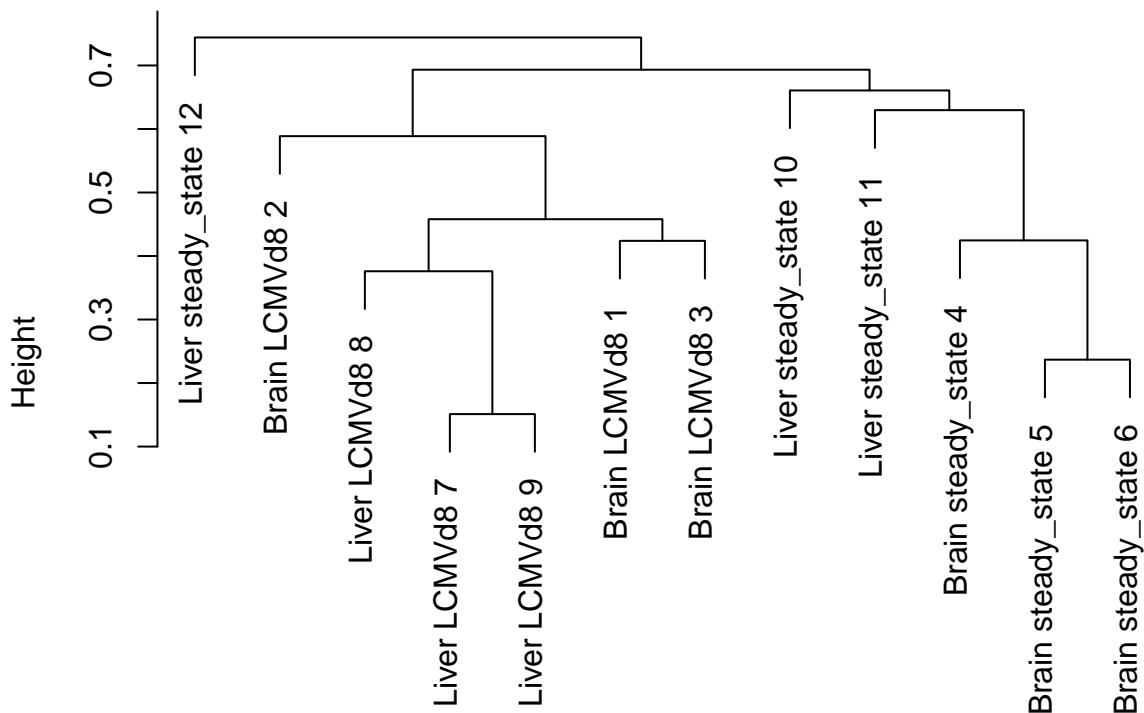


Cluster Dendrogram



as.dist(1 - corr_spearman_combat)
hclust (*, "complete")