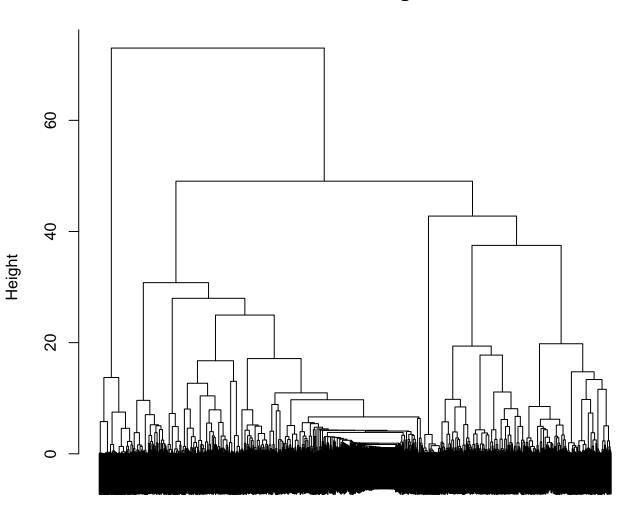
Cluster Dendrogram



as.dist(1 - cors.numeric) hclust (*, "ward.D")