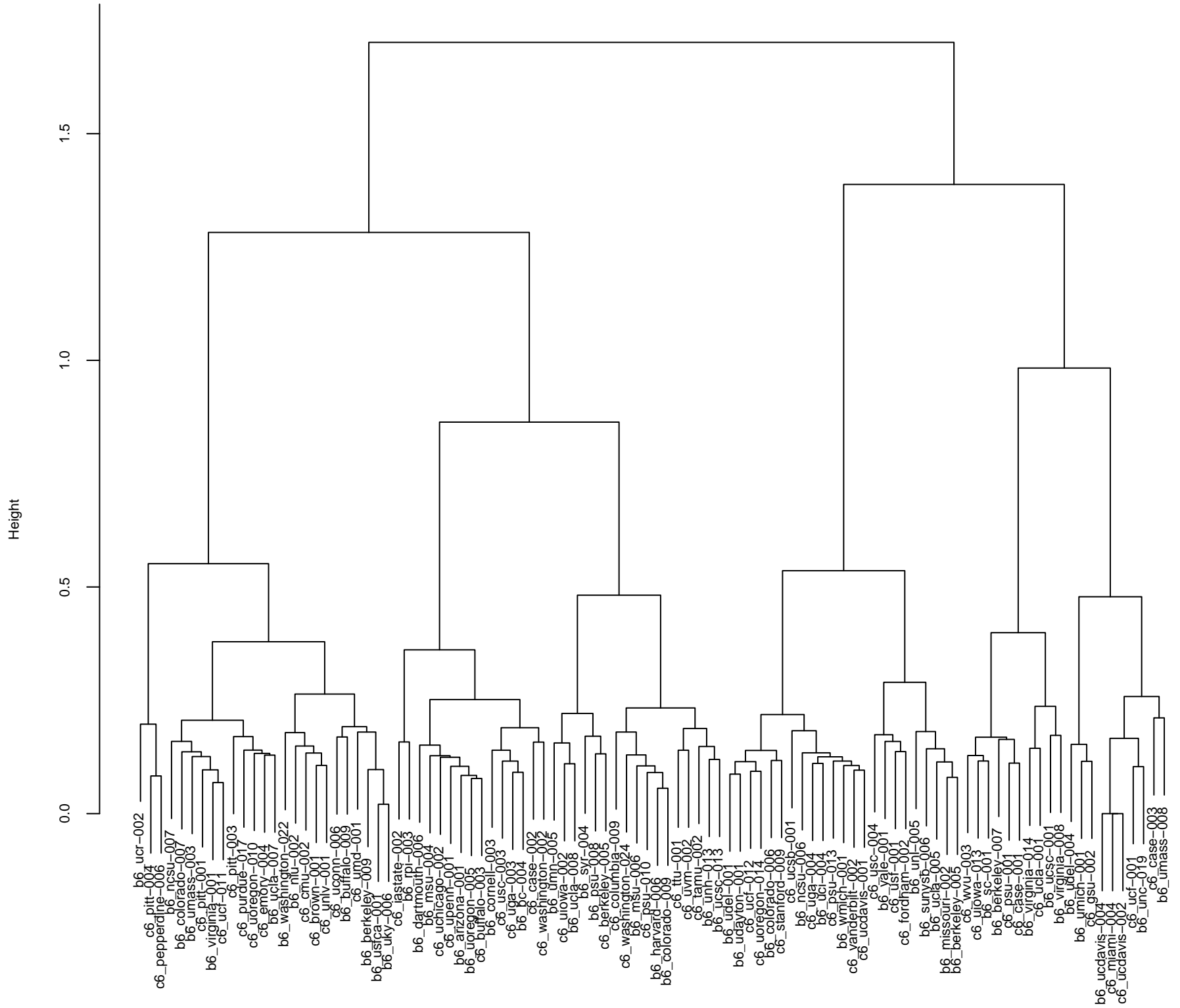


Cluster Dendrogram



as.dist(1 - cor(t(scalematr), method = "pearson"))

hclust (*, "complete")