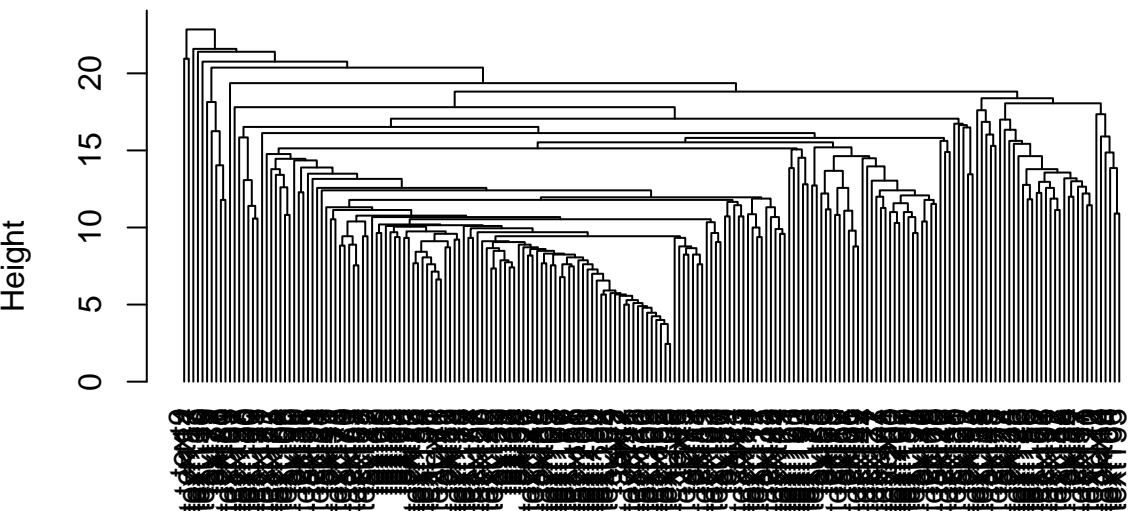


Cluster Dendrogram



```
as.dist(corpus_distances)  
hclust (*, "complete")
```