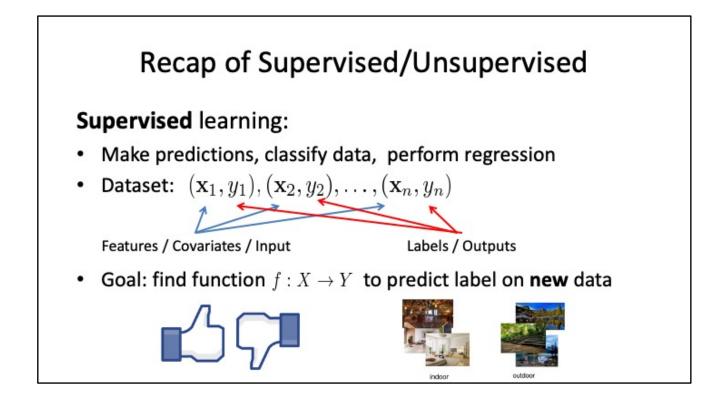
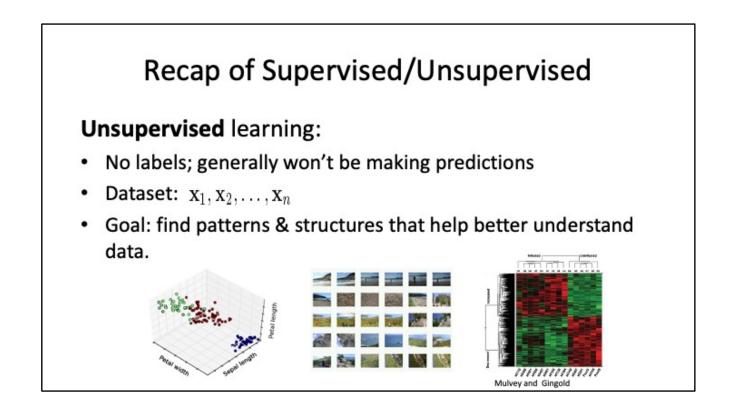
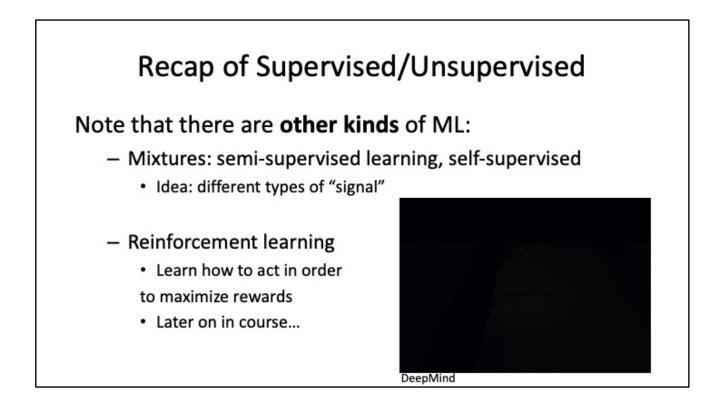


CS 540 Introduction to Artificial Intelligence Unsupervised Learning I

Yingyu Liang University of Wisconsin-Madison Oct 5, 2021 Based on slides by Fred Sala







Outline

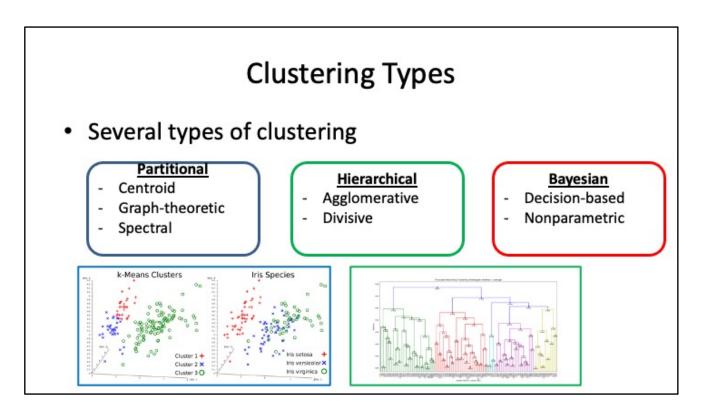
- Intro to Clustering
 - Clustering Types, Centroid-based, k-means review
- Hierarchical Clustering
 - Divisive, agglomerative, linkage strategies

Unsupervised Learning & Clustering

- Note that clustering is just one type of unsupervised learning (UL)
 - PCA is another unsupervised algorithm
- Estimating probability distributions also UL (GANs)
- Clustering is popular & useful!



