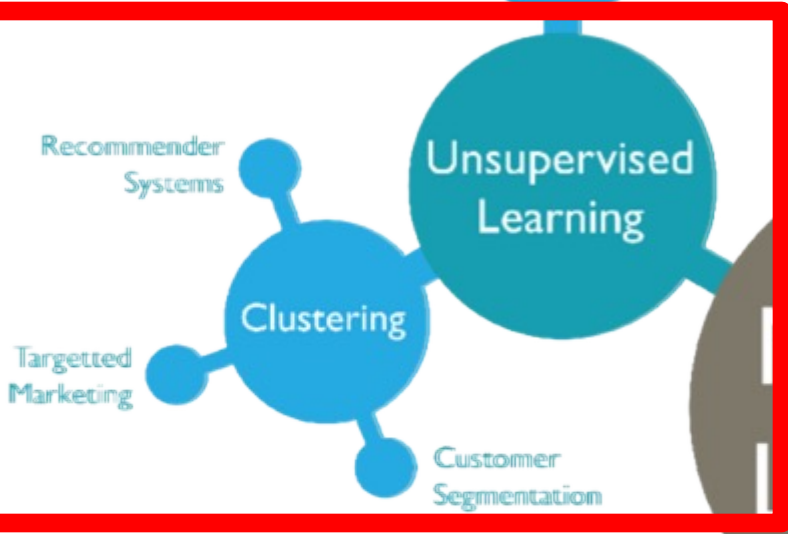
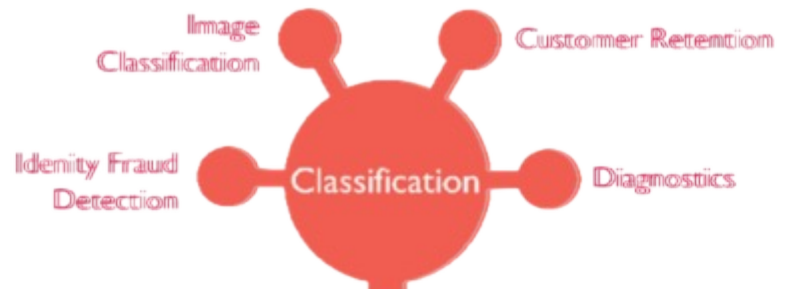


Unsupervised Learning: Clustering

Beyond K-means



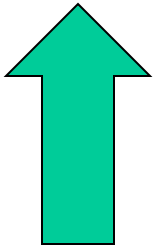
Machine Learning



(2) Hierarchical clustering

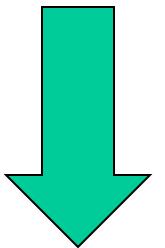
- **Agglomerative**

- **Bottom-up** approach: elements start as individual clusters & clusters are merged as one moves up the hierarchy



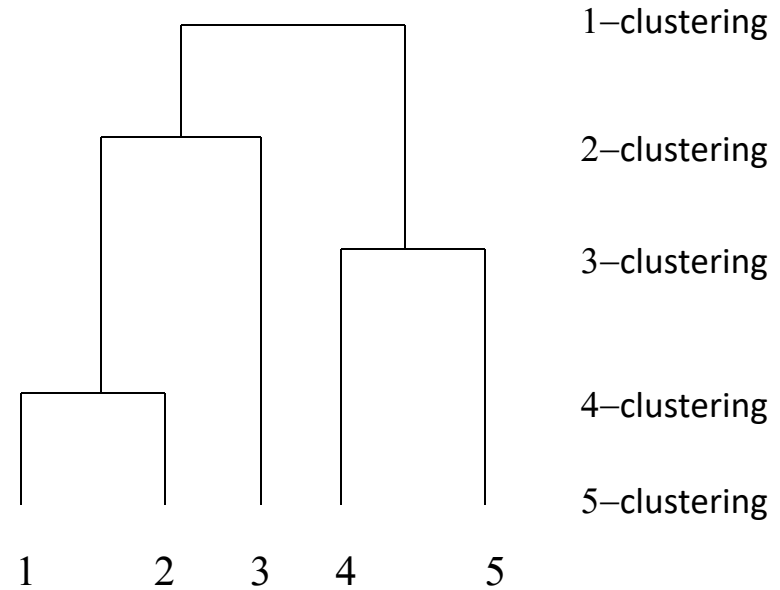
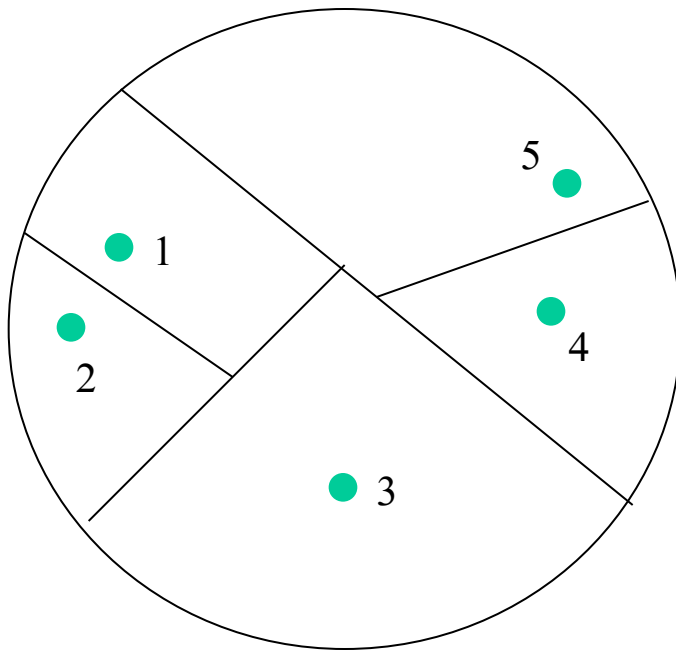
- **Divisive**

- **Top-down** approach: elements start as a single cluster & clusters are split as one moves down the hierarchy



