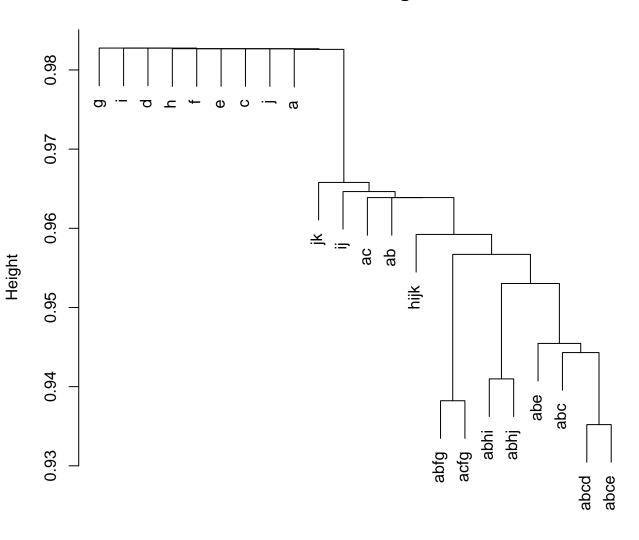
Cluster Dendrogram



as.dist(distance_matrix) hclust (*, "single")