Hierarchical Clustering of Genes
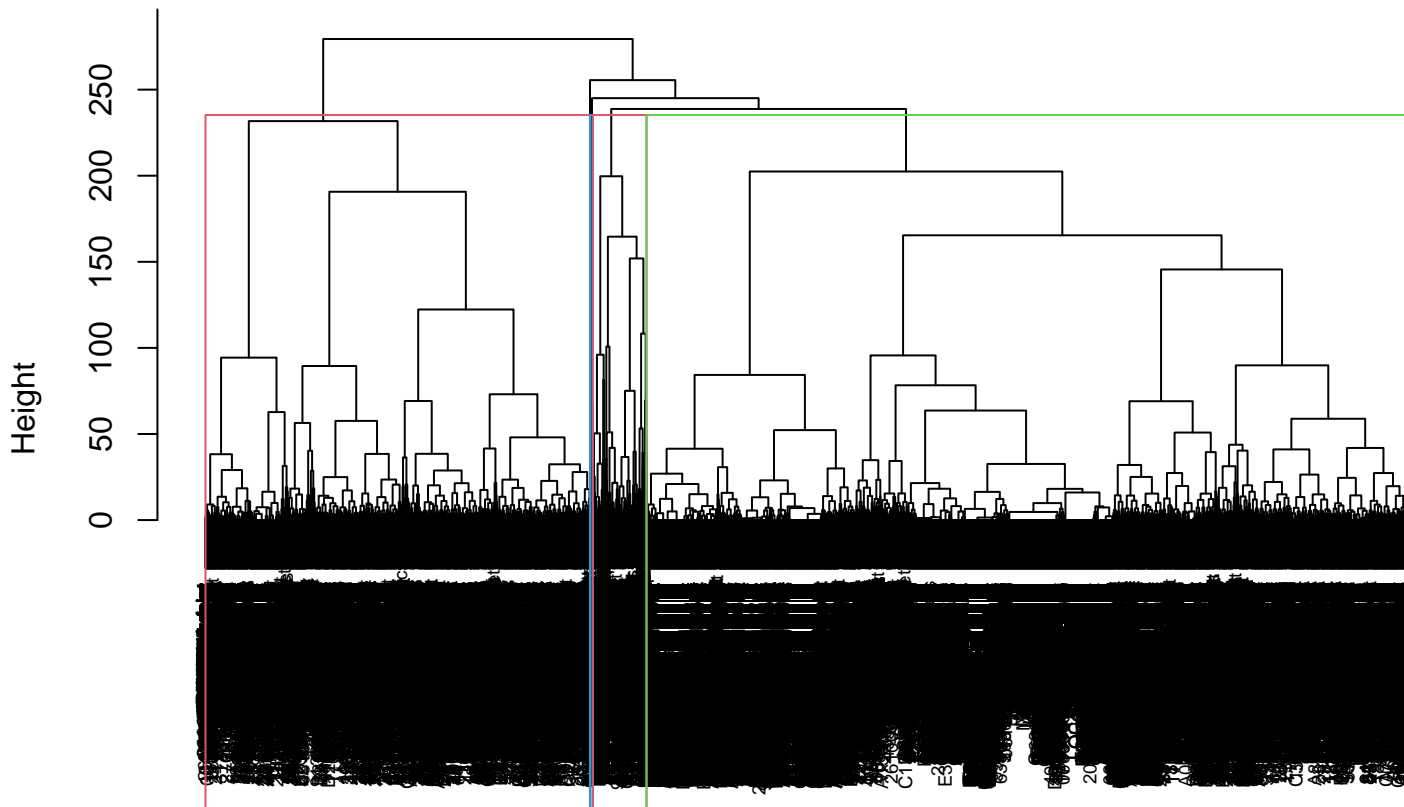
Height

dist_mat
hclust (*, "ward.D2")