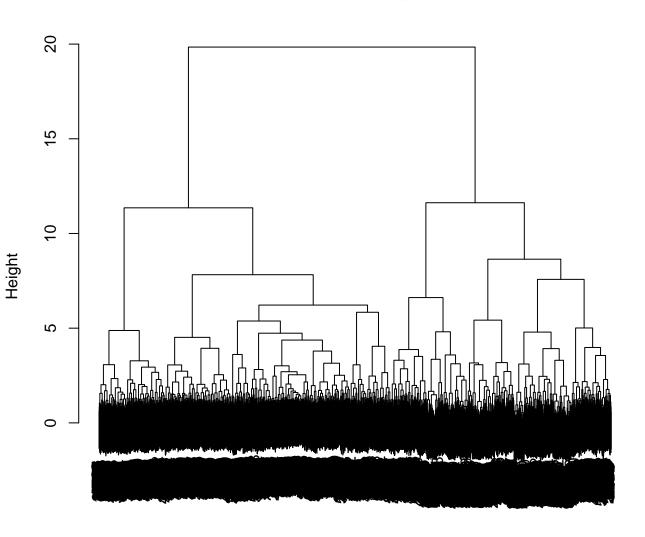
Cluster Dendrogram



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