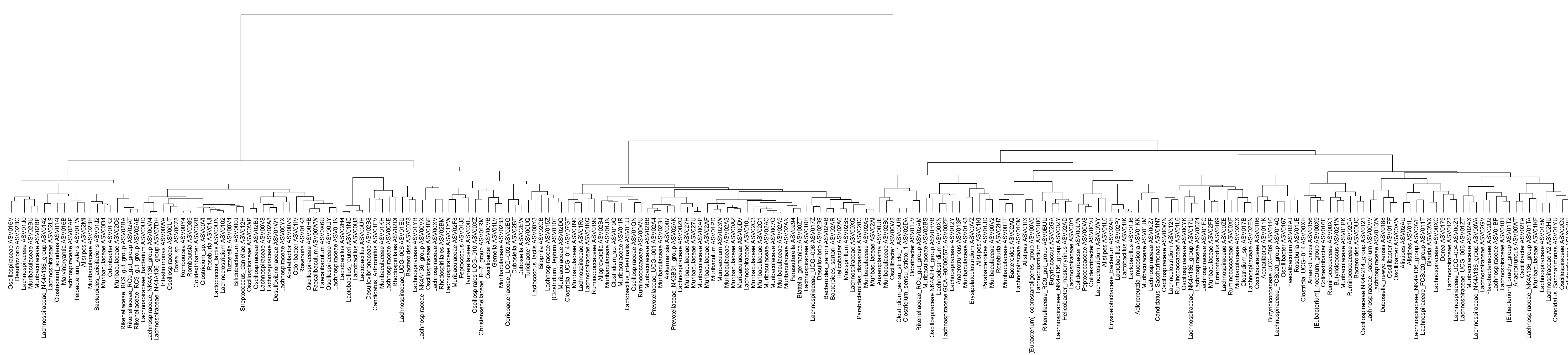


Cluster Dendrogram

Height



as.dist(mydist)

hclust (*, "ward.D2")