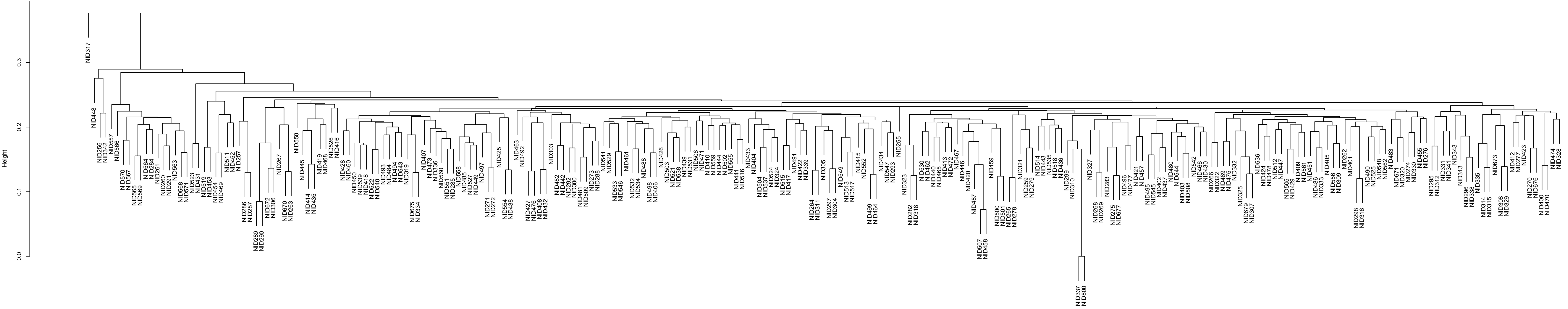


Cluster Dendrogram



as.dist(1 - kmatrix)
hclust (*, "complete")