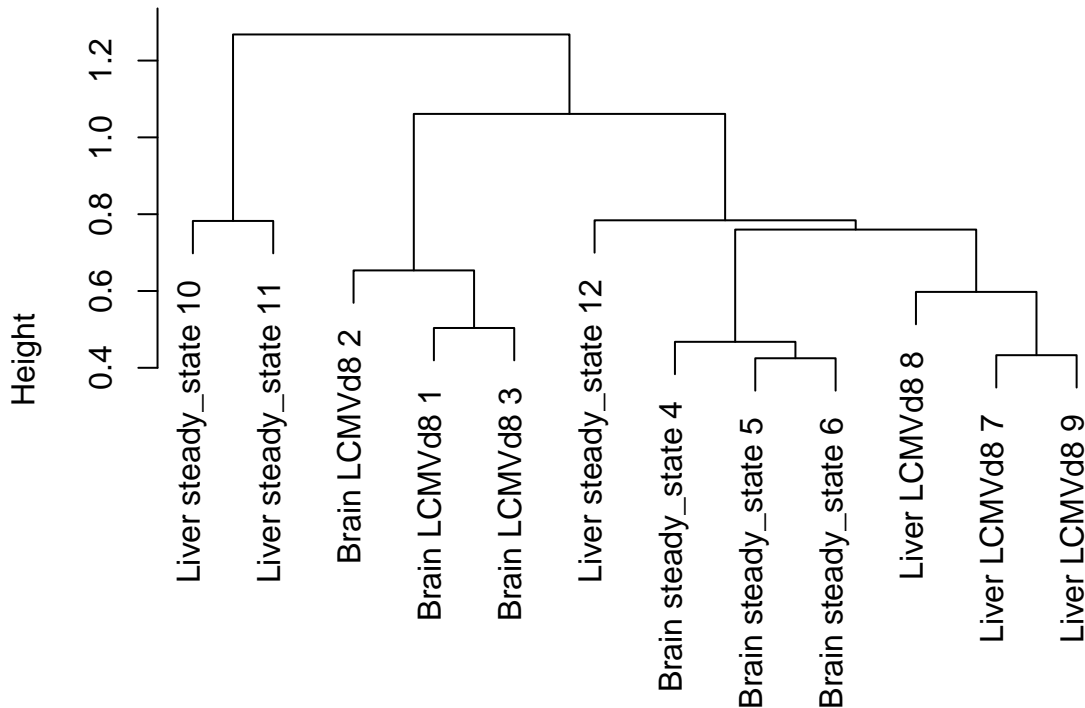


Cluster Dendrogram



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