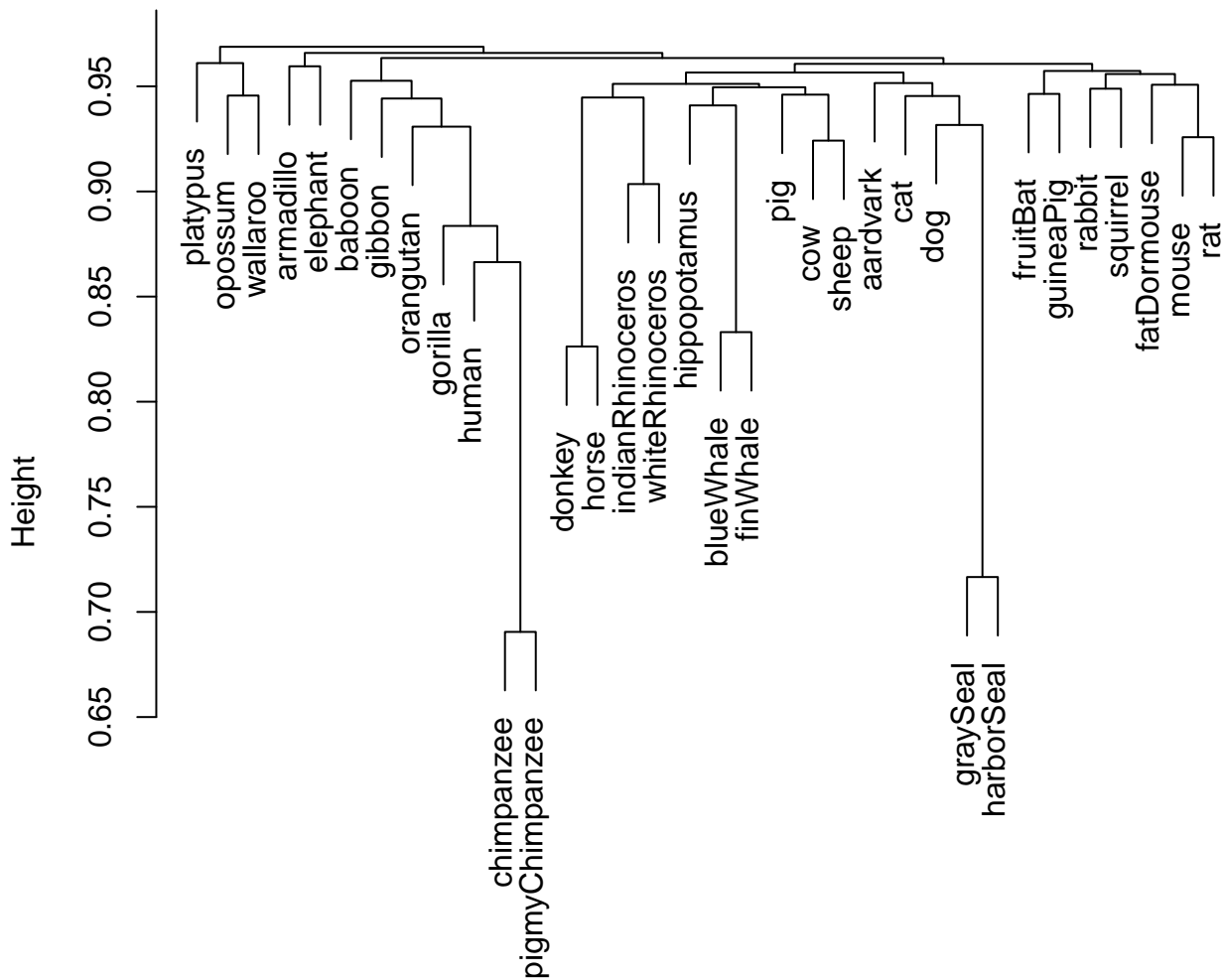


Cluster Dendrogram



as.dist(distance_matrix)
hclust (*, "complete")