Cluster Dendrogram

Height
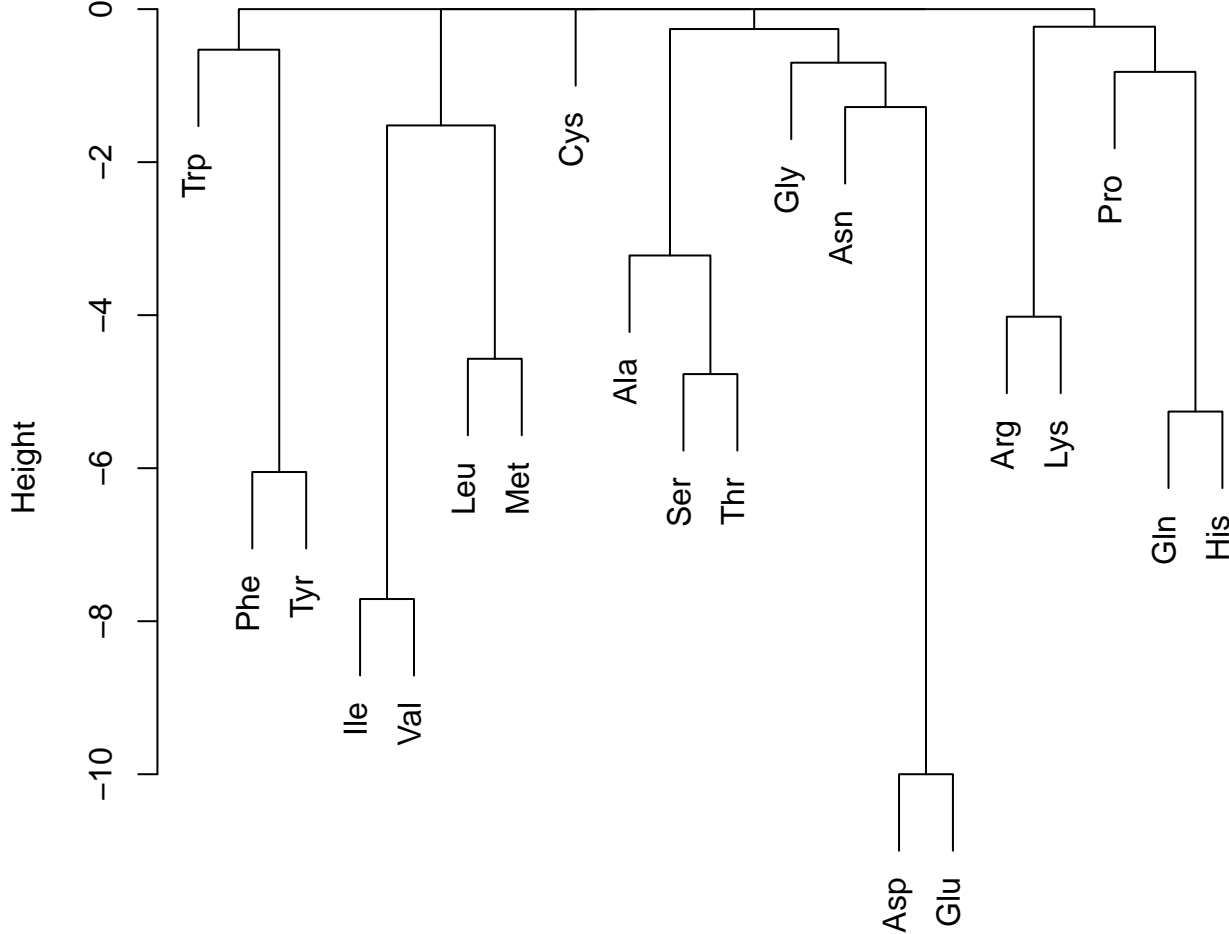
as.dist(read.table(args, head = TRUE))
hclust (*, "complete")