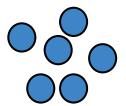




Random would have been better

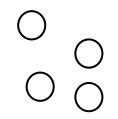


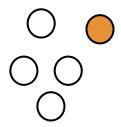


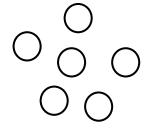


Initialize with a combination of the two methods:

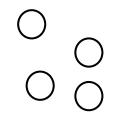
- 1. Start with a random center
- 2. Let D(x) be the distance between x and the closest of the centers picked so far. Choose the next center with probability proportional to $D(x)^2$

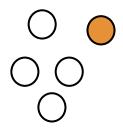


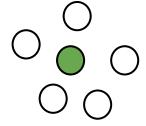




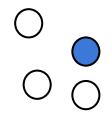


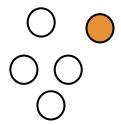


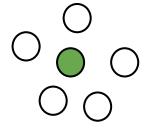




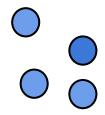


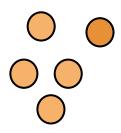




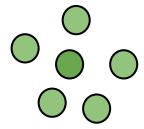




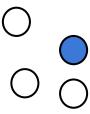


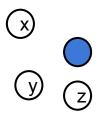


No reason to use k-means over k-means++







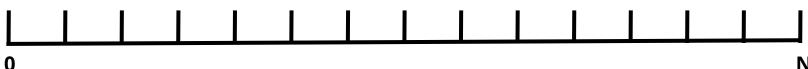


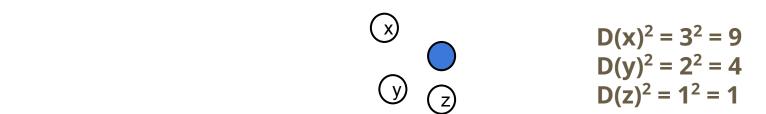
$$D(x)^2 = 3^2 = 9$$

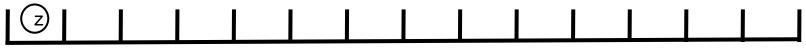
 $D(y)^2 = 2^2 = 4$
 $D(z)^2 = 1^2 = 1$

Suppose we are given a black box that will generate a uniform random number between 0 and any **N**. How can we use this black box to select points

with probability proportional to $D(x)^2$? 9+4+1=14so x gets 9 out of 14 spots y gets 4 out of 14 z gets 1 out of 14 $D(x)^2 = 3^2 = 9$ $D(y)^2 = 2^2 = 4$ $D(z)^2 = 1^2 = 1$







$$D(x)^{2} = 3^{2} = 9$$

$$D(y)^{2} = 2^{2} = 4$$

$$D(z)^{2} = 1^{2} = 1$$



$$D(x)^{2} = 3^{2} = 9$$

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$$D(z)^{2} = 1^{2} = 1$$

$$D(x)^{2} = 3^{2} = 9$$

$$D(y)^{2} = 2^{2} = 4$$

$$D(z)^{2} = 1^{2} = 1$$

K-means++

Q3: the black box returns "12" as the random number generated. Which point do we choose for the next center (x, y, or z)?



0

K-means++

Q4: the black box returns "4" as the random number generated. Which point do we choose for the next center (x, y, or z)?

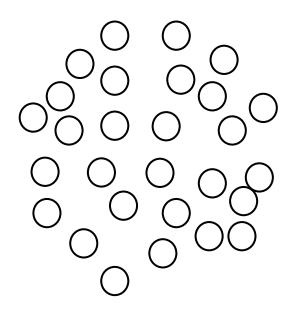


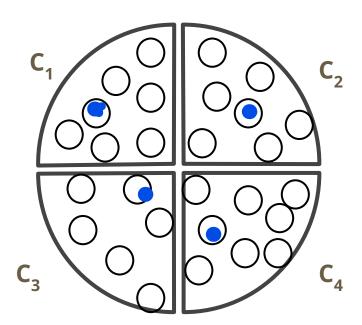
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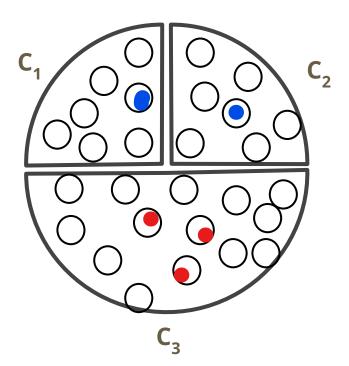
K-means++

