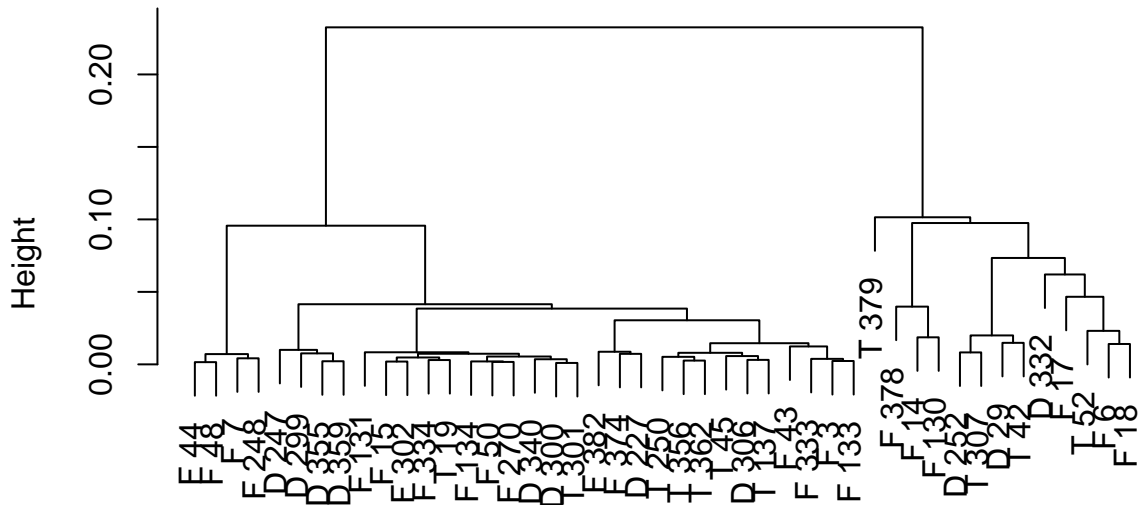


# Cluster Dendrogram



```
dist(clust_DCM, method = "maximum")  
hclust (*, "ward.D")
```