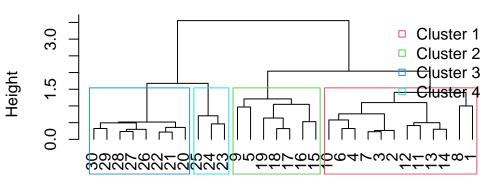
Reordered dendrogram from hclust(d = spe.ch, method = "ward.D2")



30 sites 4 clusters