StyleGAN2 (Kerras et al '20)



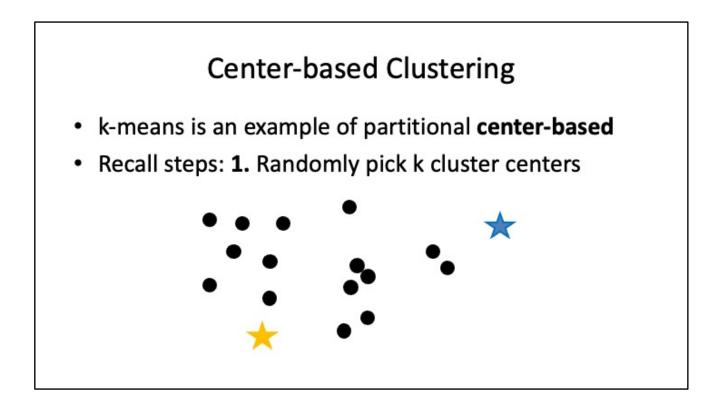
Partitional: to get a partition (ie, a set of disjoint clusters whose union is the whole dataset)

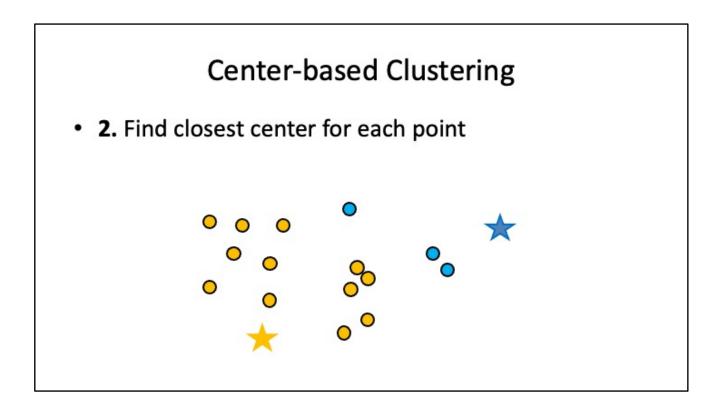
- 1) centroid: use centers and assign data points to centers to form clusters
- 2) Graph-theoretical: the input is a graph (instead of a set of numeric vectors), and would like to partition the nodes into clusters
- 3) Spectral: an approach for doing graph clustering

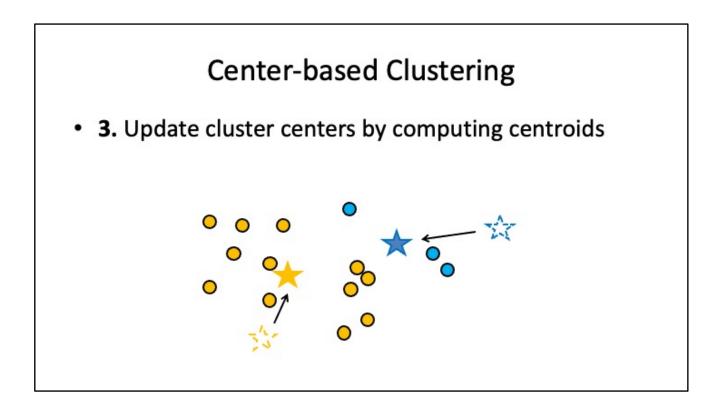
Hierarchical: to get a tree on the data points