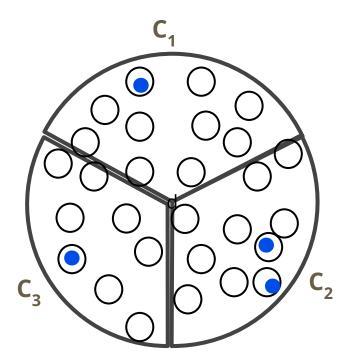
What happens if the black box can only generate numbers between 0 and 1?

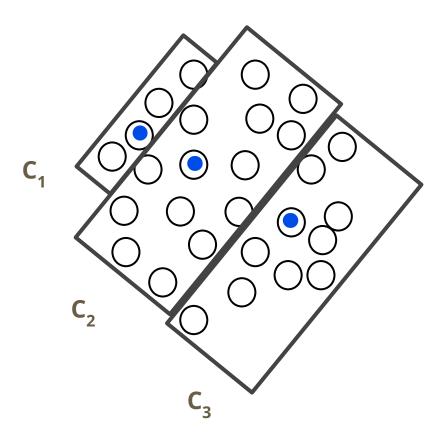
Kmeans Quizz (take 2)

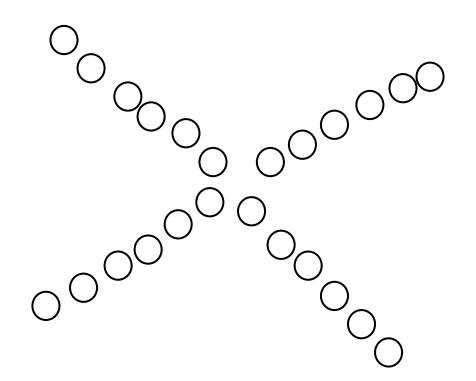


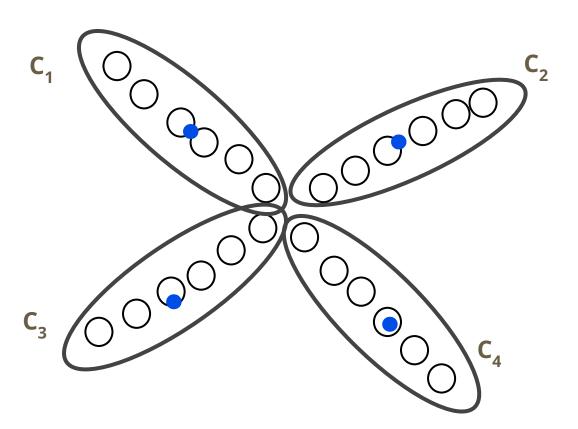


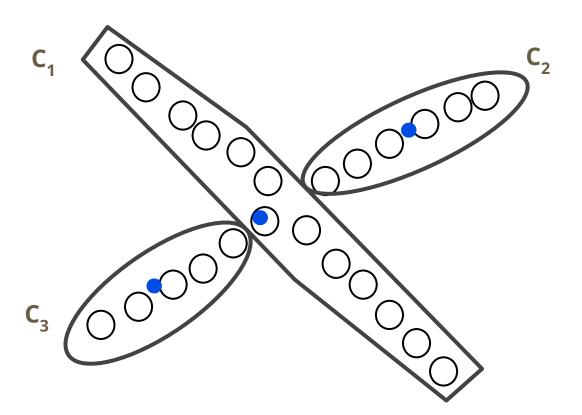


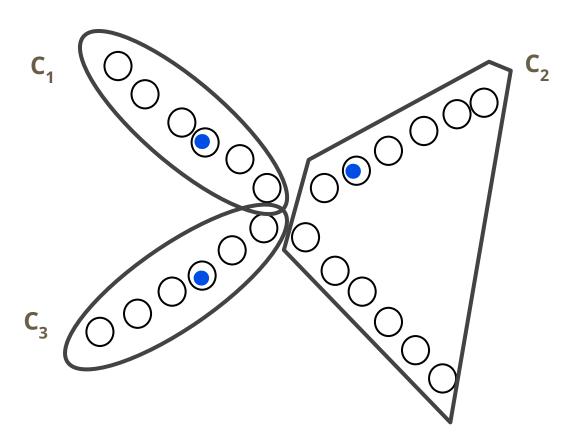


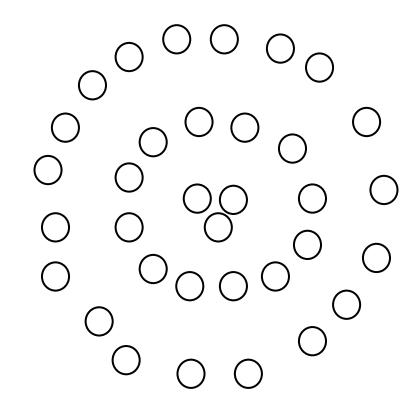


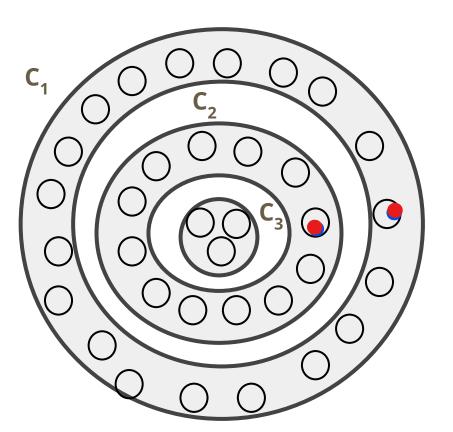


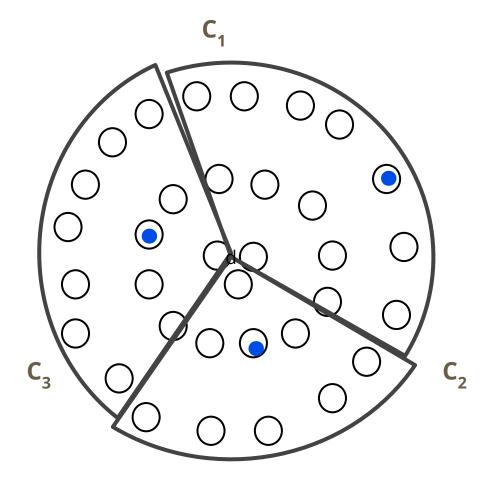


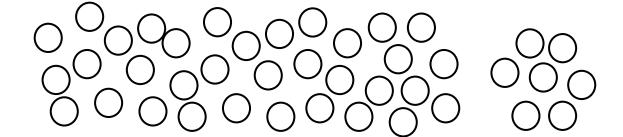


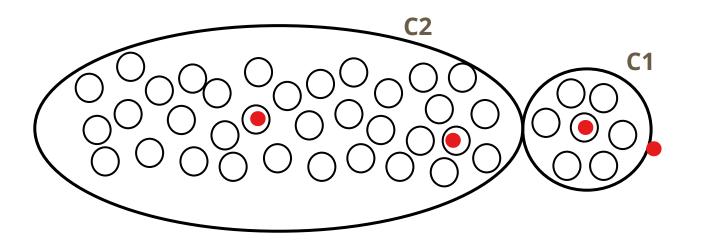






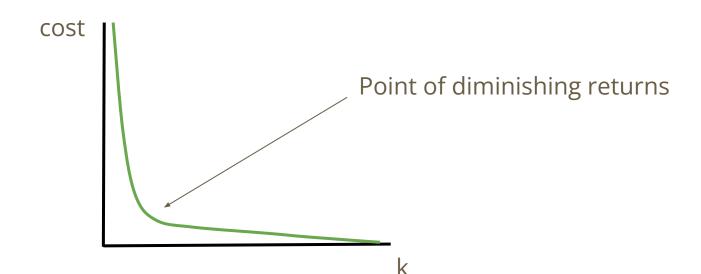






How to choose the right k?

1. Iterate through different values of k (elbow method)



How to choose the right k?