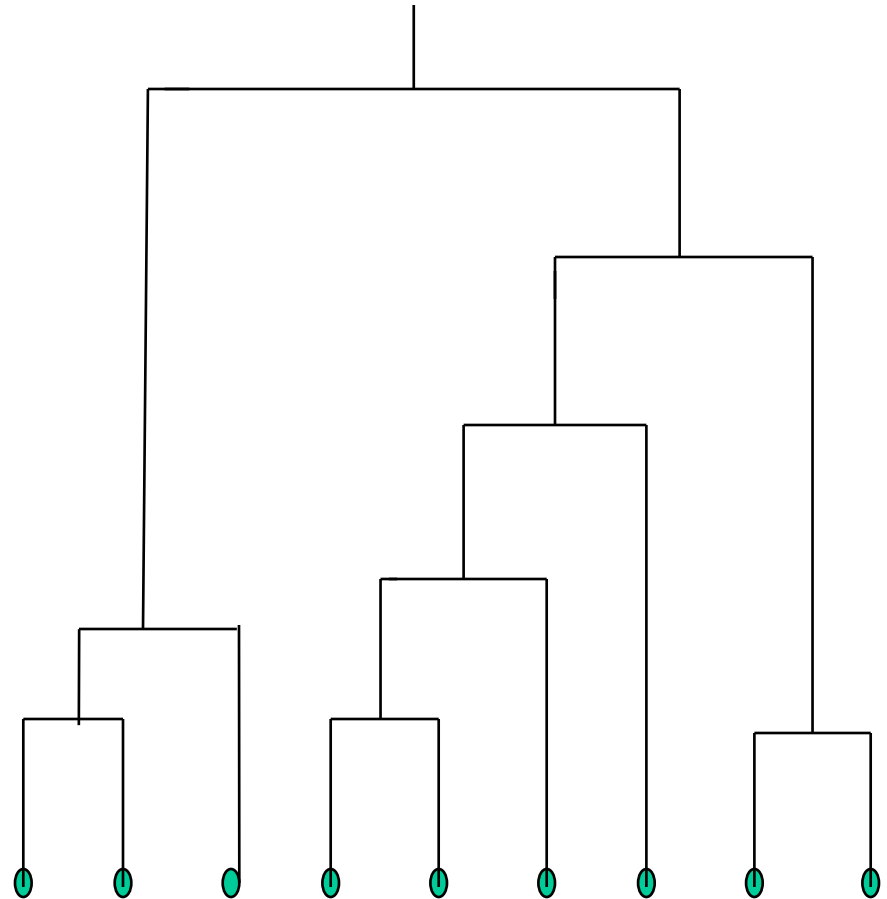
Hierarchical Clustering

Recursive partitioning/merging of a data set



Dendrogram

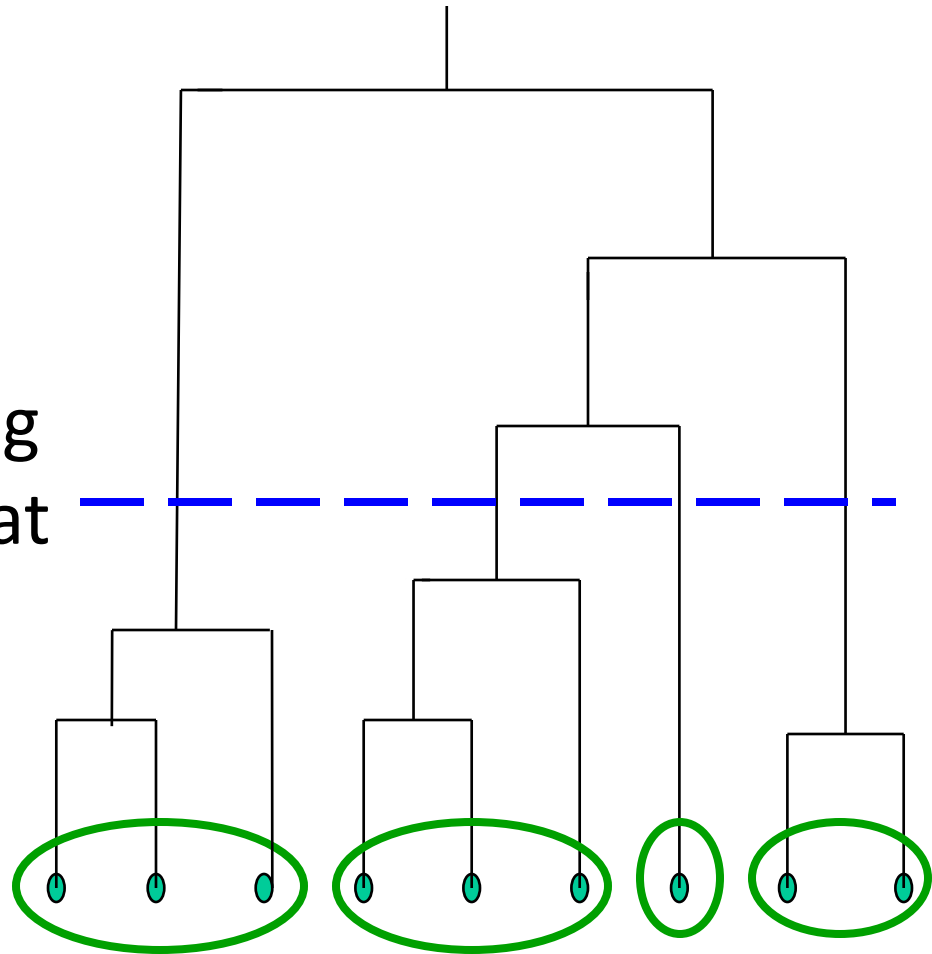
- Tree structure representing all data partitionings
- Constructed as clustering proceeds



Nine items

Dendrogram

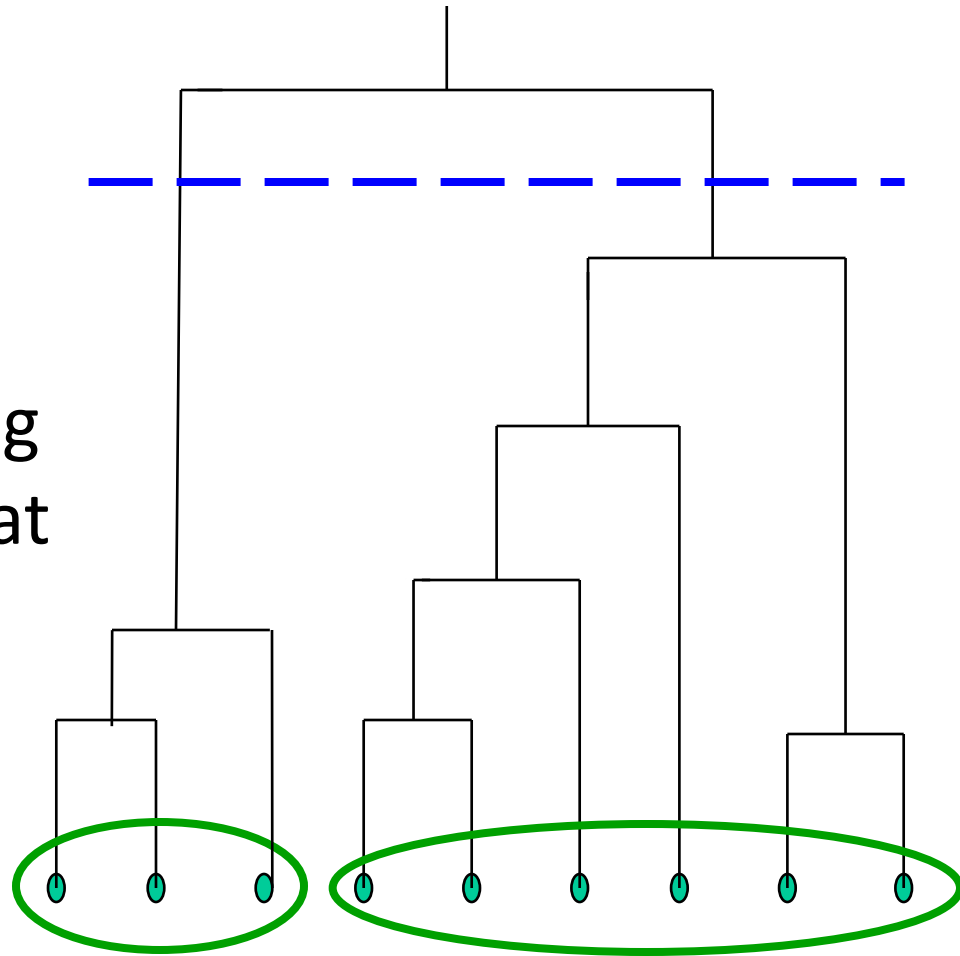
- Tree structure representing all data partitionings
- Constructed as clustering proceeds
- Get a K-clustering by looking at **connected** components at any given level
- Often binary dendograms, but n-ary ones easy to get with minor algorithm changes



Four clusters at this level

Dendrogram

- Tree structure representing all data partitionings
- Constructed as clustering proceeds
- Get a K-clustering by looking at **connected** components at any given level
- Often binary dendograms, but n-ary ones easy to get with minor algorithm changes

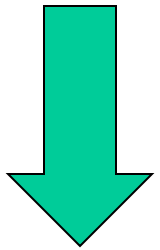


Two clusters at this level

Hierarchical clustering advantages

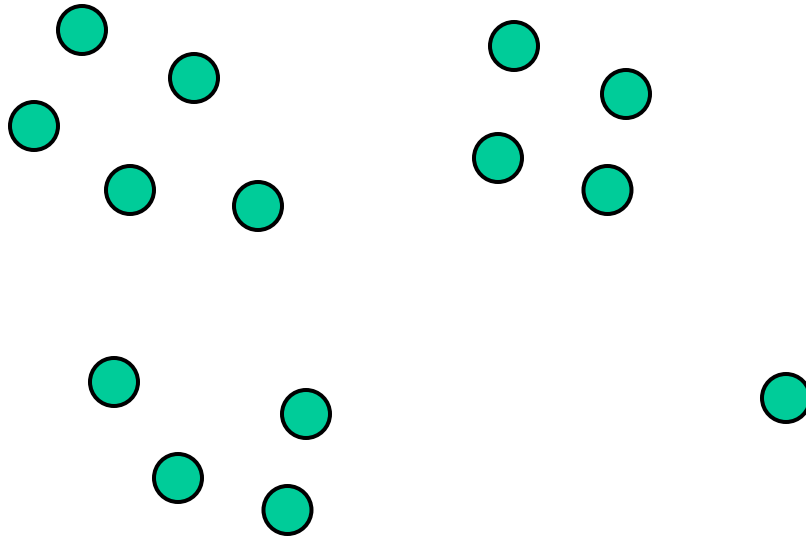
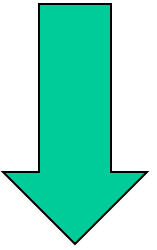
- Need not specify number of clusters
- Good for data visualization
 - See how data points interact at many levels
 - Can view data at multiple granularity levels
 - Understand how all points interact
- Specifies all of the K clusterings/partitions

