

A dendrogram shows the mergers which have been made at each successive level.

Ex: $n=4$

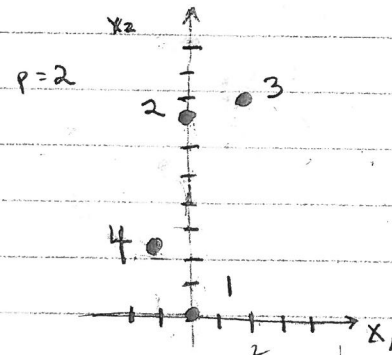
(i) Single Linkage

$$D = \{d_{ik}\} = \begin{matrix} & \begin{matrix} 1 & 2 & 3 & 4 \end{matrix} \\ \begin{matrix} 1 \\ 2 \\ 3 \\ 4 \end{matrix} & \begin{bmatrix} 0 & & & \\ 7 & 0 & & \\ 8 & 2 & 0 & \\ 3 & 5 & 6 & 0 \end{bmatrix} \end{matrix}$$

↑ Just use approximate values here for simplicity.

$$\min_{i \neq k} d_{ik} = 2$$

Objects 2 and 3 form the cluster (23).



Cluster Distances

$$d_{(23)1} = \min \{d_{21}, d_{31}\} = 7$$

$$d_{(23)4} = \min \{d_{24}, d_{34}\} = 5$$

Delete the row and column for objects 2 and 3 and add a row and column for cluster (23).

$$\begin{matrix} & \begin{matrix} (23) & 1 & 4 \end{matrix} \\ \begin{matrix} (23) \\ 1 \\ 4 \end{matrix} & \begin{bmatrix} 0 & & \\ 7 & 0 & \\ 5 & 3 & 0 \end{bmatrix} \end{matrix}$$

min distance in above matrix = 3

Merge objects 1 and 4 to form the cluster (14).

```

> TX <- c(0, 0,
+         0, 7,
+         2, 8,
+        -1, 2.5)
>
> X <- matrix(data=TX, ncol=2, byrow=TRUE)
>
> library(ecodist) # assumes package ecodist is installed
>
> # Distances
> de <- distance(X, method = "euclidean")
> dm <- distance(X, method = "mahalanobis")
> d <- de
> d
      1      2      3
2 7.000000
3 8.246211 2.236068
4 2.692582 4.609772 6.264982
>
> # Perform cluster analysis
> fit <- hclust(d, method="single")
>
> # Display dendrogram
> plot(fit)
> groups <- cutree(fit, k=2)
> groups
[1] 1 2 2 1
> #rect.hclust(fit,k=2,border="red")
> #rect.hclust(fit,h=3,border="blue")

```