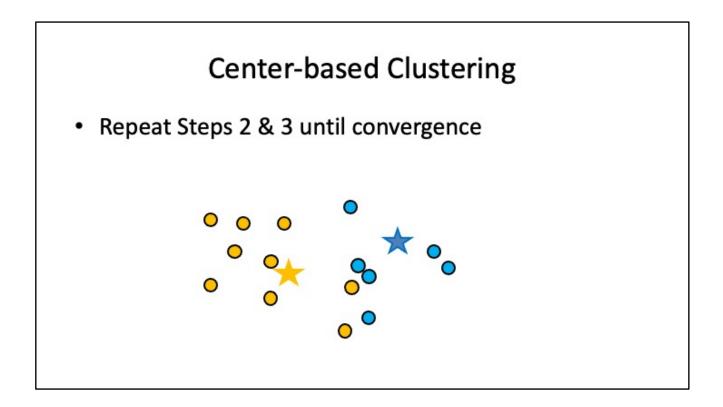
- 1) Agglomerative: begin with each point as a singleton cluster, and keep merging them until all are merged into one cluster
- 2) Divisive: begin with all points in one cluster, and keep splitting the clusters to smaller ones until containing only one point (or satisfying some other stopping criteria)

Bayesian: a family of methods using Bayes' rule to do clustering. Can produce a partition or a tree. Not covered in this course.









Q 1.1: You have seven 2-dimensional points. You run 3-means on it, with initial clusters

 $C_1 = \{(2, 2), (4, 4), (6, 6)\}, C_2 = \{(0, 4), (4, 0)\}, C_3 = \{(5, 5), (9, 9)\}$

Cluster centroids at the next iteration are?

- A. C₁: (4,4), C₂: (2,2), C₃: (7,7)
- B. C₁: (6,6), C₂: (4,4), C₃: (9,9)
- C. C₁: (2,2), C₂: (0,0), C₃: (5,5)
- D. C₁: (2,6), C₂: (0,4), C₃: (5,9)

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- D. C₁: (2,6), C₂: (0,4), C₃: (5,9)

The average of points in C1 is (4,4). The average of points in C2 is (2,2). The average of points in C3 is (7,7).

Q 1.2: We are running 3-means again. We have 3 centers, C_1 (0,1), C_2 , (2,1), C_3 (-1,2). Which cluster assignment is possible for the points (1,1) and (-1,1), respectively? Ties are broken arbitrarily:

(i) C₁, C₁ (ii) C₂, C₃ (iii) C₁, C₃

- A. Only (i)
- B. Only (ii) and (iii)
- C. Only (i) and (iii)
- D. All of them

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- A. Only (i)
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For the point (1,1): square-Euclidean-distance to C1 is 1, to C2 is 1, to C3 is 5 So it can be assigned to C1 or C2

For the point (-1,1): square-Euclidean-distance to C1 is 1, to C2 is 9, to C3 is 1 So it can be assigned to C1 or C3

Q 1.3: If we run K-means clustering twice with random starting cluster centers, are we guaranteed to get same clustering results? Does K-means always converge?