Divisive hierarchical clustering

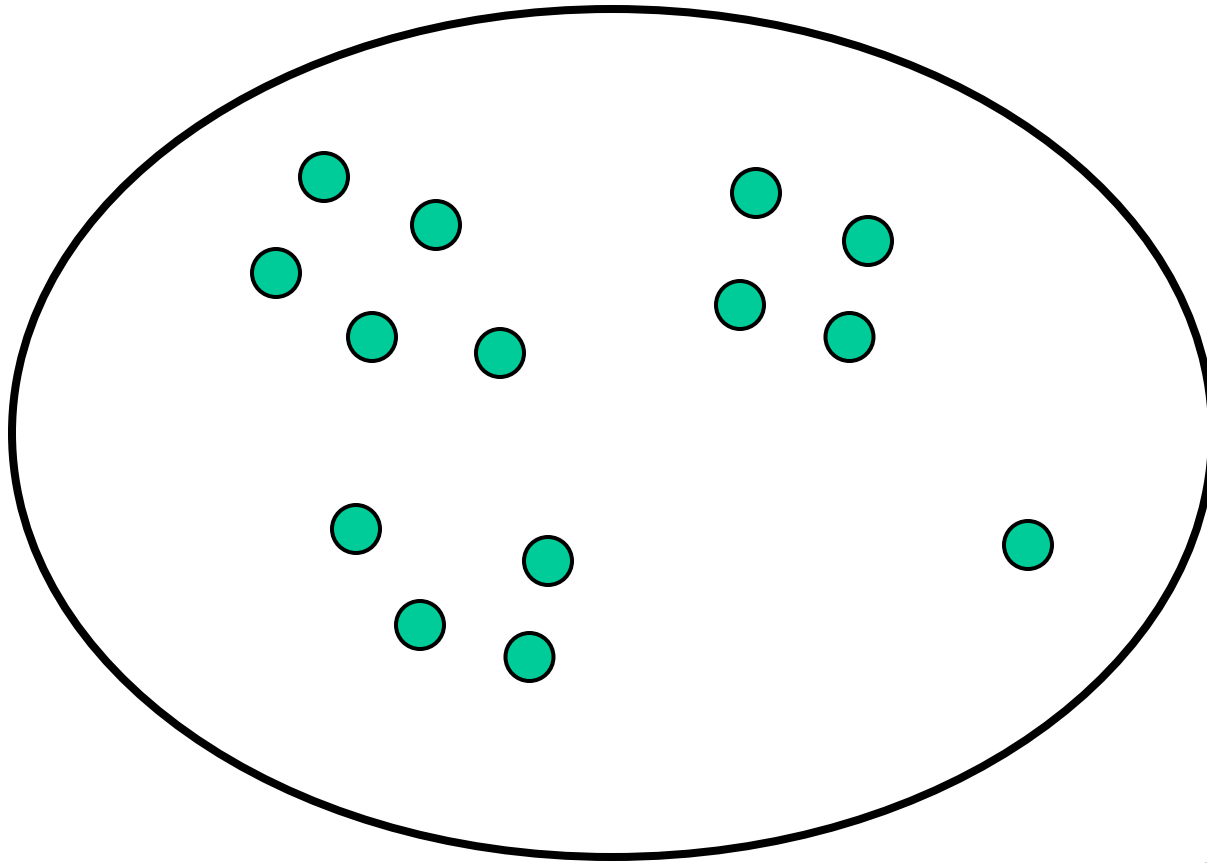
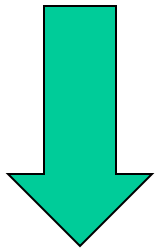


- Top-down technique to find best partitioning of data, generally exponential in time
- Common approach:
 - Let \mathbf{C} be a set of clusters
 - Initialize \mathbf{C} to be a one-clustering of data
 - While there exists a cluster c in \mathbf{C}
 - remove c from \mathbf{C}
 - partition c into 2 clusters (c_1 and c_2) using a flat clustering algorithm (e.g., k-means with $k=2$)
 - Add to c_1 and c_2 \mathbf{C}

Divisive clustering

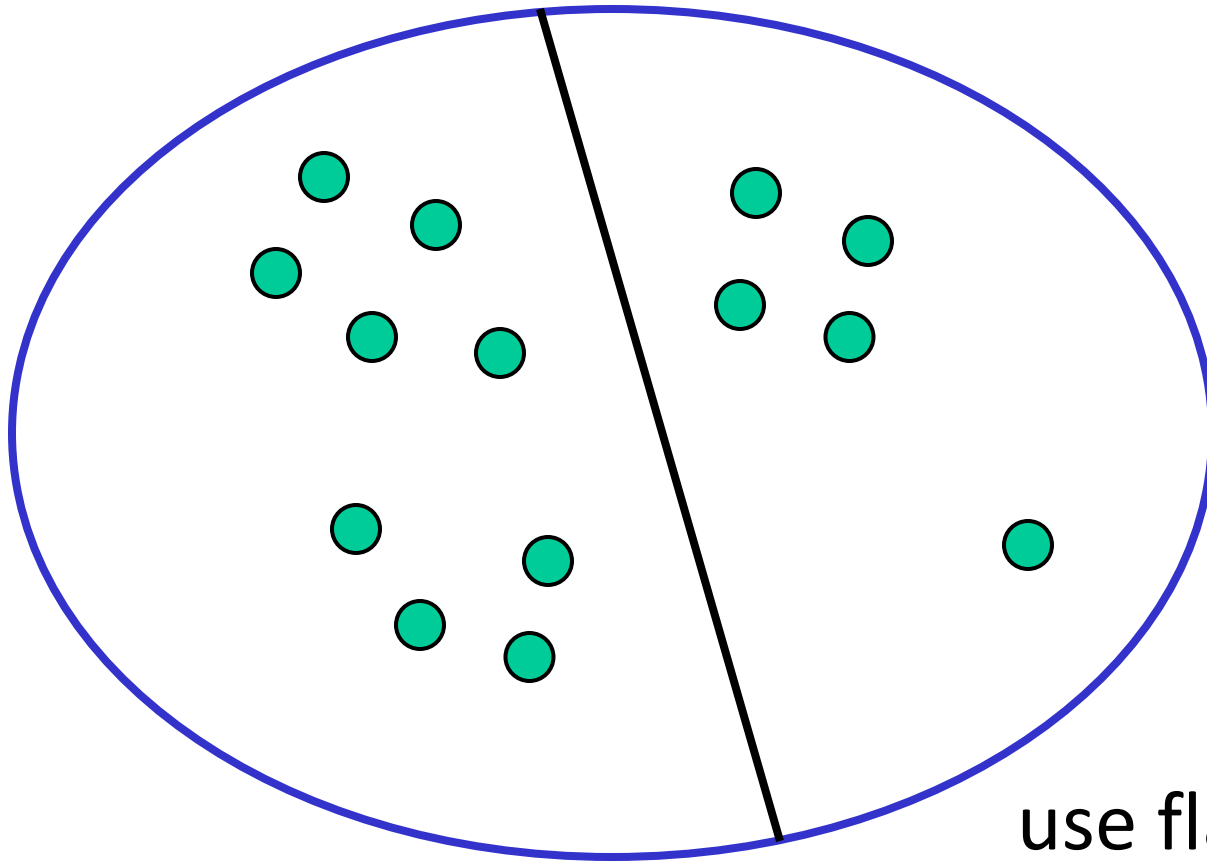
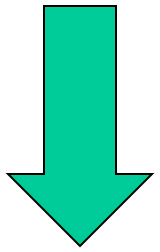


