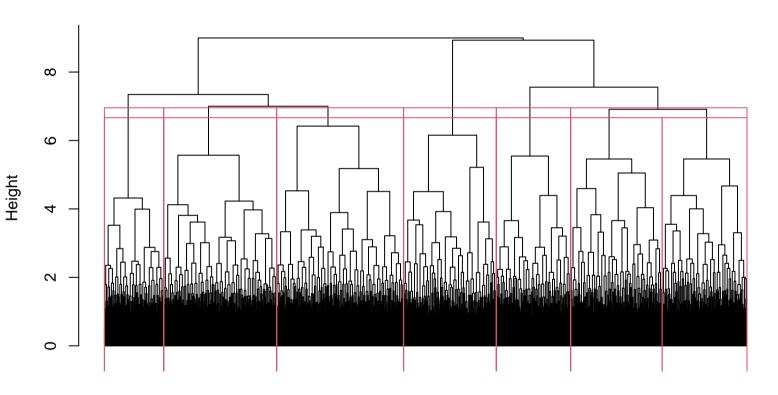
Cluster Dendrogram



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