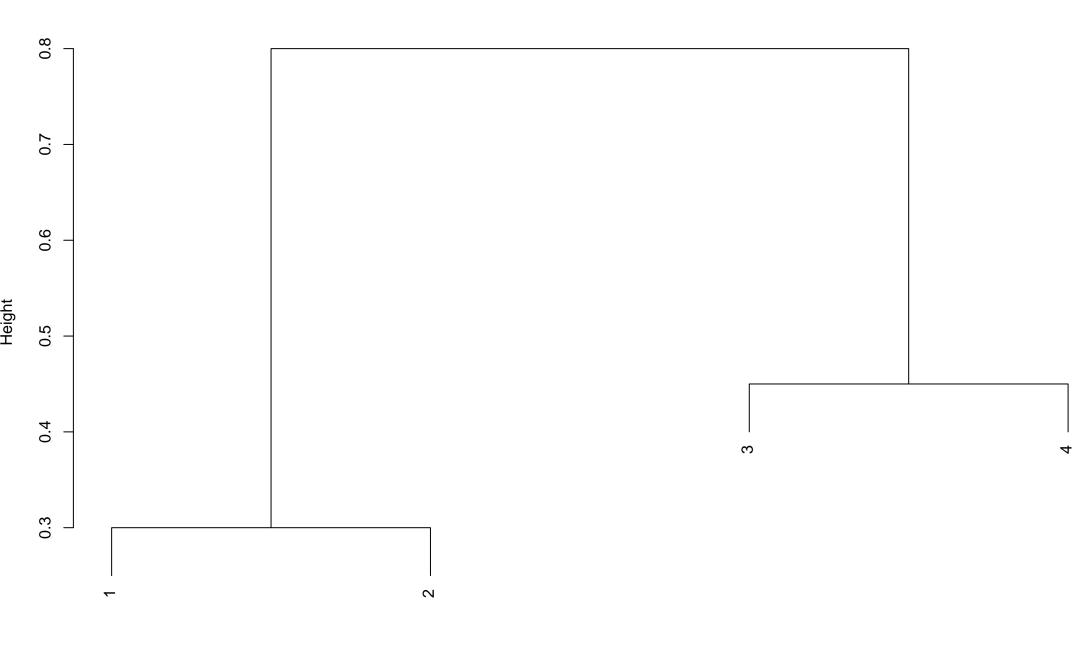
Cluster Dendrogram



as.dist(dissmilarity) hclust (*, "complete")