Divisive clustering



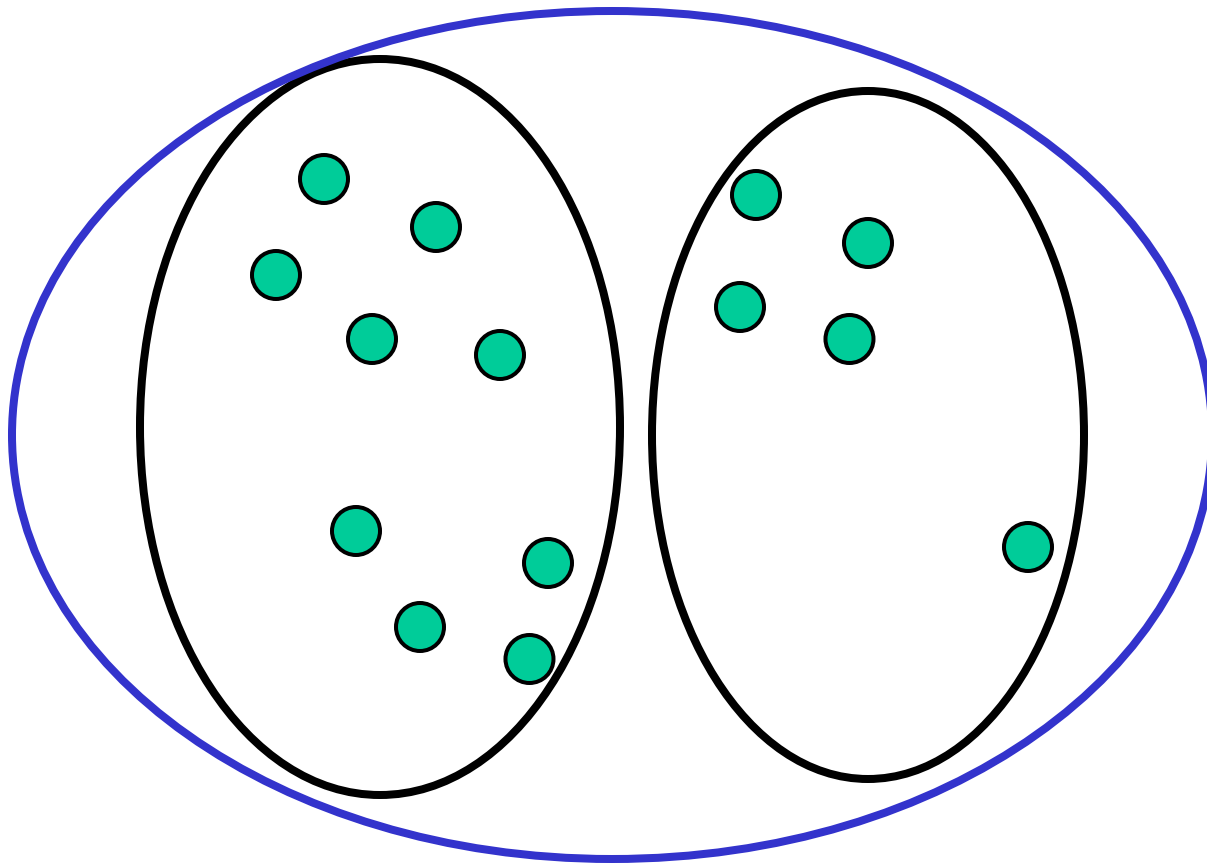
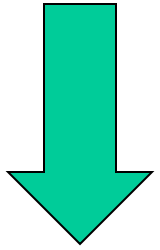
start with one
cluster

Divisive clustering

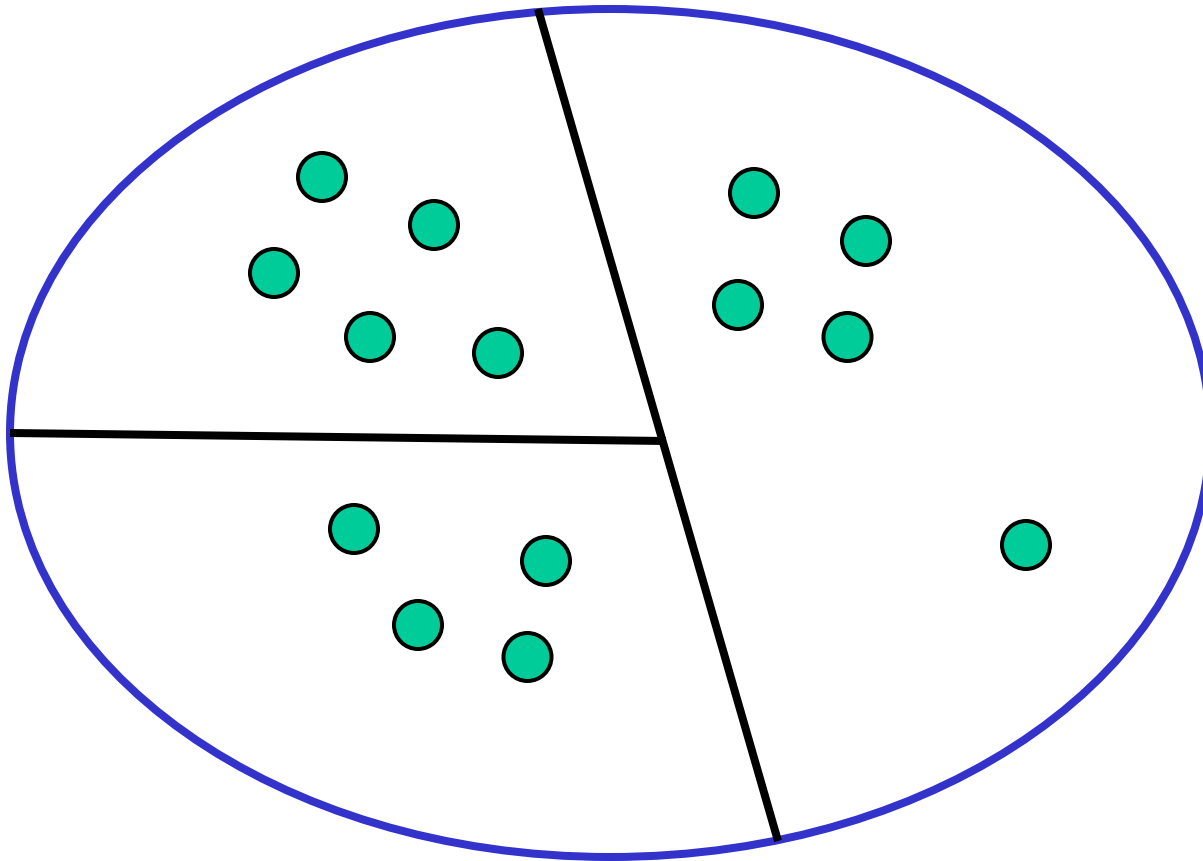
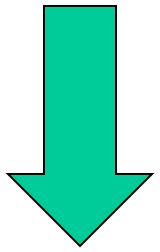


use flat clustering to
split into two clusters (e.g.,
using K-means with $k=2$)

