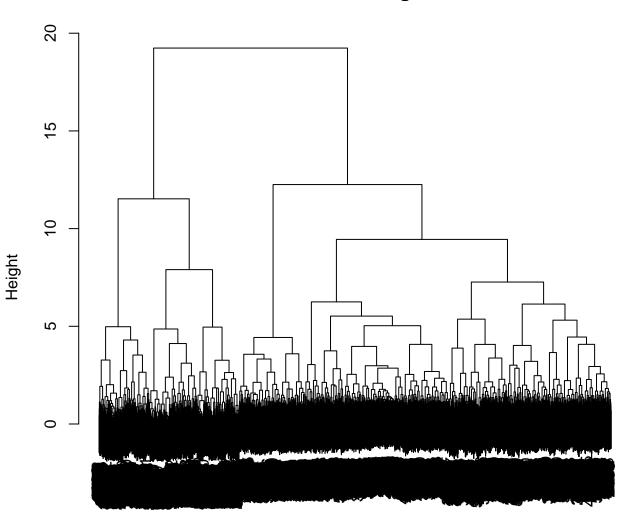
## **Cluster Dendrogram**



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