- A. Yes, Yes
- B. No, Yes
- C. Yes, No
- D. No, No

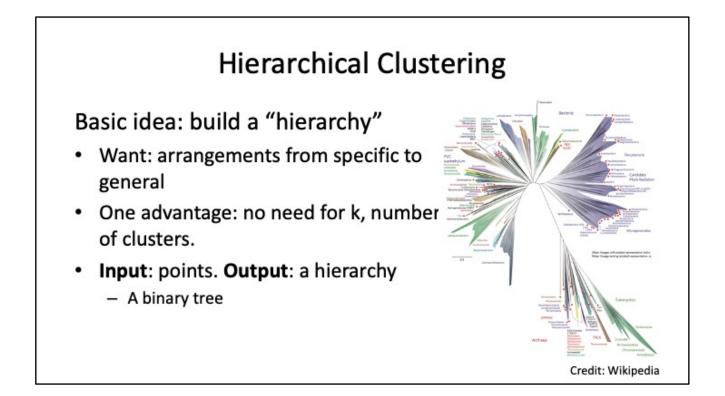
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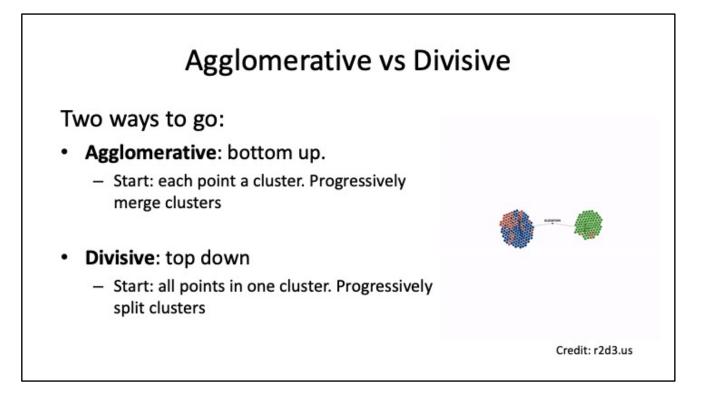
The clustering from k-means will depend on the initialization. Different initialization can lead to different outcomes.

K-means will always converge on a finite set of data points:

- 1. There are finite number of possible partitions of the points
- 2. The assignment and update steps of each iteration will only decrease the sum of the distances from points to their corresponding centers.
- 3. If it run forever without convergence, it will revisit the same partition, which is contradictory to item 2.

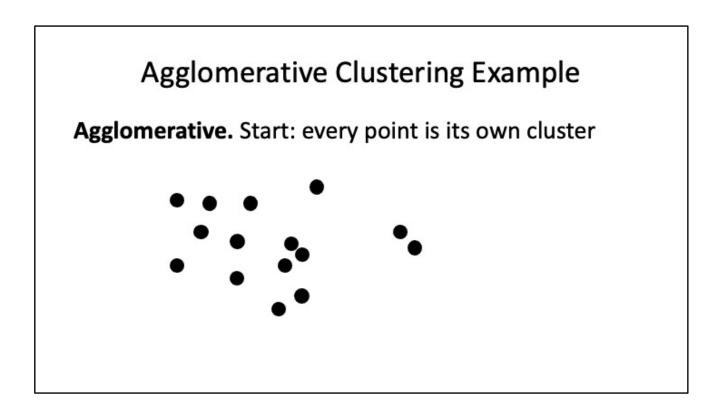


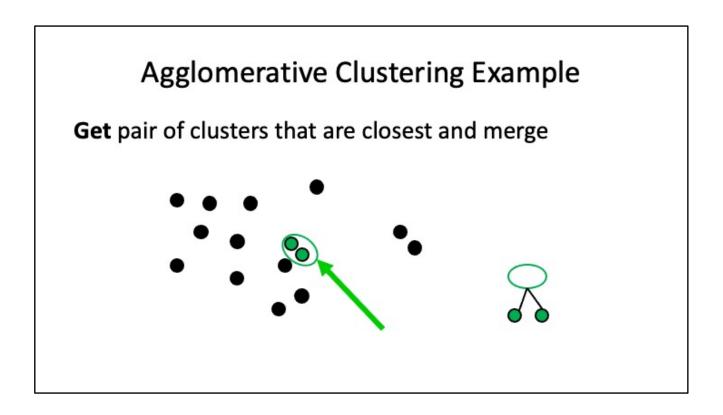
Typically, the algorithms build a binary tree (each node only has 2 children). Sometimes can be a tree with branching factor more than 2.