Divisive clustering

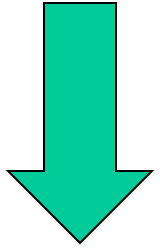


Divisive clustering

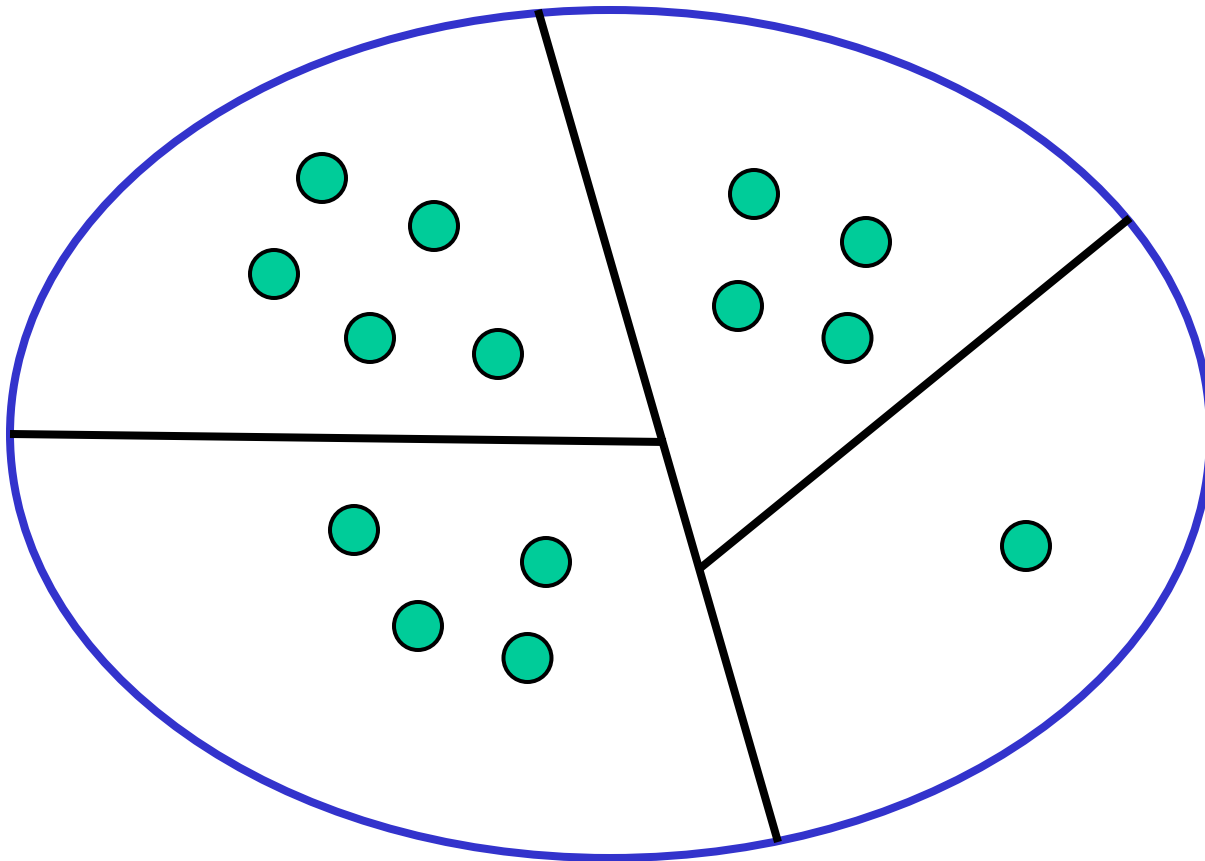


split using flat
clustering,
e.g., K-means

Divisive clustering

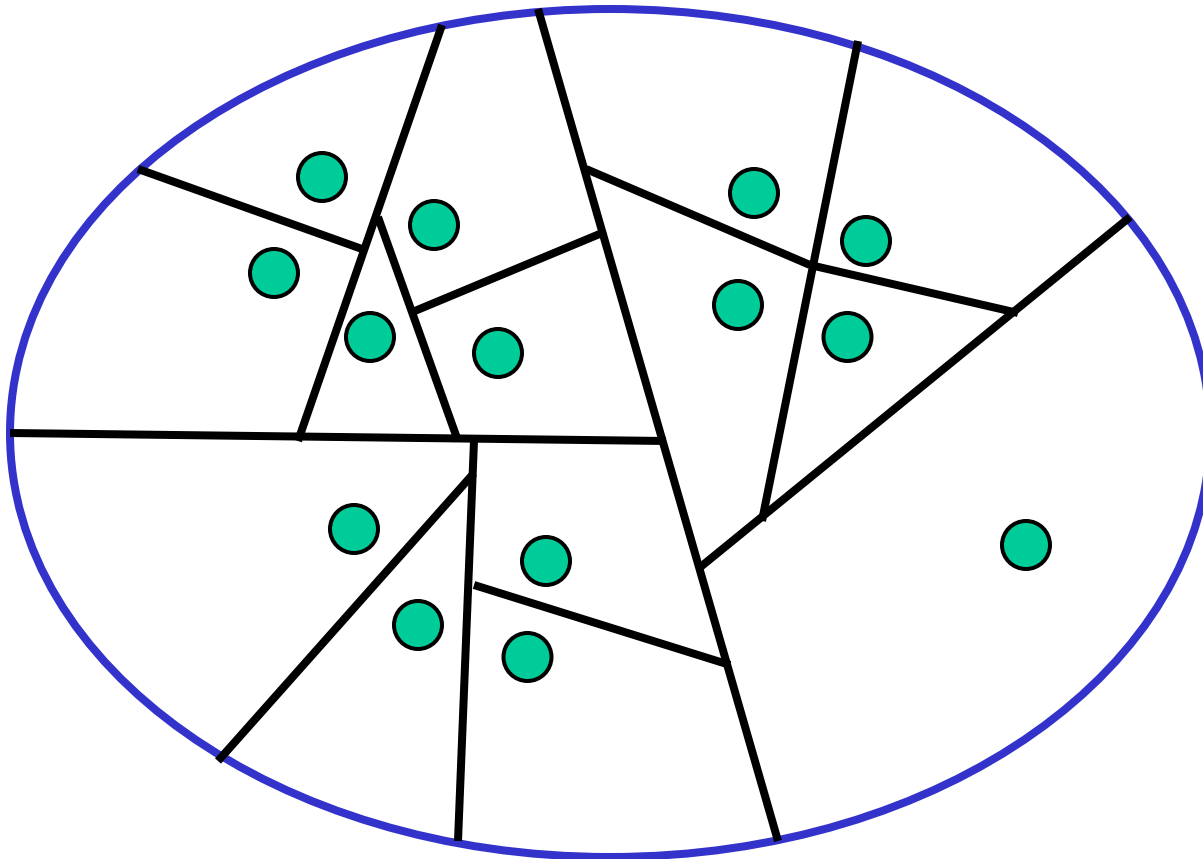
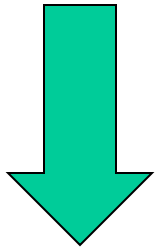


split using flat clustering



split using flat
clustering,
e.g., K-means

Divisive clustering



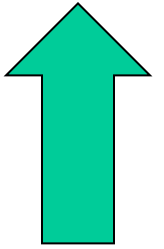
Stop when clusters reach some constraint,
e.g., all of size 1

AGGLOMERATIVE CLUSTERING

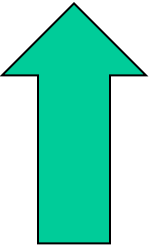
All observations start as their own cluster. Clusters meeting some criteria are merged. This process is repeated, growing clusters until some end point is reached.

