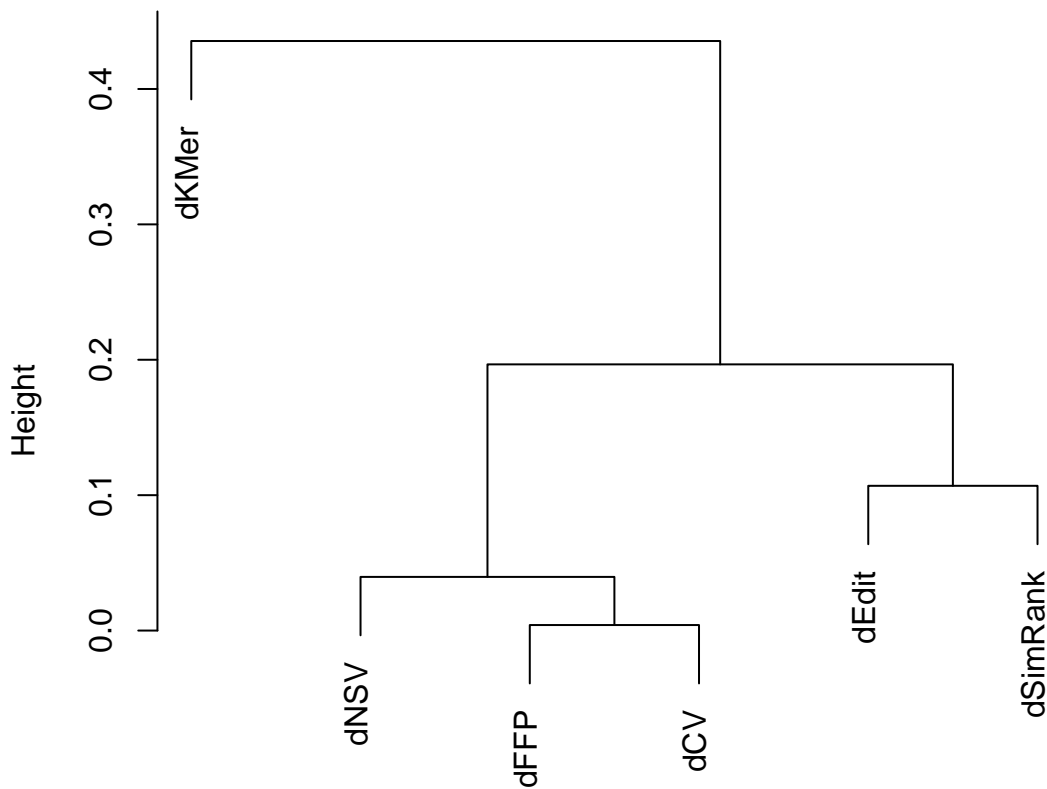


Cluster Dendrogram



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