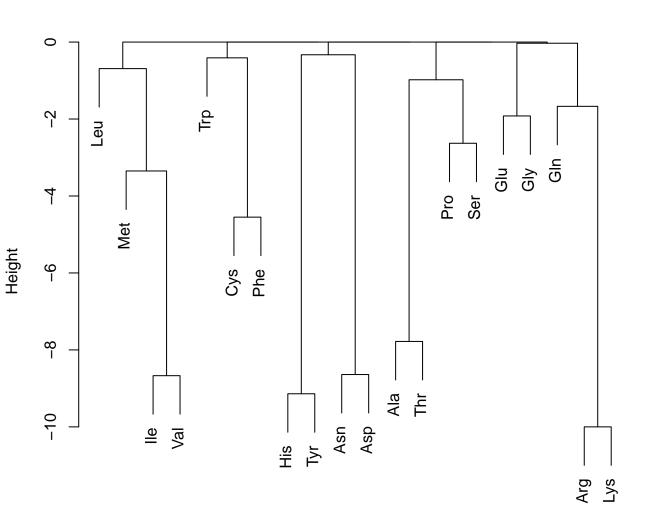
## **Cluster Dendrogram**



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