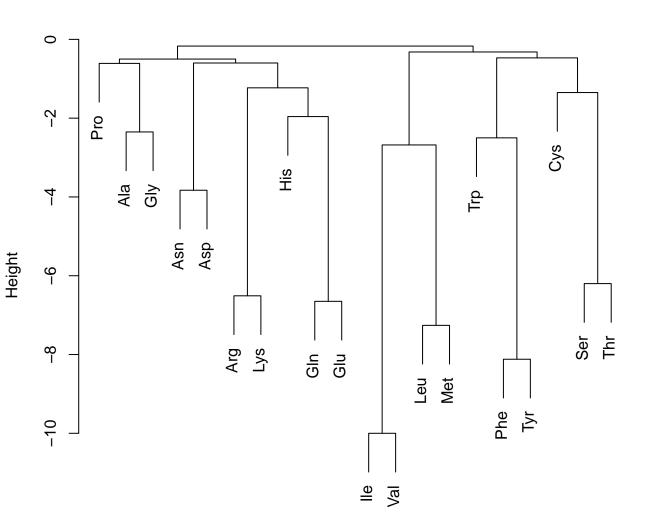
Cluster Dendrogram



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