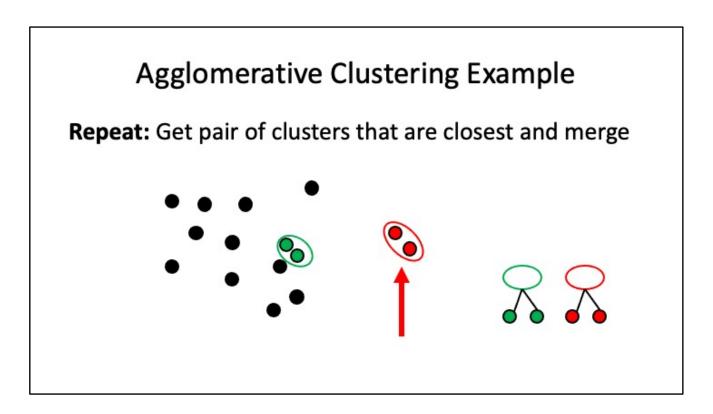


Hierarchical: to get a tree on the data points

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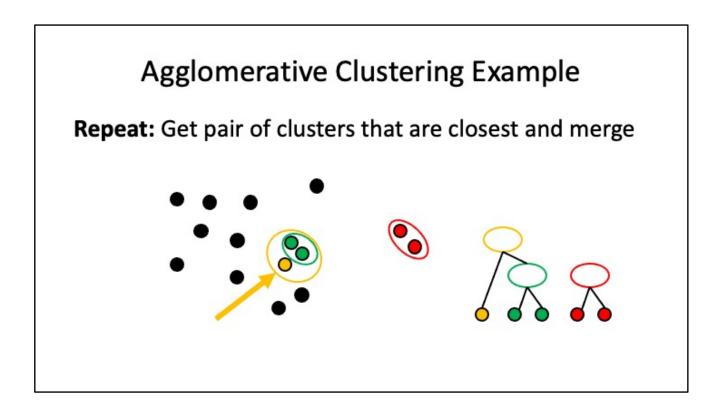


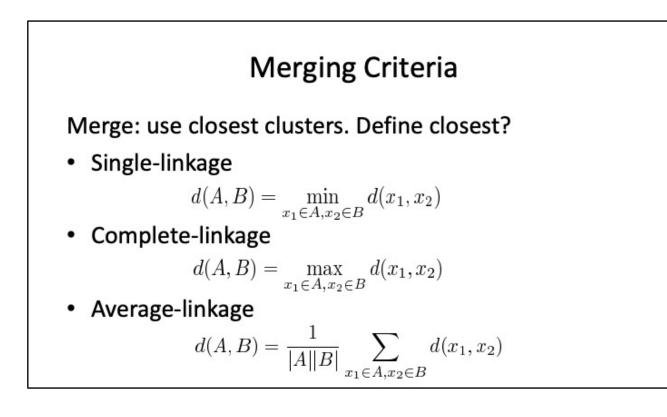




Keep merging the closest pair of clusters.

We only have a definition of distance between data points. Need a definition of distance between clusters!





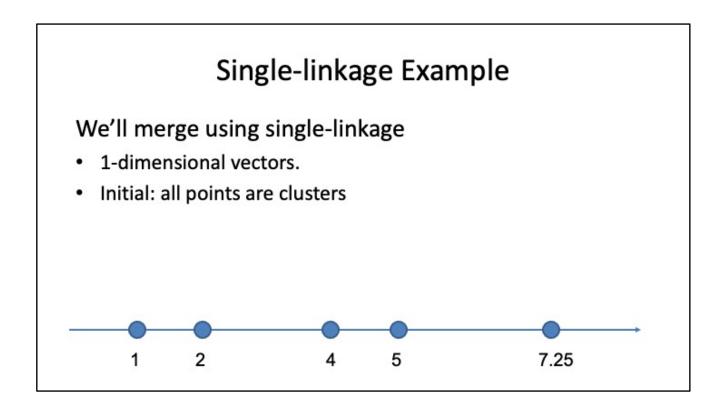
We can have different definitions of distances between clusters, which lead to different algorithms.