- 1. Iterate through different values of k (elbow method)
- Use empirical / domain-specific knowledge
 Example: Is there a known approximate distribution of the data? (K-means is good for spherical gaussians)
- 3. Metric for evaluating a clustering output

Evaluation

Recall our goal: Find a clustering such that

- Similar data points are in the same cluster
- Dissimilar data points are in different clusters

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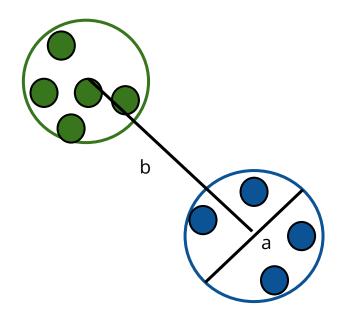
Evaluation

K-means cost function tells us the within-cluster distances between points will be small overall.

But what about the intra-cluster distance? Are the clusters we created far? How far? Relative to what?

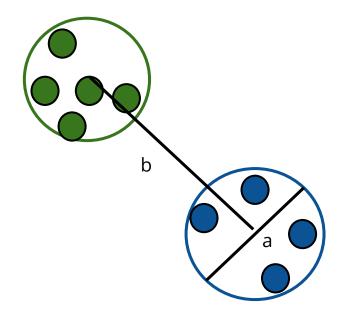
Discuss - 5min

Define a metric that evaluates how spread out the clusters are from one another.



a: average within-cluster distance

b: average intra-cluster distance



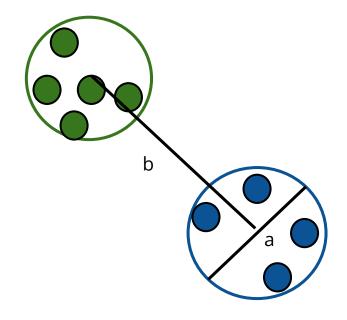
The distance within the cluster is the same as the distance between the clusters

a: average within-cluster distance

b: average intra-cluster distance

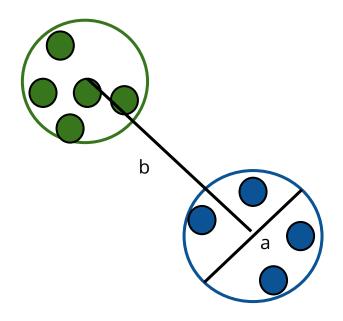
What does it mean for (b - a) to be 0?

means clusters are right next to each other

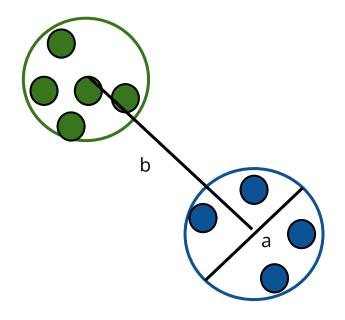


a: average within-cluster distanceb: average intra-cluster distance

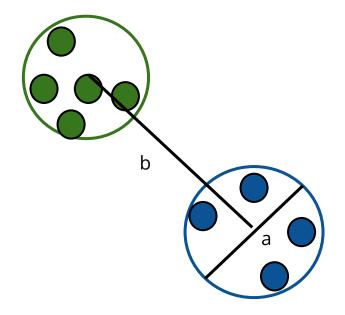
What does it mean for (b - a) to be large?



