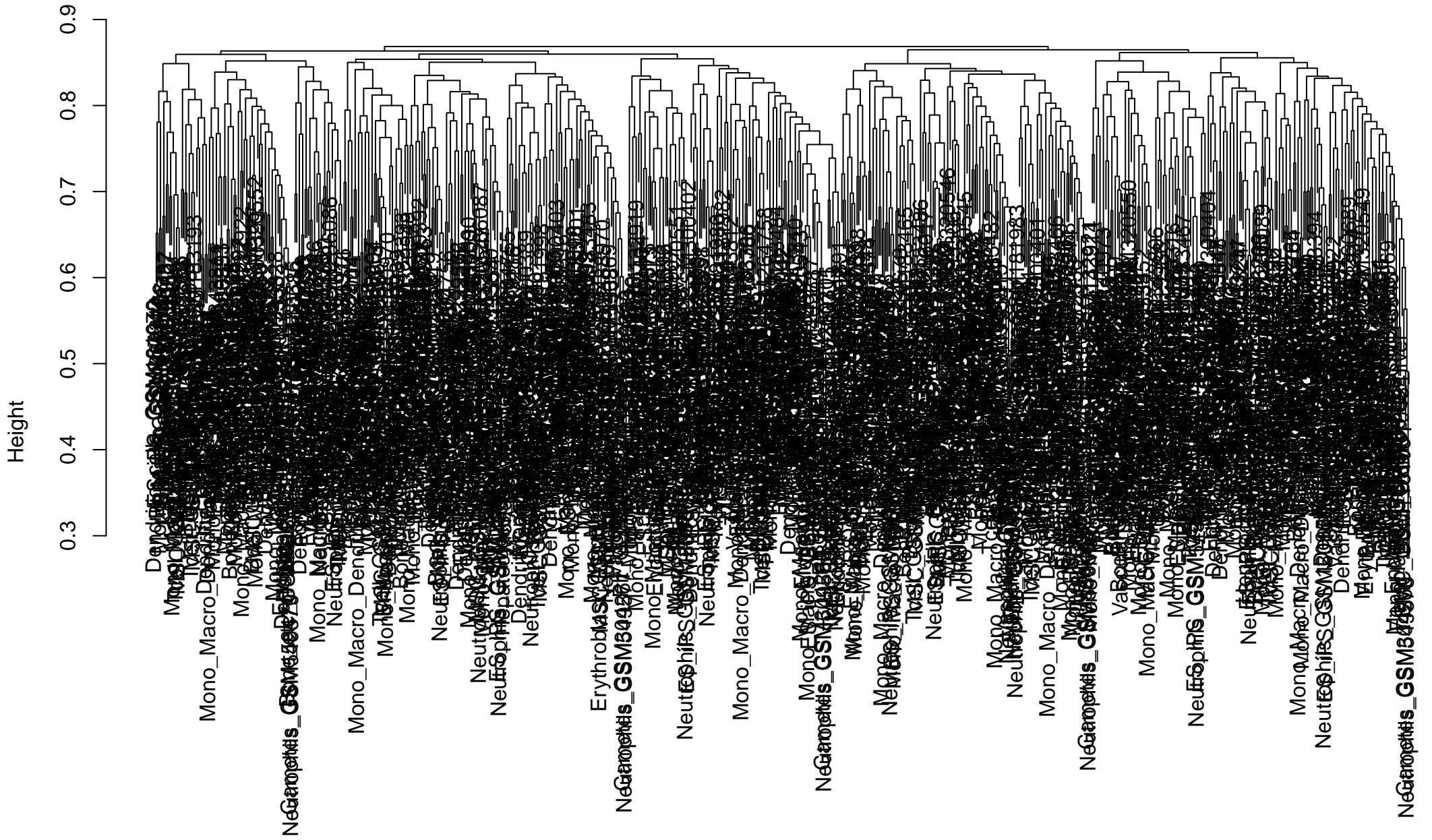


Cluster Dendrogram



```
as.dist(d)
hclust (*, "ward.D2")
```