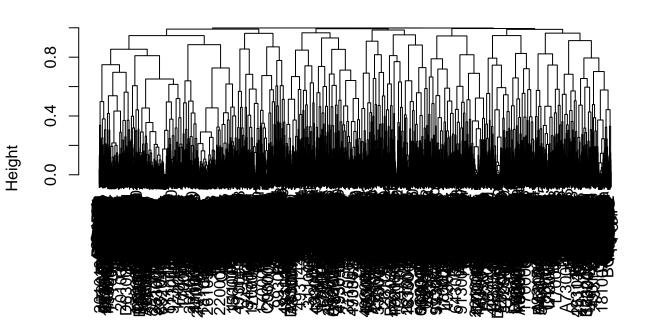
Cluster Dendrogram



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