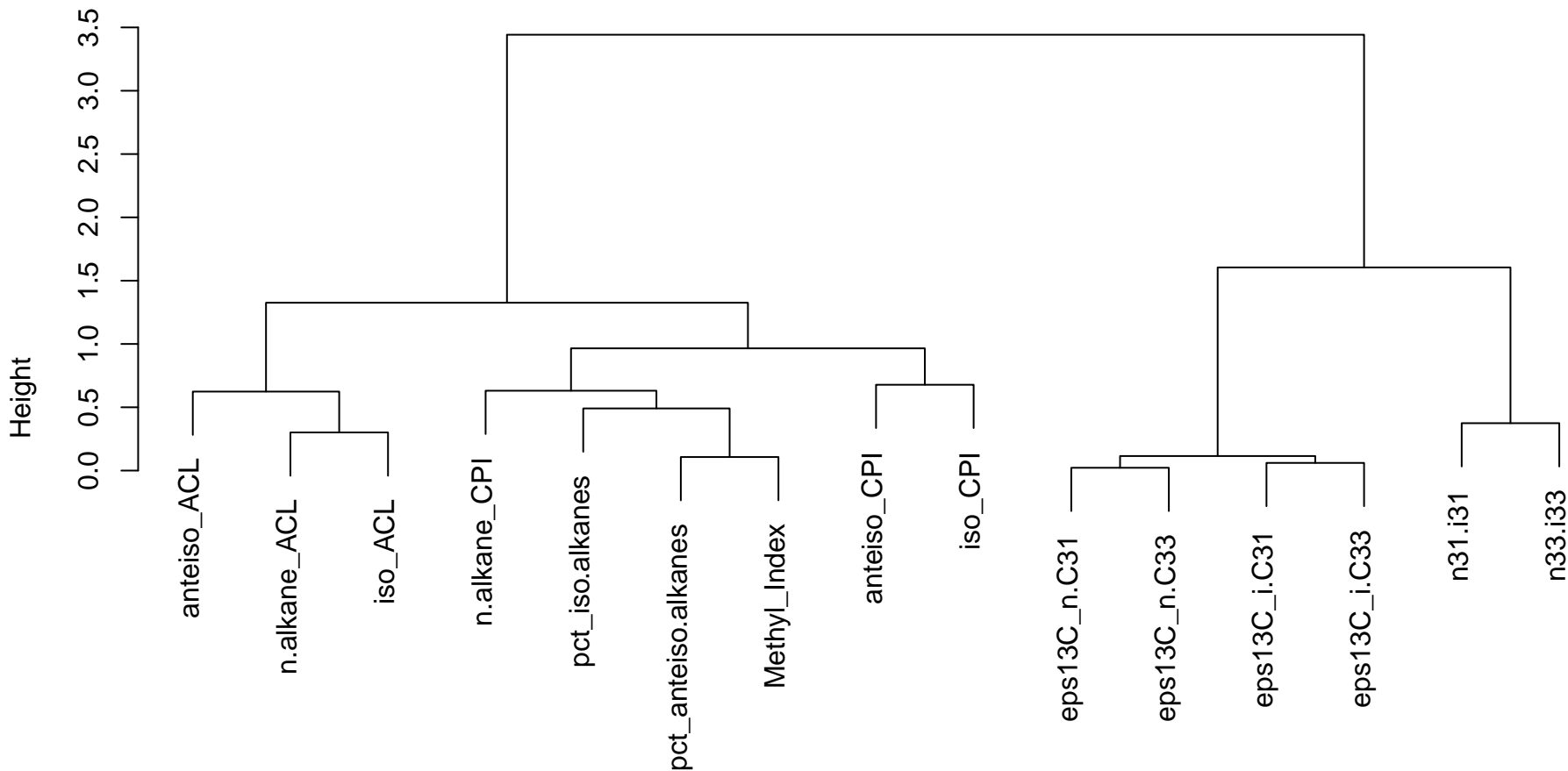


Cluster Dendrogram



as.dist(1 - abs(cor(snd, method = "spearman", use = "pairwise.complete.obs")))
hclust (*, "ward.D")