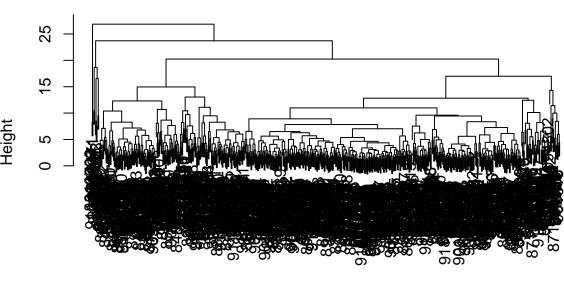# Cluster Dendrogram

data.dist
hclust (*, "complete")