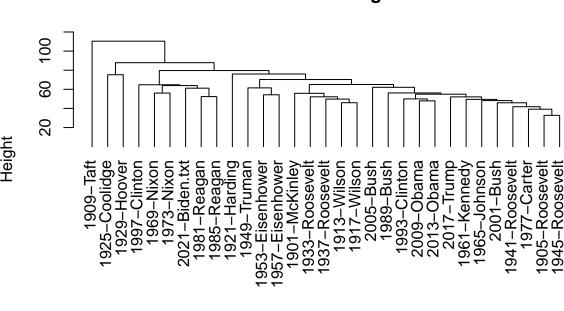
Cluster Dendrogram



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