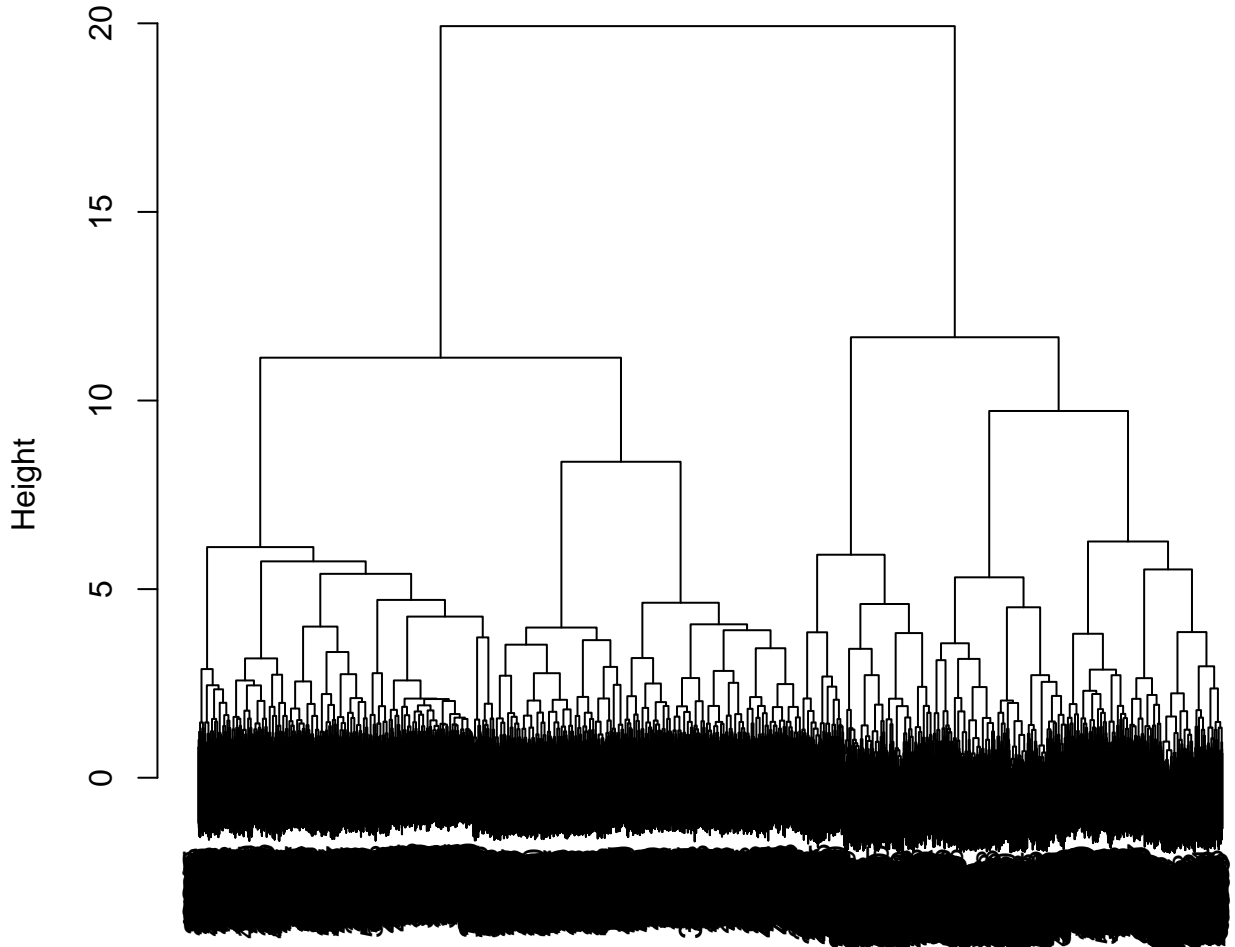


Cluster Dendrogram



as.dist(1 - modelo\$proximity)
hclust (*, "ward.D2")