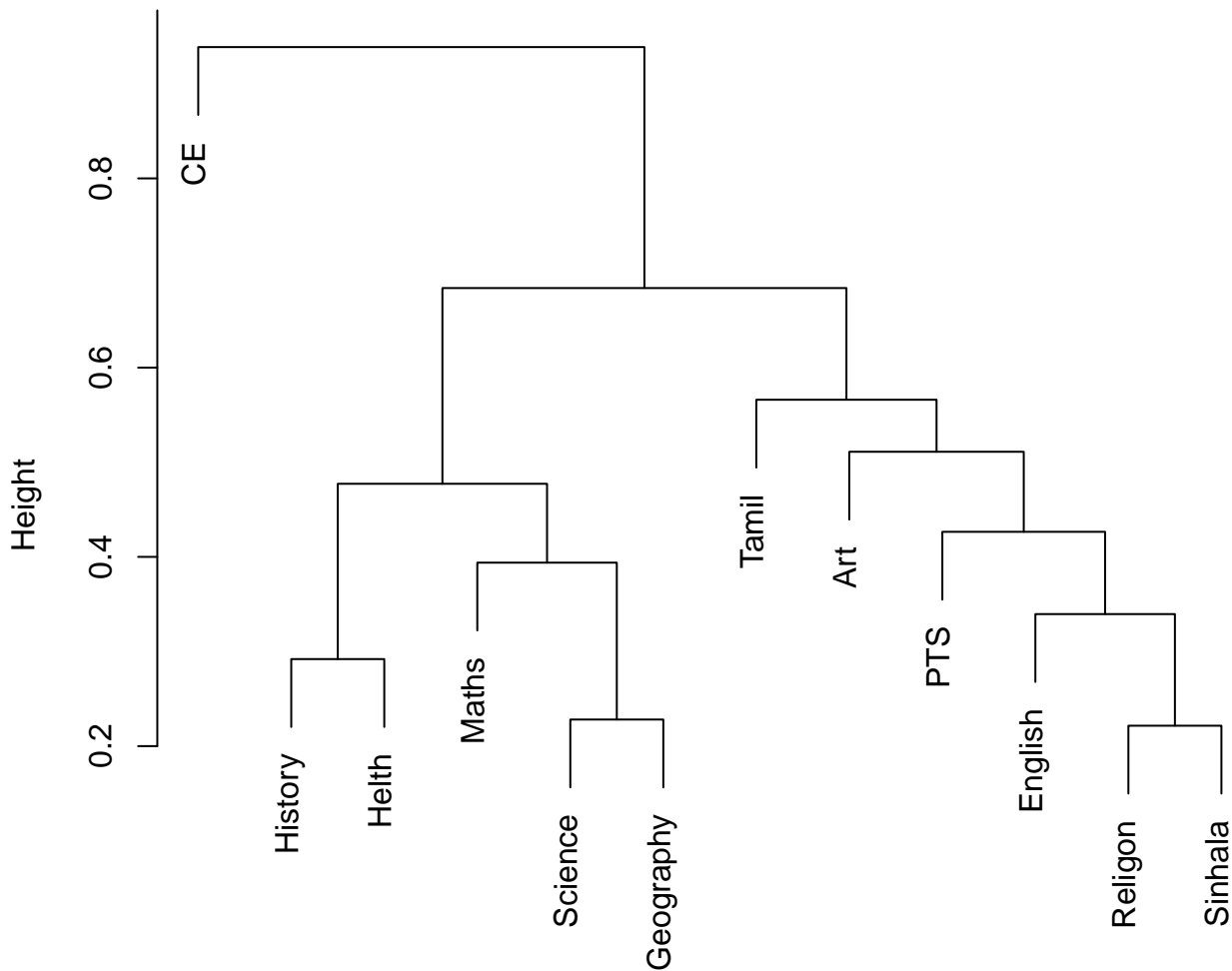


Cluster Dendrogram



dist(abs(cor(na.omit(grade_6))))
hclust (*, "complete")