





- It is time to explore another clustering method!
- Hierarchical clustering is very common in biology and lends itself nicely to visualizing clusters.
- It can also help the user decide on an appropriate number of clusters.





- Section Overview:
  - Theory and Intuition of Hierarchical Clustering
  - Coding Example of Hierarchical Clustering





Theory and Intuition





- Like most clustering algorithms,
   Hierarchical Clustering simply relies on
   measuring which data points are most
   "similar" to other data points.
- "Similarity" is defined by choosing a distance metric.





#### So why use Hierarchical Clustering?

- Easy to understand and visualize.
- Helps users decide how many clusters to choose.
- Not necessary to choose cluster amount **before** running the algorithm.





#### So why use Hierarchical Clustering?

- o Divides points into *potential* clusters:
  - Agglomerative Approach:
    - Each point begins as its own cluster, then clusters are joined.
  - Divisive Approach:
    - All points begin in the same cluster, then clusters are split.



- Hierarchical Clustering
  - Divides points into potential clusters:

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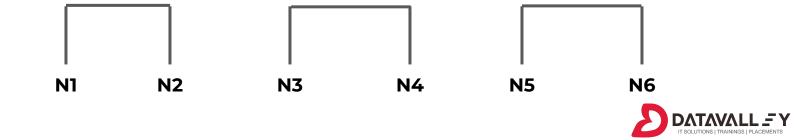


- Hierarchical Clustering
  - Agglomerative:



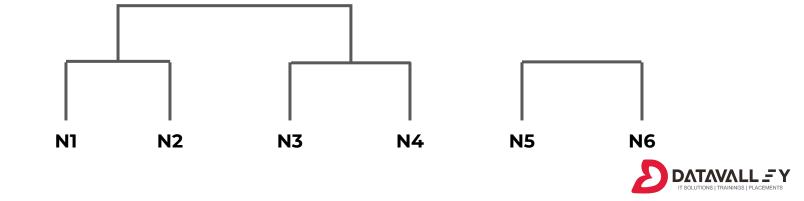


- Hierarchical Clustering
  - Agglomerative:



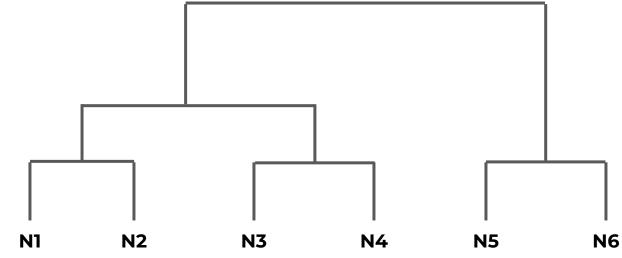


- Hierarchical Clustering
  - Agglomerative:





- Hierarchical Clustering
  - Agglomerative:







 Opposite of the Agglomerative approach is a **Divisive** approach, which starts with all points belonging to the same cluster, and the begins divisions to separate out clusters.



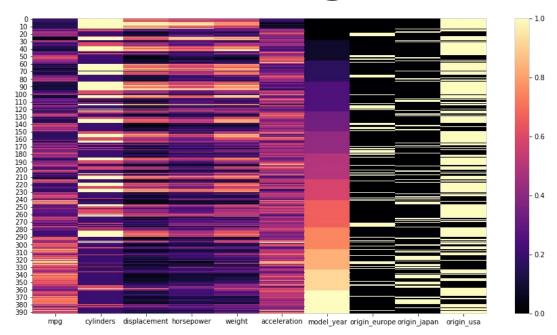


#### Hierarchical Clustering Process

- Compare data points to find most similar data points to each other.
- Merge these to create a cluster.
- Compare clusters to find most similar clusters and merge again.
- Repeat until all points in a single cluster.