ChrisAlbon

Hierarchical Agglomerative Clustering

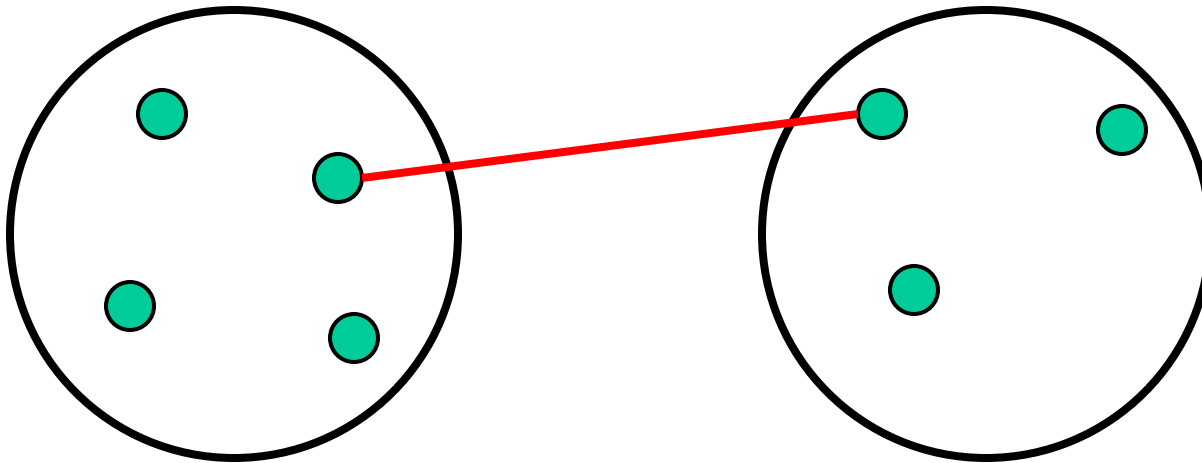


- Let \mathbf{C} be a set of clusters
- Initialize \mathbf{C} to all points/docs as separate clusters
- While \mathbf{C} contains more than one cluster
 - find c_1 and c_2 in \mathbf{C} that are **closest together**
 - remove c_1 and c_2 from \mathbf{C}
 - merge c_1 and c_2 and add resulting cluster to \mathbf{C}
- Merging history forms a binary tree or hierarchy
- **Q: How to measure distance between clusters?**



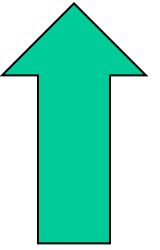
Distance between clusters

Single-link: Similarity of the *most* similar (single-link)



$$\max_{l \in L, r \in R} \text{sim}(l, r)$$

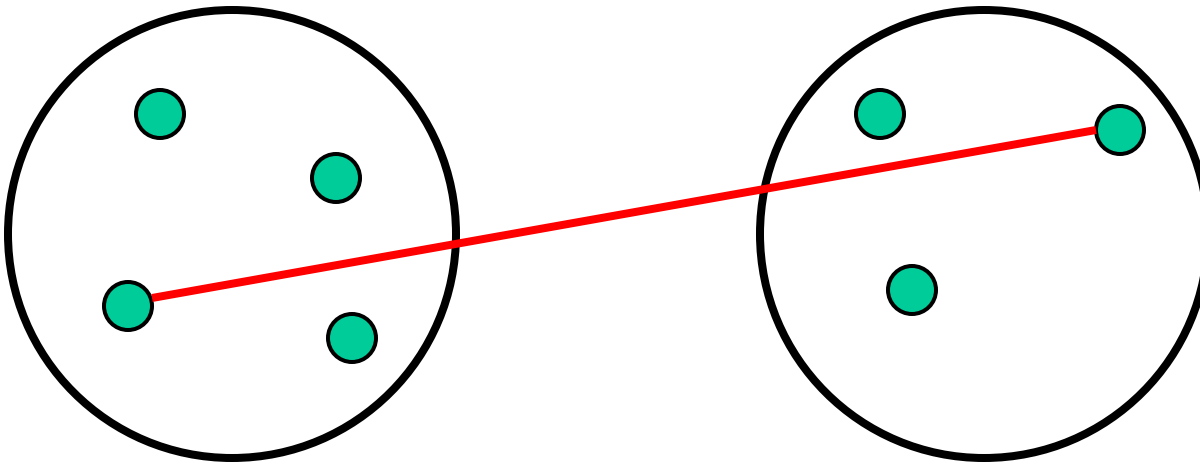
Weka: linkType=SINGLE



Distance between clusters

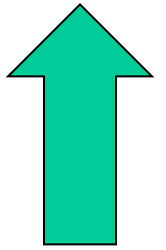
Complete-link: Similarity of the “furthest” points, the *least* similar

$$\min_{l \in L, r \in R} sim(l, r)$$

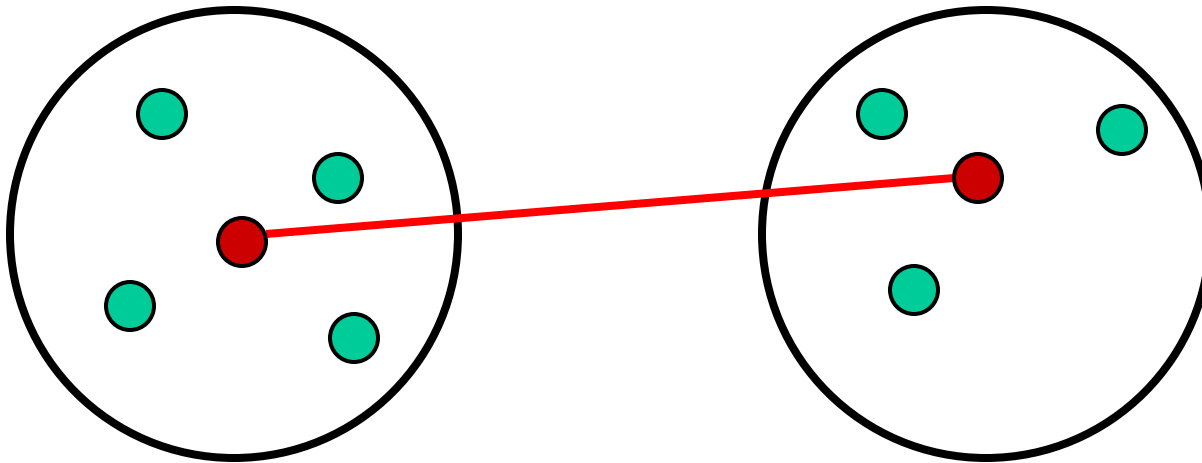


Weka: linkType=COMPLETE

Distance between clusters



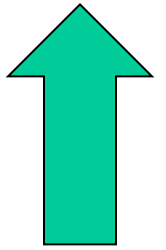
Centroid: Clusters whose centroids (centers of gravity) are the most similar



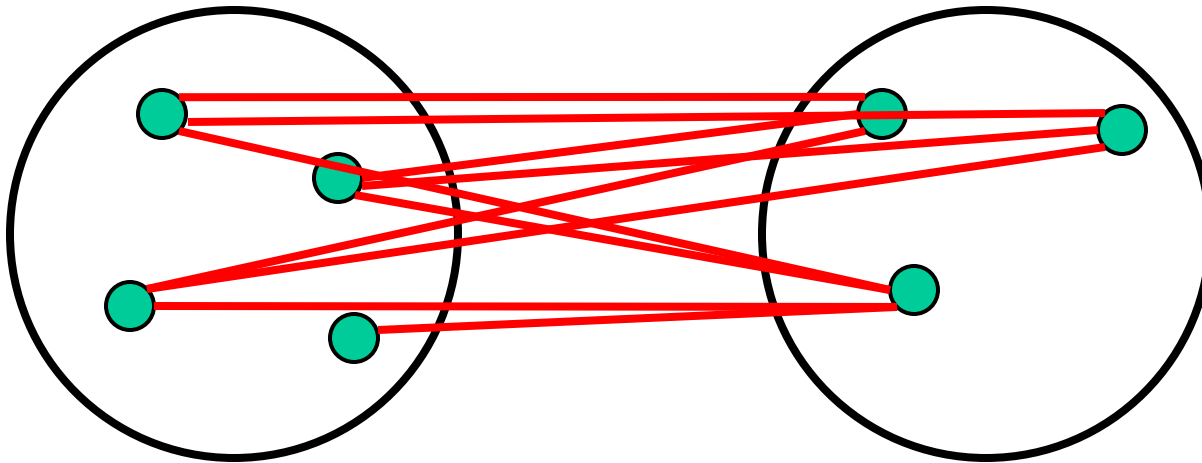
$$\|\mu(L) - \mu(R)\|^2$$

Weka: linkType=CENTROID

Distance between clusters



Average-link: Average similarity between all pairs of elements



$$\frac{1}{|L| \cdot |R|} \sum_{x \in L, y \in R} \|x - y\|^2$$

Weka: linkType=AVERAGE

Preprocess Classify Cluster Associate Select attributes Visualize

Clusterer

Choose HierarchicalClusterer -N 3 -L SINGLE -P -A "weka.core.EuclideanDistance -R first-last"

Cluster mode

- Use training set
 Supplied test set Set...
 Percentage split % 66
 Classes to clusters evaluation
 (Nom) class
 Store clusters for visualization

