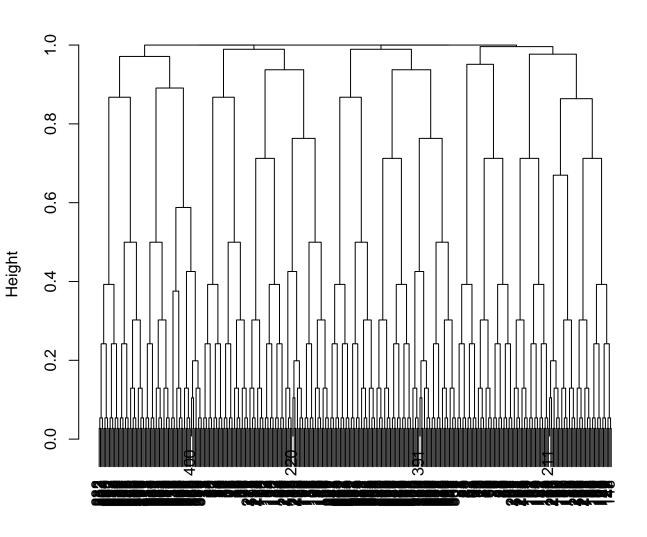
## **Cluster Dendrogram**



as.dist(1 – Sigma) hclust (\*, "complete")