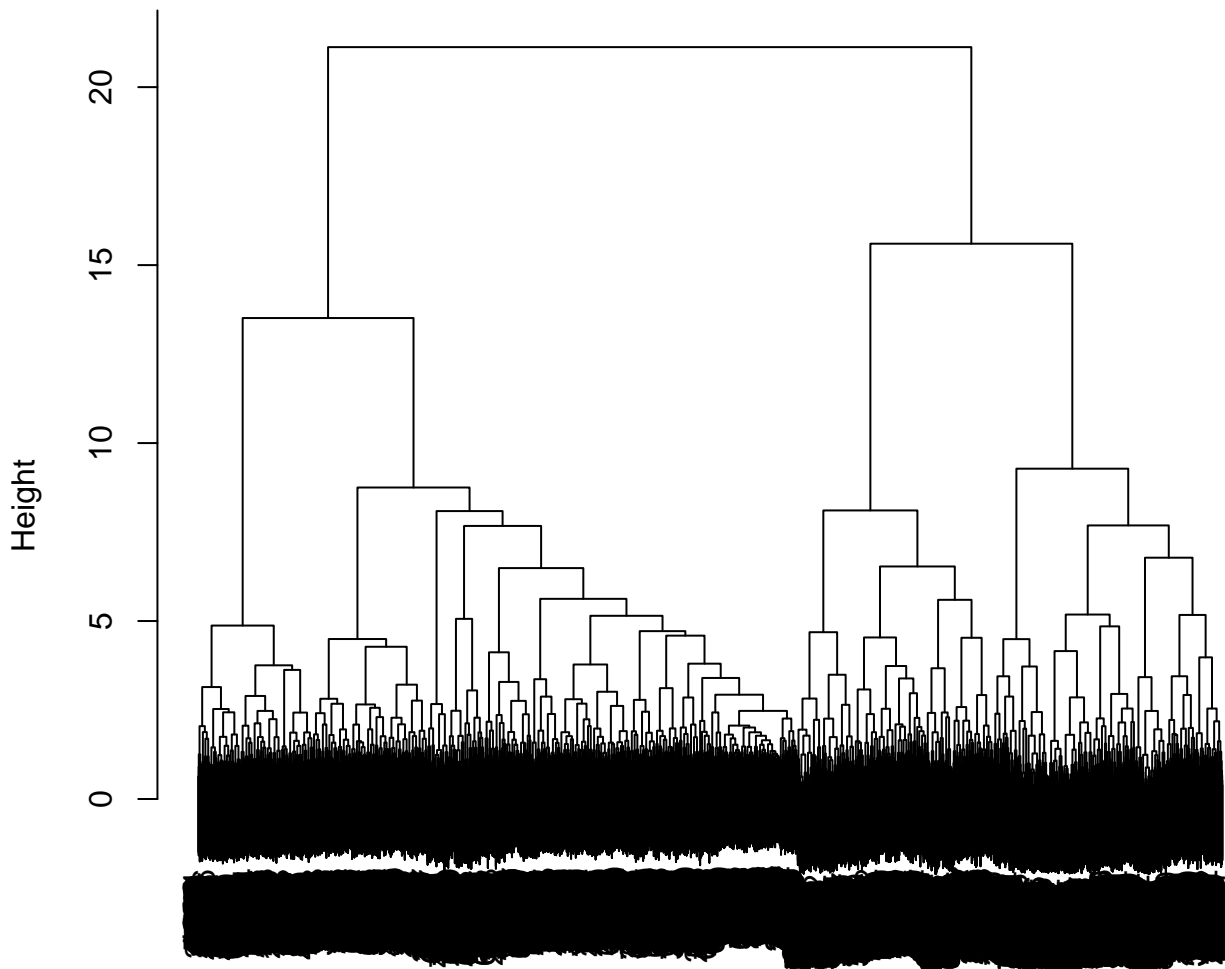


# Cluster Dendrogram



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