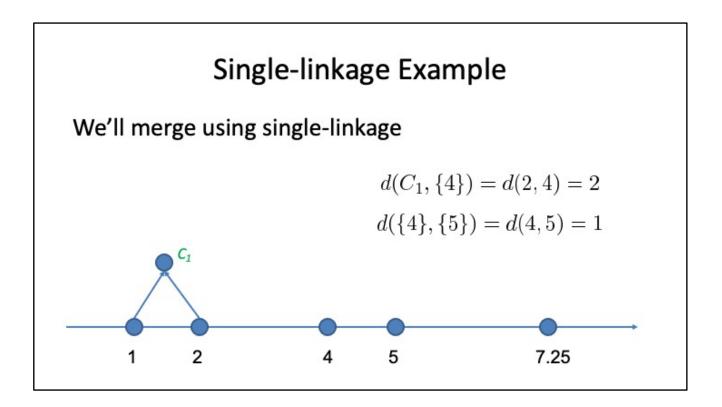
Once we have the definition, we can compute the distances and find the closest pair of clusters and merge them.

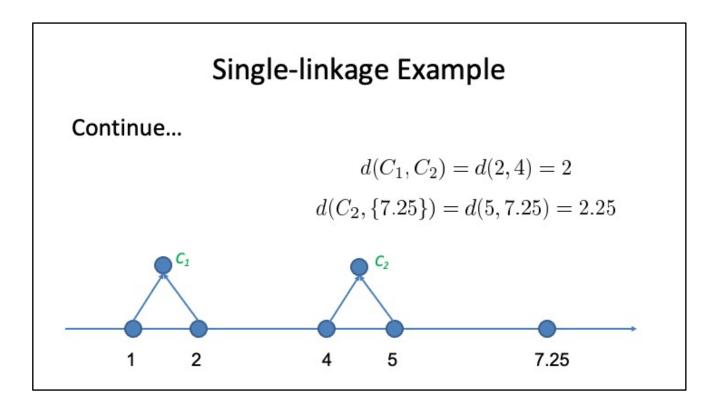
Note: in complete-linkage, we find the closest pair of clusters by

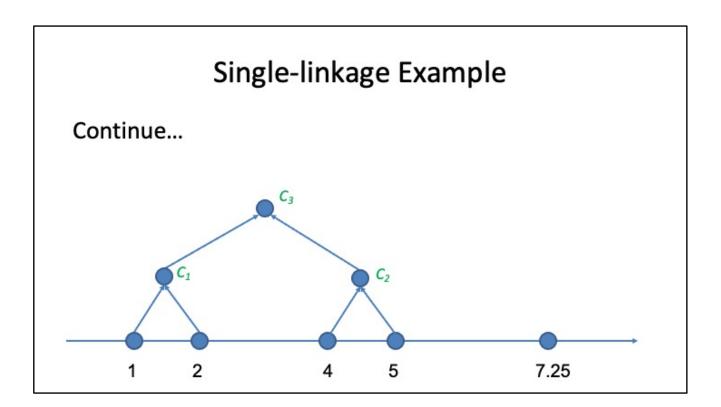
 $(A^*, B^*) = argmin_{clusters A B} d(A,B) = argmin_{clusters A B} max_{x1 \in A, x2} \in B d(x1, x2)$

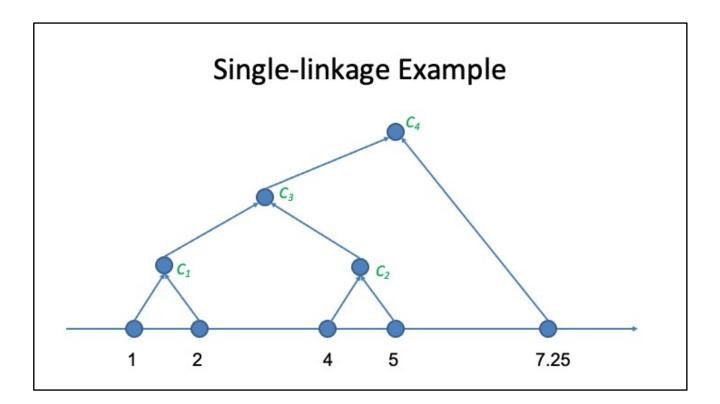
Do not confuse the max over data points with the min over clusters. That is, while we compute the distance between clusters, we take the maximum over the points; but we are still looking for the closest pair of clusters, not the farthest pair of clusters.

