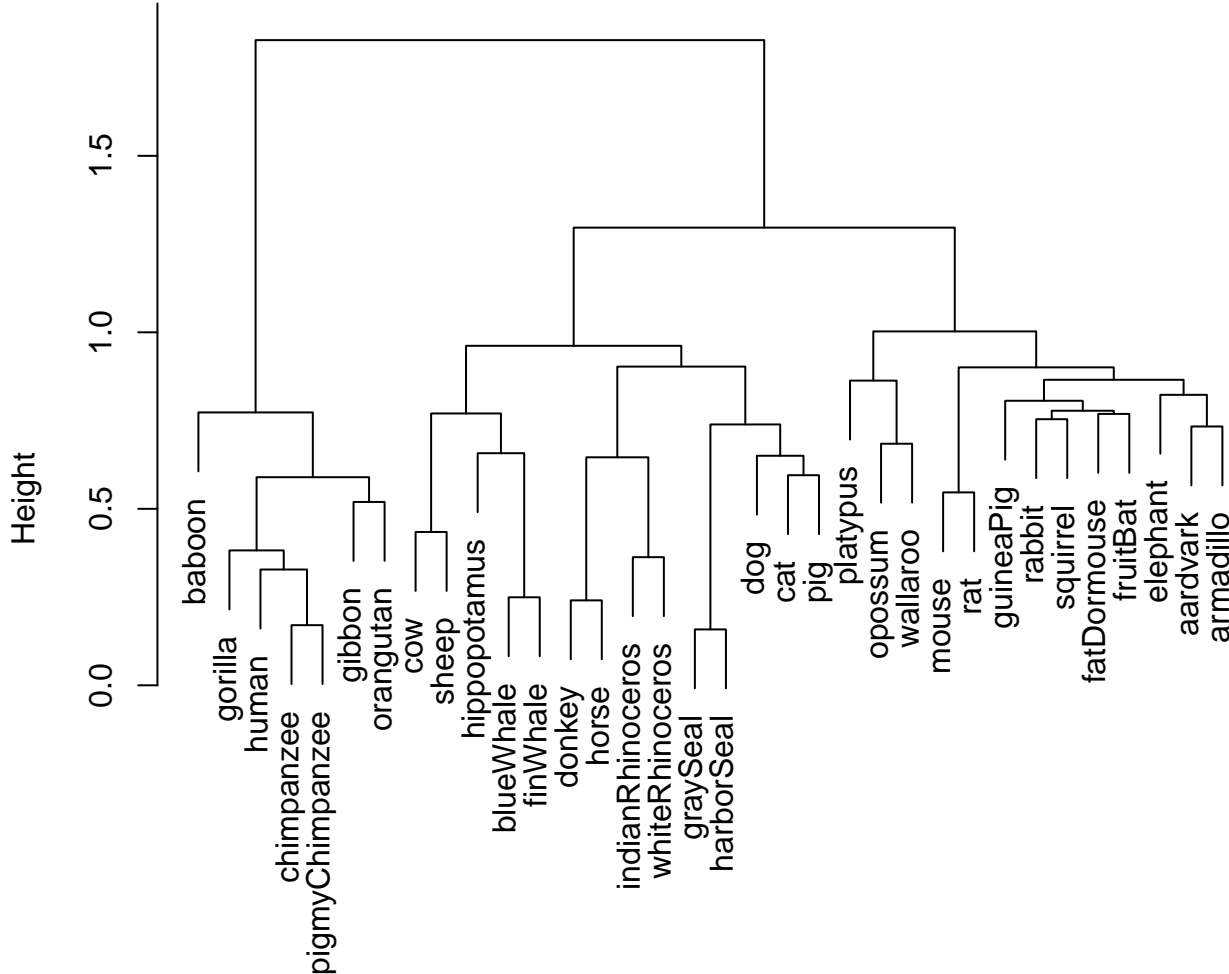


# Cluster Dendrogram



as.dist(distance\_matrix)  
hclust (\*, "ward.D2")