Ignore attributes

Start

Ignore attributes during clustering

Result list (right-click for options)

10:09:16 - HierarchicalClusterer

Clusterer output

```
Cluster 0
((((((((((((((((((((((0.2:0.03254,0.2:0.03254):0.00913,(0.3:0.03254,0.3:0.03254):0.00913):0.0
Cluster 2
((((((((((((((((((((((((((((((((((((((((((1.4:0.07344,(((1.5:0.06508,1.5:0.06508):0.00066,(1.4:0.05008,1
```

Time taken to build model (full training data) : 0.01 seconds

=== Model and evaluation on training set ===

Clustered Instances

```
0      49 ( 33%)
1       1 (  1%)
2     100 ( 67%)
```

Class attribute: class

Classes to Clusters:

```
  0  1  2  <-- assigned to cluster
49  1  0  | Iris-setosa
  0  0 50 | Iris-versicolor
  0  0 50 | Iris-virginica
```

Cluster 0 <-- Iris-setosa

Cluster 1 <-- No class

Cluster 2 <-- Iris-versicolor

Incorrectly clustered instances : 51 0 24 %

Default **SINGLE** cluster distance gives poor results here

Preprocess Classify Cluster Associate Select attributes Visualize

Clusterer

Choose HierarchicalClusterer -N 3 -L AVERAGE -P -A "weka.core.EuclideanDistance -R first-last"

Cluster mode

- Use training set
 Supplied test set
 Percentage split % 66
 Classes to clusters evaluation

 Store clusters for visualization

Ignore attributes

Start

Stop

Result list (right-click for options)

10:09:16 - HierarchicalClusterer
 10:09:58 - HierarchicalClusterer

Clusterer output

```
Cluster 1
((((((1.4:0.08775,(1.5:0.06508,1.5:0.06508):0.02267):0.04395,1.7:0.1317):0.01307,((1.5:0.0
Cluster 2
((((((2.5:0.12797,(2.3:0.10565,(2.4:0.06047,2.3:0.06047):0.04518):0.02232):0.06295,(((2.1:0.
```

Time taken to build model (full training data) : 0.01 seconds

=== Model and evaluation on training set ===

Clustered Instances

```
0      50 ( 33%)
1      67 ( 45%)
2      33 ( 22%)
```

Class attribute: class

Classes to Clusters:

```
0 1 2 <-- assigned to cluster
50 0 0 | Iris-setosa
0 50 0 | Iris-versicolor
0 17 33 | Iris-virginica
```

```
Cluster 0 <-- Iris-setosa
Cluster 1 <-- Iris-versicolor
Cluster 2 <-- Iris-virginica
```

Incorrectly clustered instances : 17 0 11 3333 %

Using **AVERAGE** cluster distance measure improves results

Knowing when to stop



- General issue is knowing when to stop merging/splitting a cluster
- We may have a problem specific desired range of clusters (e.g., 3-6)
- There are some general metrics for assessing quality of a cluster
- There are also domain specific heuristics for cluster quality

(3) DBSCAN Algorithm

- Density-Based Spatial Clustering of Applications with Noise
- It clusters close points based on a distance and a minimum number of points
 - Key parameters: ϵ =maximum distance between two points; minPoints= minimal cluster size
- Marks as outliers points in low-density regions
- Needn't specify number of clusters expected
- Fast

DBSCAN

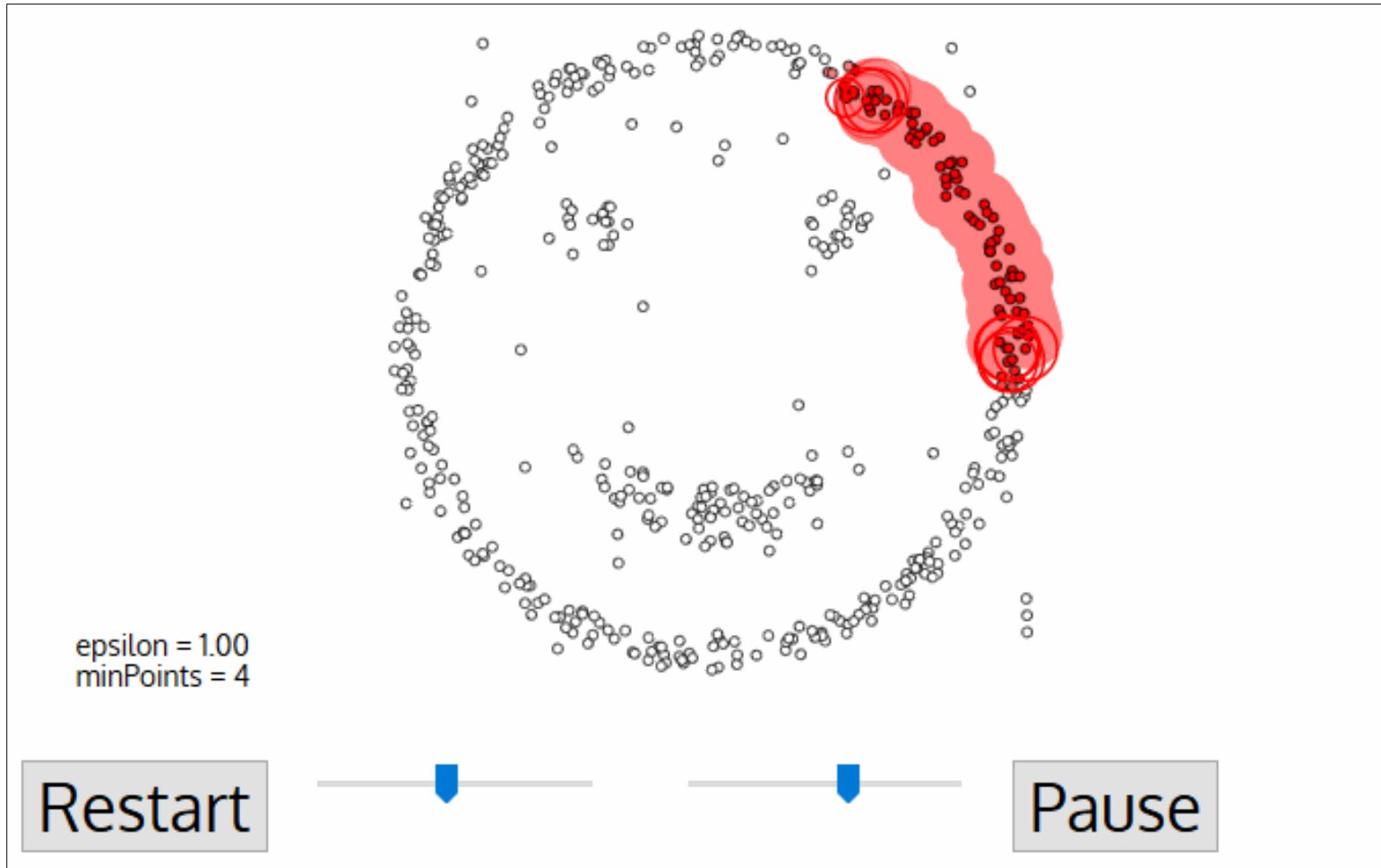
DBSCAN looks for densely packed observations and makes no assumptions about the number or shape of clusters.

1. A random observation, x_i , is selected
2. If x_i has a minimum of close neighbors, we consider it part of a cluster.
3. Step 2 is repeated recursively for all of x_i 's neighbors, then neighbors' neighbors etc... These are the cluster's core members.
4. Once Step 3 runs out of observations, a new random point is chosen

Afterwards, observations not part of a core are assigned to a nearby cluster or marked as outliers.