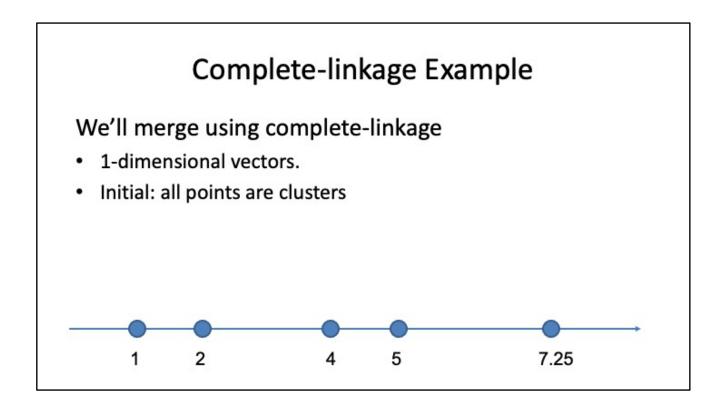


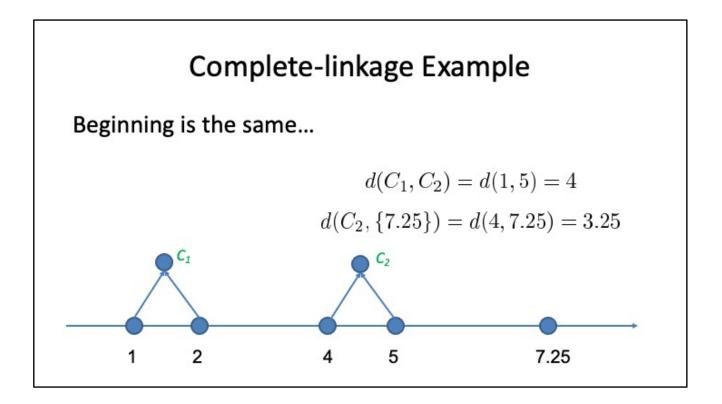


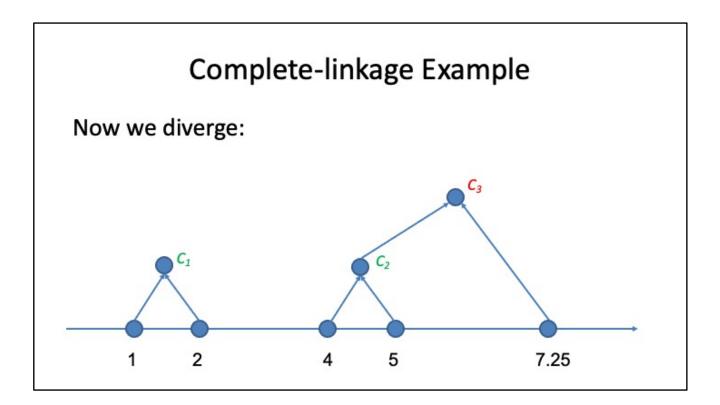


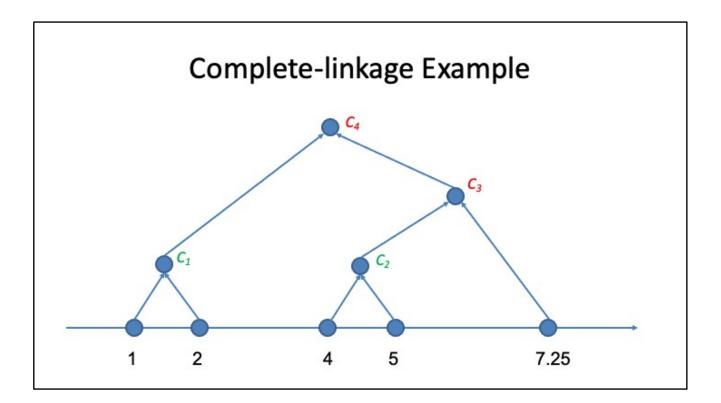


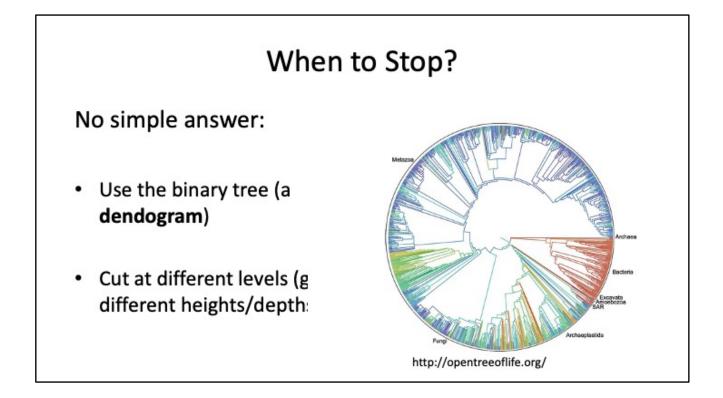




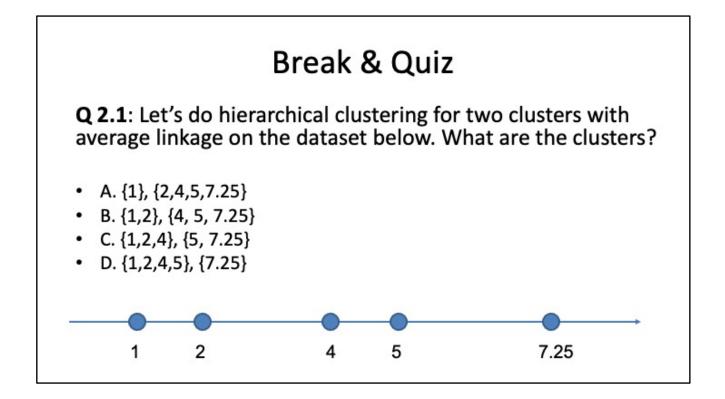


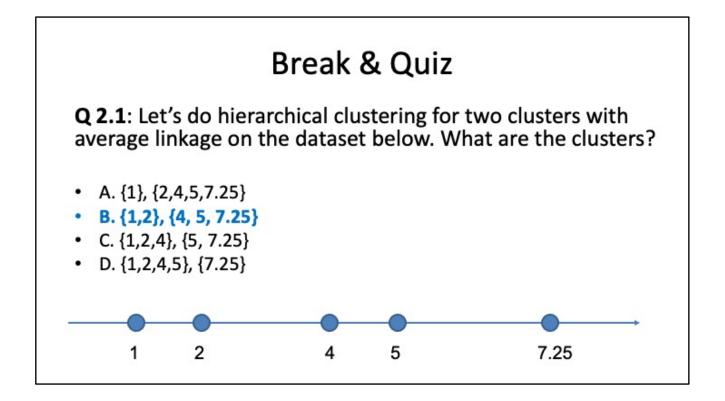






Typical in practice: merge until only one cluster (the root). Then cut at different levels to get different partitions; number of clusters or the cut level is application-dependent.





Iteration 1: merge 1 and 2 Iteration 2: merge 4 and 5 Iteration 3: Now we have clusters $\{1,2\}, \{4,5\}, \{7.25\}$. distance($\{1,2\}, \{4,5\}$)= 3 distance($\{4,5\}, \{7.25\}$) = 2.75 distance($\{1,2\}, \{7.25\}$) is clearly larger than the above two. So average linkage will merge $\{4,5\}$ and $\{7.25\}$

Q 2.2: If we do hierarchical clustering on n points, the maximum depth of the resulting tree is

- A. 2
- B. log n
- C. n/2
- D. n-1

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Denote the points as x_1, x_2, ..., x_n

```
Suppose:
in iteration 1, we merge points x_1 and x_2
in iteration 2, we merge \{x_1, x_2\} with x_3
...
in iteration t, we merge \{x_1, x_2, ..., x_t\} with x_\{t+1\}
...
in iteration n-1, we merge \{x_1, x_{n-1}\} with x_n
```

Then we will get a tree with depth n-1.