Cluster Dendrogram

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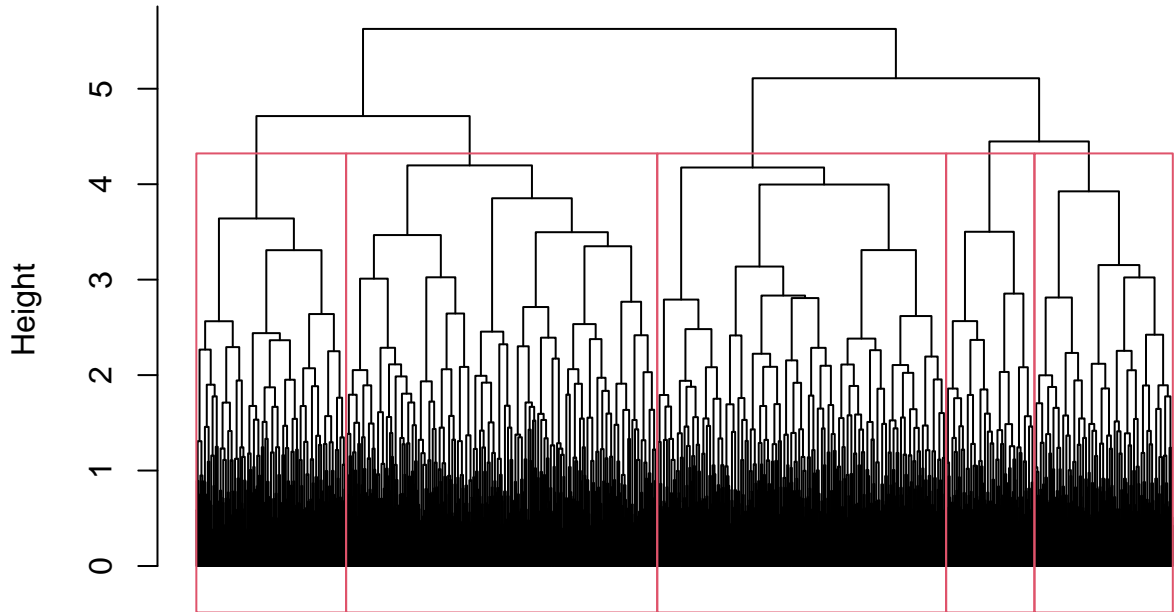
as.dist(1 − matrix_similarity)
fastcluster::hclust (*, "ward.D2")