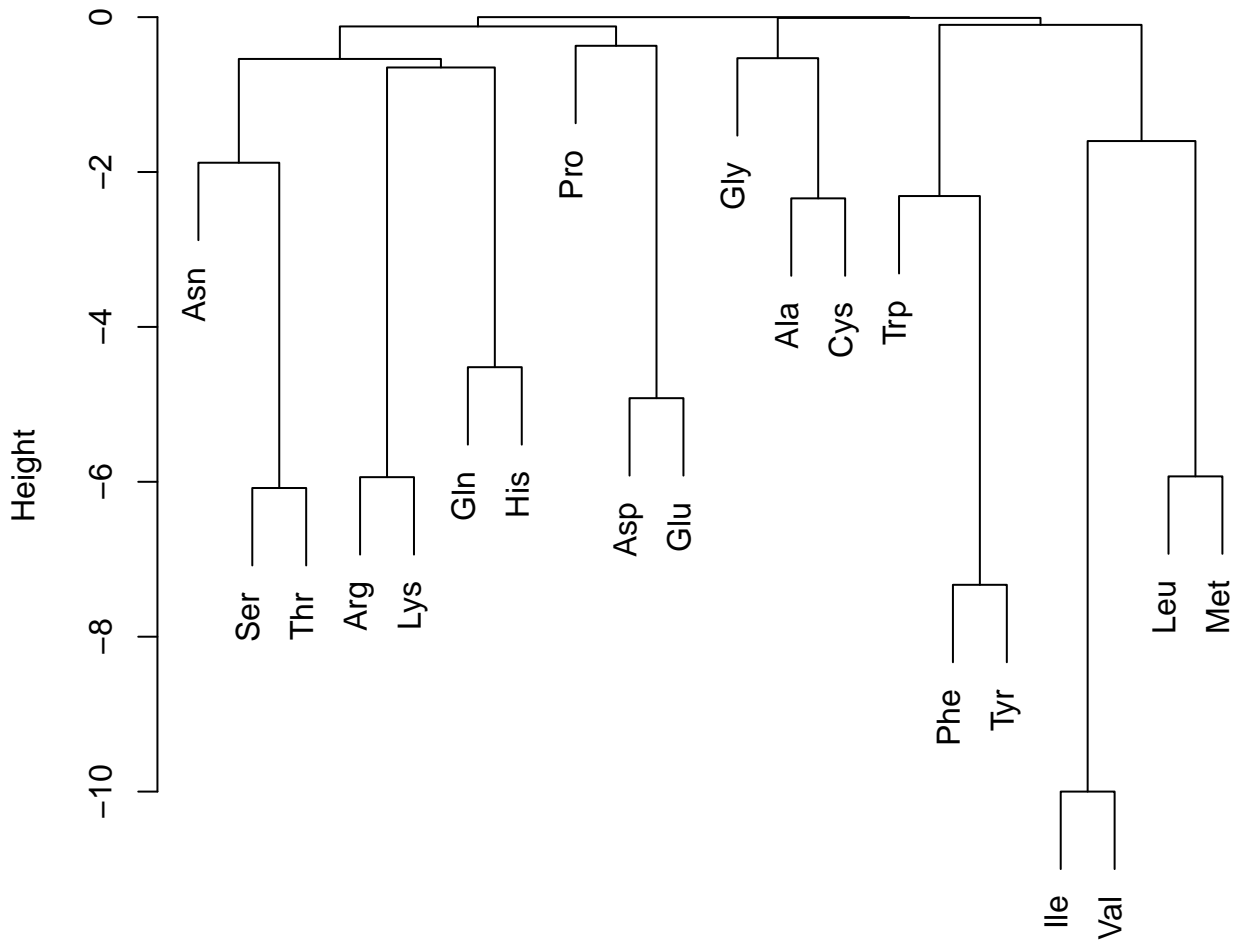


# Cluster Dendrogram



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