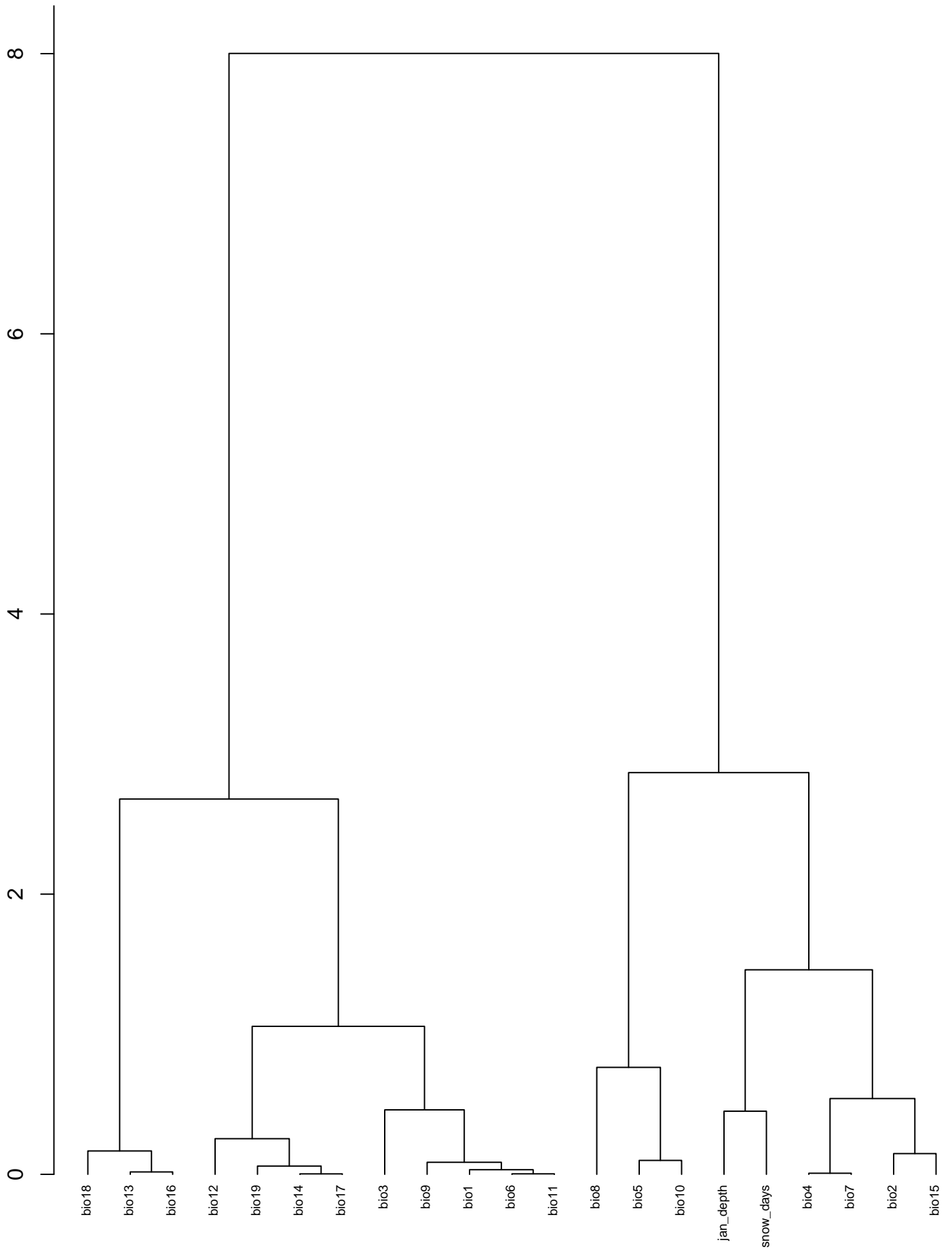


# Cluster Dendrogram



as.dist(1 – corplot2)  
hclust (\*, "ward.D")