The value of (b-a) doesn't mean much by itself. Can we compare it to something so that the ratio becomes a value between 0 and 1?

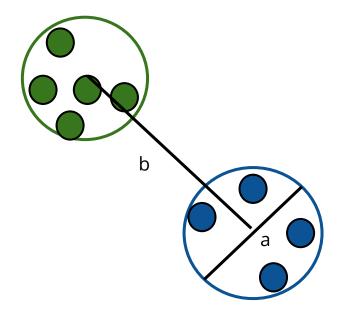


(b - a) / max(a, b)



What does it mean for (b - a) / max(a, b) to be close to 1?

A value close to 1 indicates that the data points are well-separated from each other and from other clusters, while a value close to -1 indicates that the data points are not well-separated

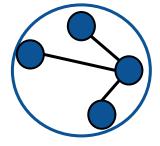


What does it mean for (b - a) / max(a, b) to be close to 0?

A value close to 1 indicates that the data points are well-separated from each other and from other clusters, while a value close to -1 indicates that the data points are not well-separated



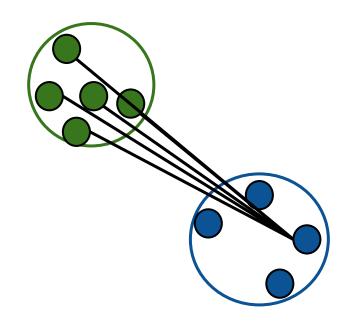
For each data point i: a_i: mean distance from point i to every other point in its cluster



For each data point i:

a_i: mean distance from point i to every other point in its cluster

b_i: smallest mean distance from point i to every point in another cluster





For each data point i:

a_i: mean distance from point i to every other point in its cluster

b_i: smallest mean distance from point i to every point in another cluster



$$s_i = (b_i - a_i) / max(a_i, b_i)$$

si should be small in theory

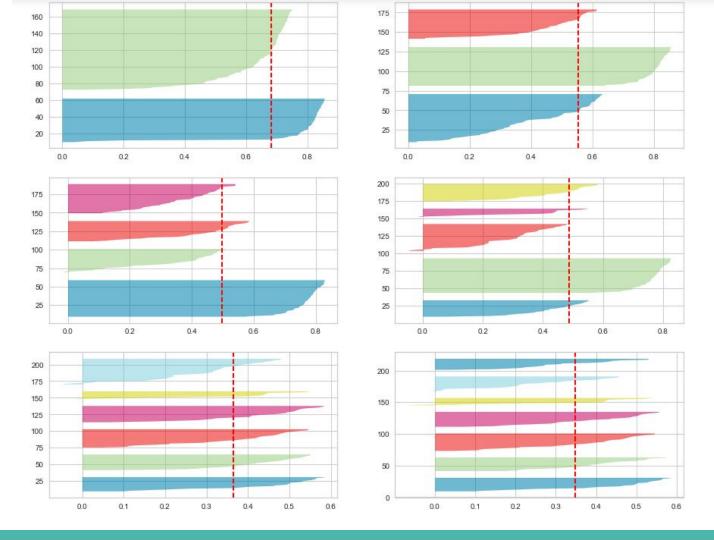
The silhouette score ranges from -1 to 1, where a score close to 1 indicates that the samples are well-clustered, a score close to 0 indicates overlapping clusters, and a negative score indicates that the samples might have been assigned to the wrong cluster

$$s_i = (b_i - a_i) / max(a_i, b_i)$$

Silhouette score plot



OR return the mean s_i over the entire dataset as a measure of goodness of fit



K-means Variations

- 1. K-medians (uses the L₁ norm / manhattan distance)
- 2. K-medoids (any distance function + the centers must be in the dataset)
- 3. Weighted K-means (each point has a different weight when computing the mean)