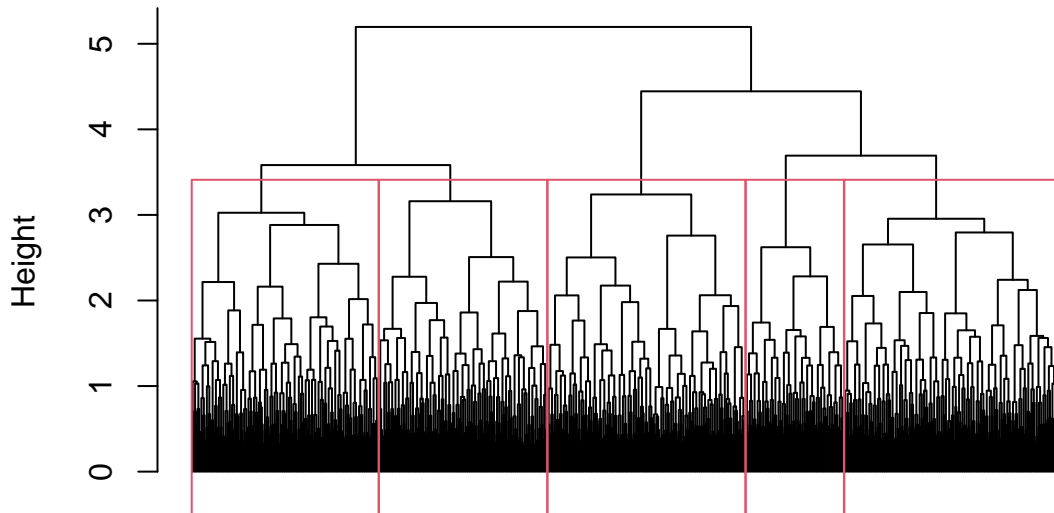


# Cluster Dendrogram



```
as.dist(1 - matrix_similarity)  
fastcluster::hclust (*, "ward.D2")
```