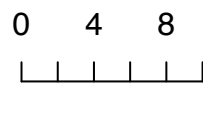
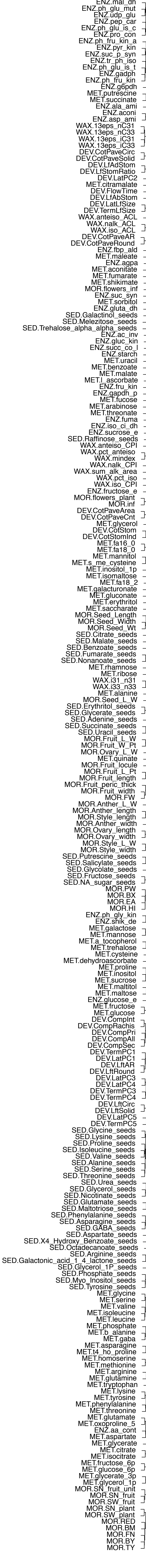


Height



Cluster Dendrogram



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