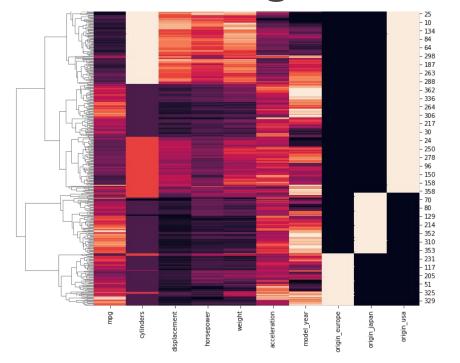
#### Hierarchical Clustering Process







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- There are a few key topics we still need to understand for Hierarchical Clustering:
  - Similarity Metric
  - Dendrogram
  - Linkage Matrix



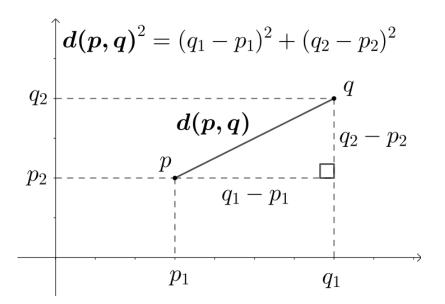


- Similarity Metric
  - Measures distance between two points.
  - Many options:
    - Euclidean Distance
    - Manhattan
    - Cosine
    - and many more...





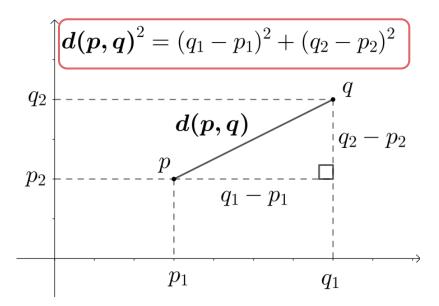
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  - Default choice is Euclidean







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- Similarity Metric
  - Each dimension would be a feature
  - For **n** data points and **p** features:





- Similarity Metric
  - Each dimension would be a feature
  - For **n** data points and **p** features:

- Using MinMaxScaler we can scale all features to be between 0 and 1.
- This allows for maximum distance between a feature to be 1.

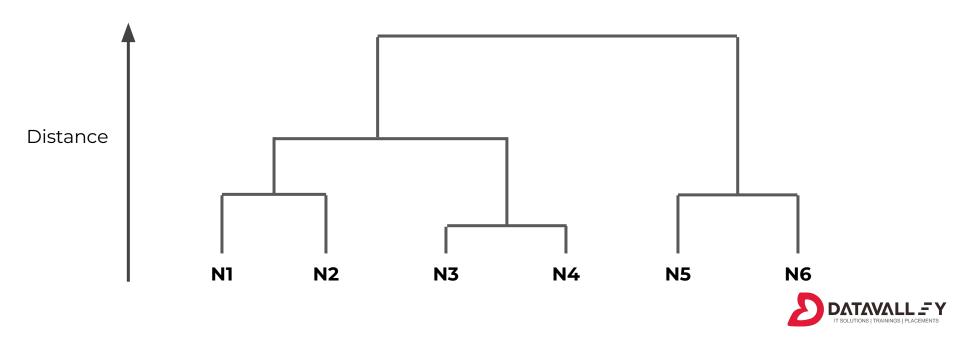


- Dendrogram:
  - Plot displaying all potential clusters.
  - Very computationally expensive to compute and display for larger data sets.
  - Very useful for deciding on number of clusters.



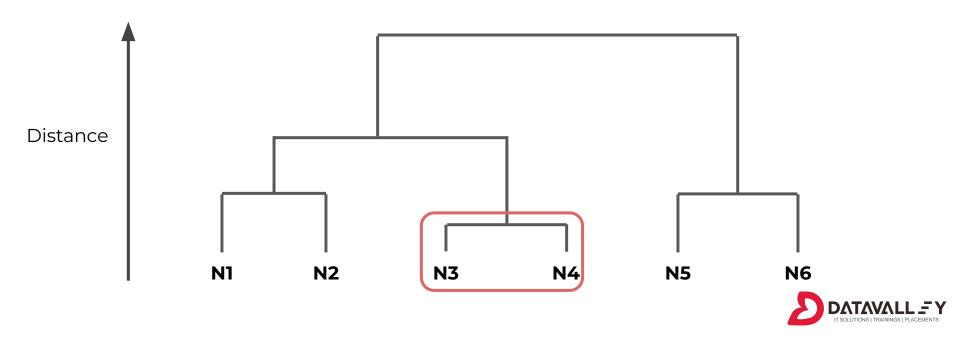


• Dendrogram:



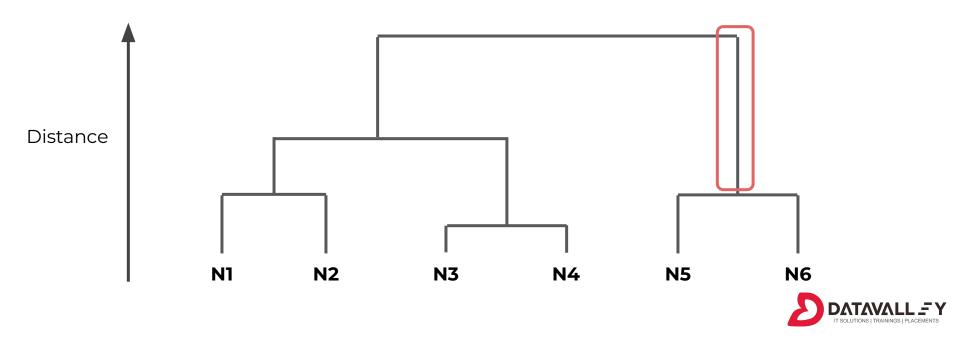


• Dendrogram:



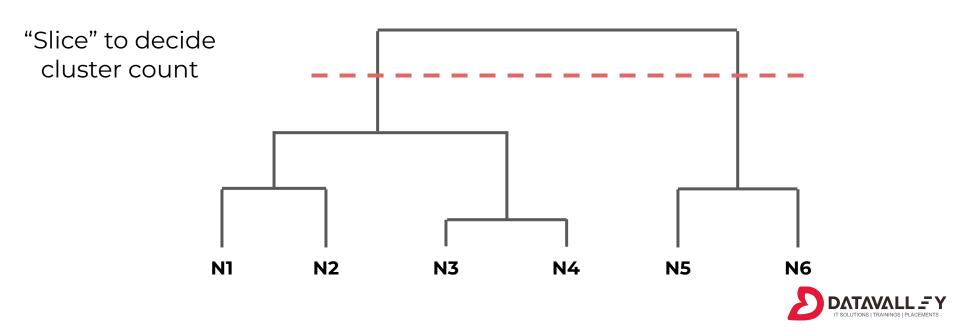


• Dendrogram:



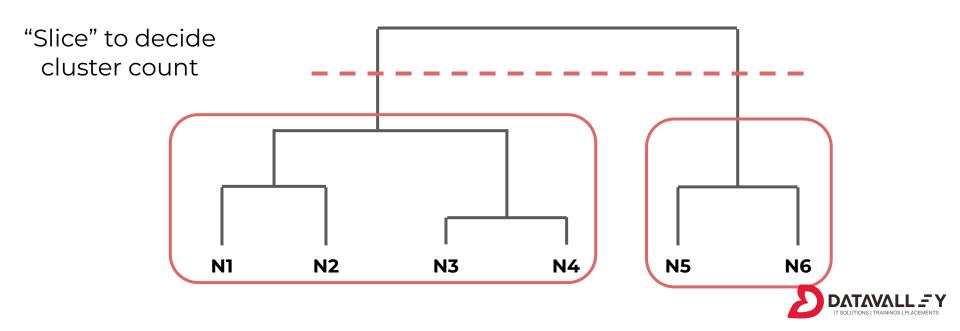


Dendrogram:



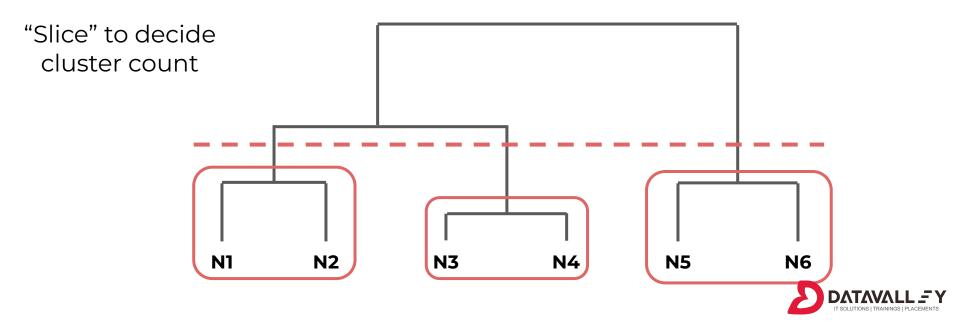


Dendrogram:





Dendrogram:





- Linkage
  - How do we measure distance from a point to an entire cluster?
  - How do we measure distance from a cluster to another cluster?

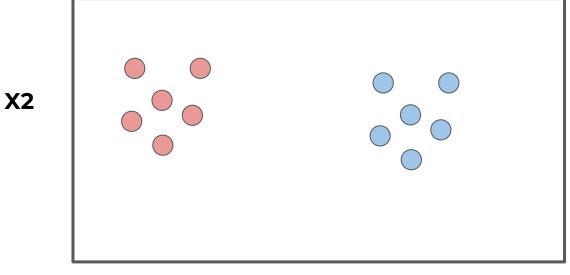




- Linkage
  - Once two or more points are together and we want to continue agglomerative clustering to join clusters, we need to decide on a linkage parameter.

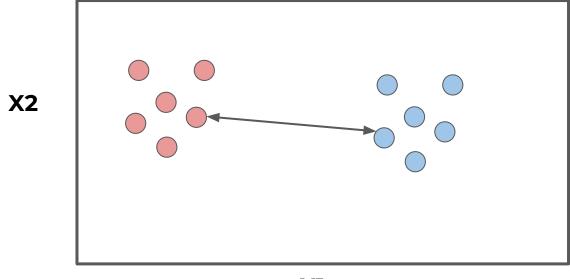






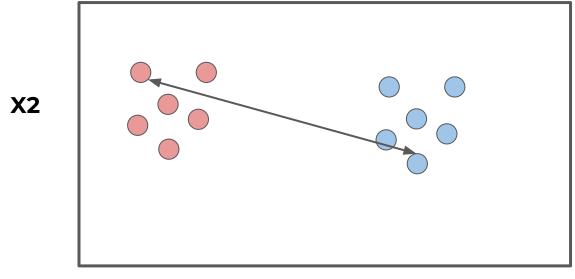






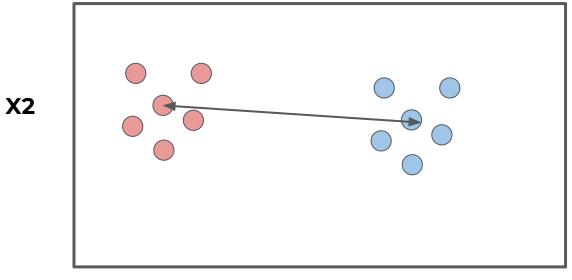






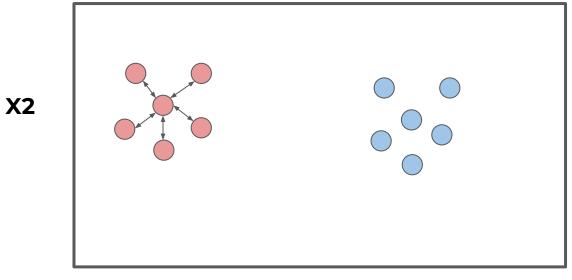
















- Linkage
  - Criterion determining which distance to use between sets of observation.
  - Algorithm will merge pairs of clusters that minimizes the criterion.





- Linkage:
  - Ward: minimizes variance of clusters being merged.
  - Average: uses average distances between two sets.
  - Minimum or Maximum distances between all observations of the two sets.