ChrisAlbon

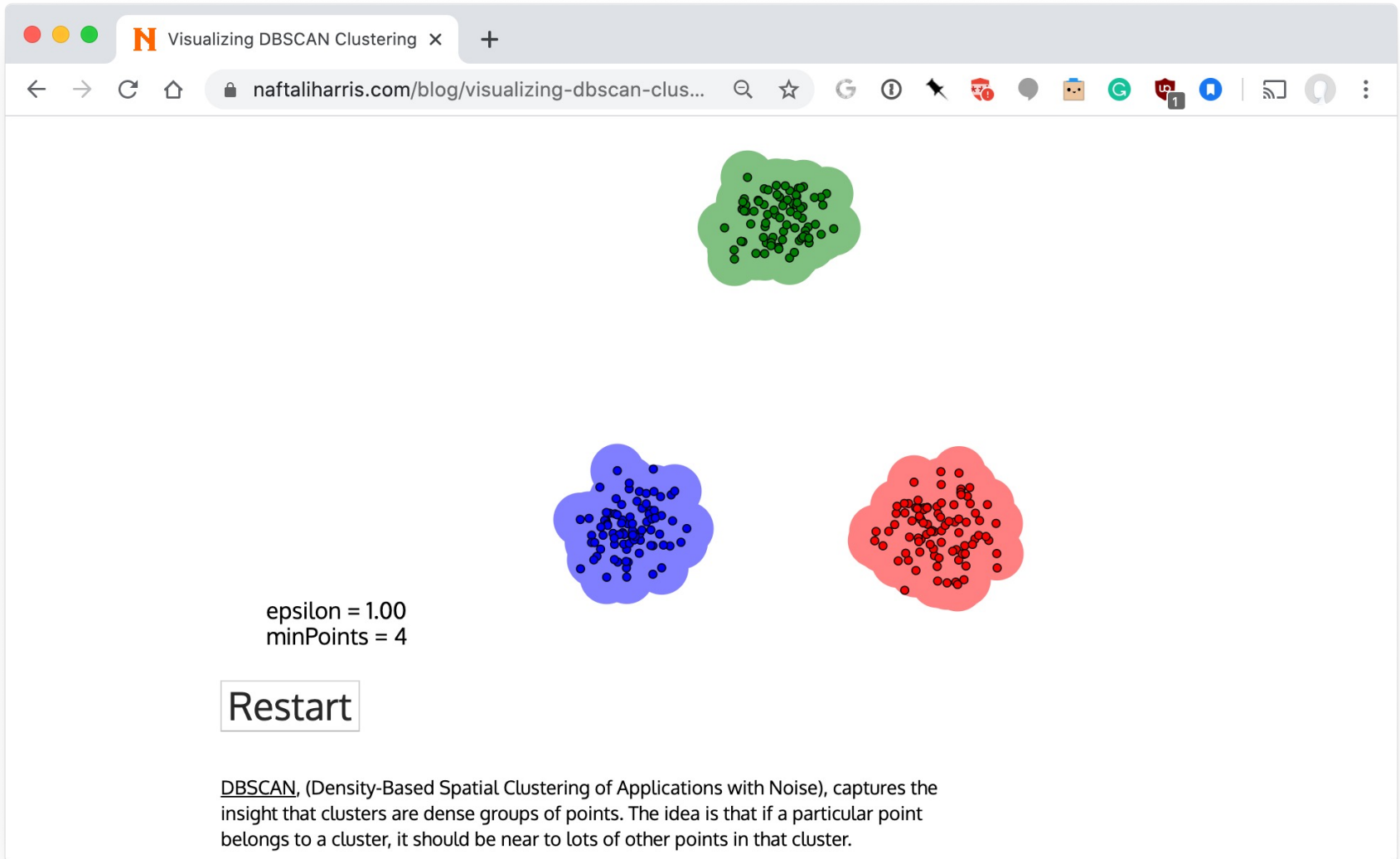
DBSCAN Example



This gif (in ppt) shows how DBSCAN grows four clusters and identifies the remaining points as outliers

Visualizing DBSCAN

<https://bit.ly/471dbscan>



Visualizing DBSCAN Clustering

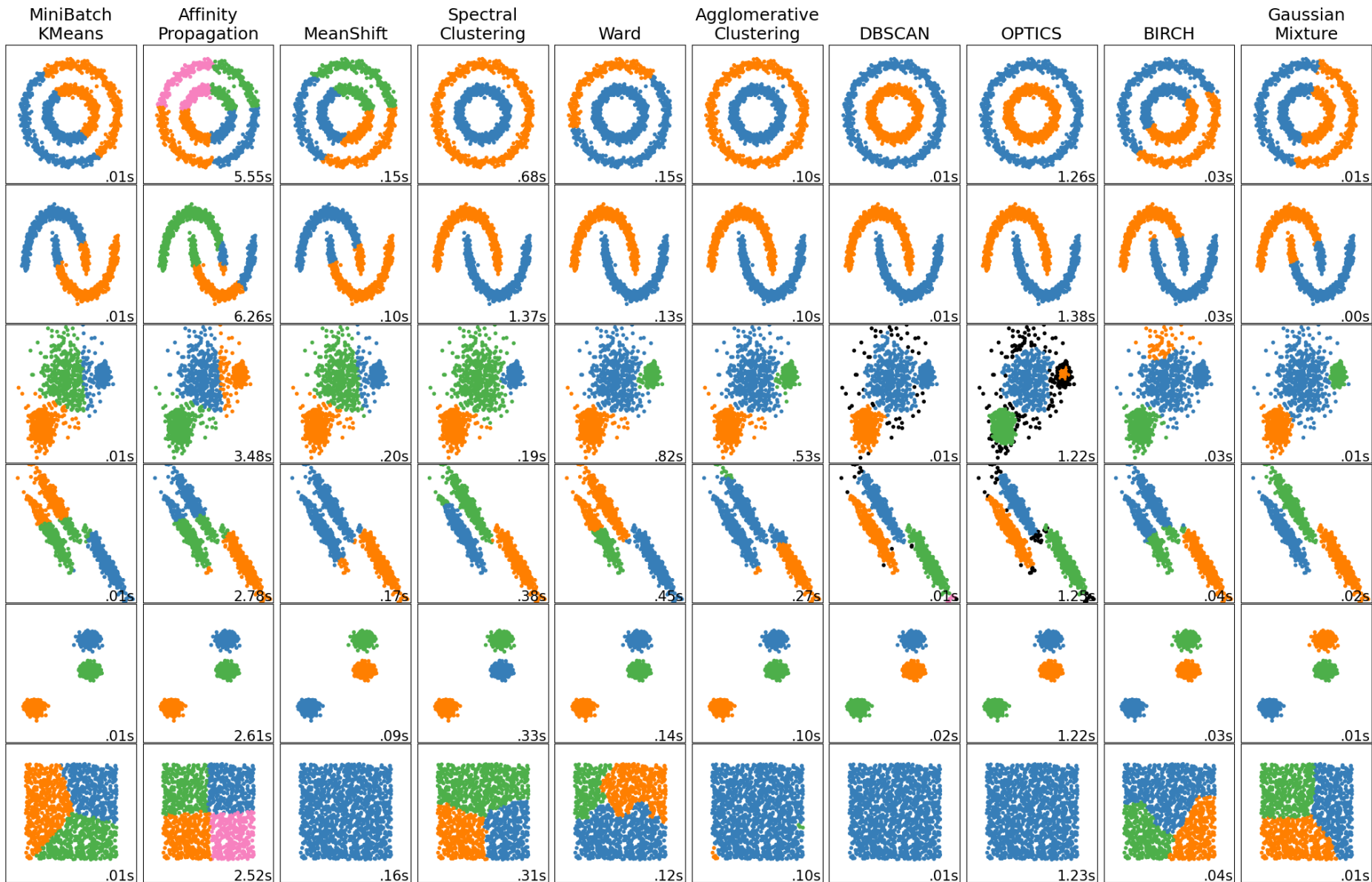
naftaliharris.com/blog/visualizing-dbscan-clus...

epsilon = 1.00
minPoints = 4

Restart

DBSCAN, (Density-Based Spatial Clustering of Applications with Noise), captures the insight that clusters are dense groups of points. The idea is that if a particular point belongs to a cluster, it should be near to lots of other points in that cluster.

Comparing clustering algorithms via Scikit Learn



Clustering Summary

- Clustering useful & effective for many tasks
- K-means clustering one of simplest & fastest techniques, but
 - Requires knowing how many clusters is right
 - Doesn't handle outliers well
- Hierarchical clustering slower and more general, but needs a metric to know when to stop
- There are many other clustering options