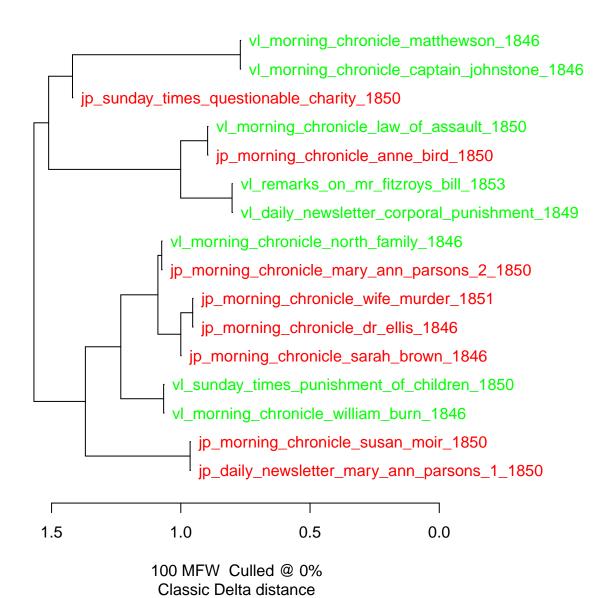
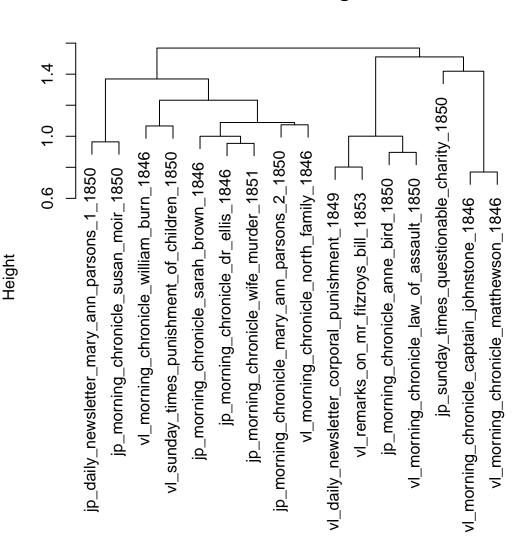
R_Script Cluster Analysis



Cluster Dendrogram



as.dist(data\$distance.table) hclust (*, "ward.D")