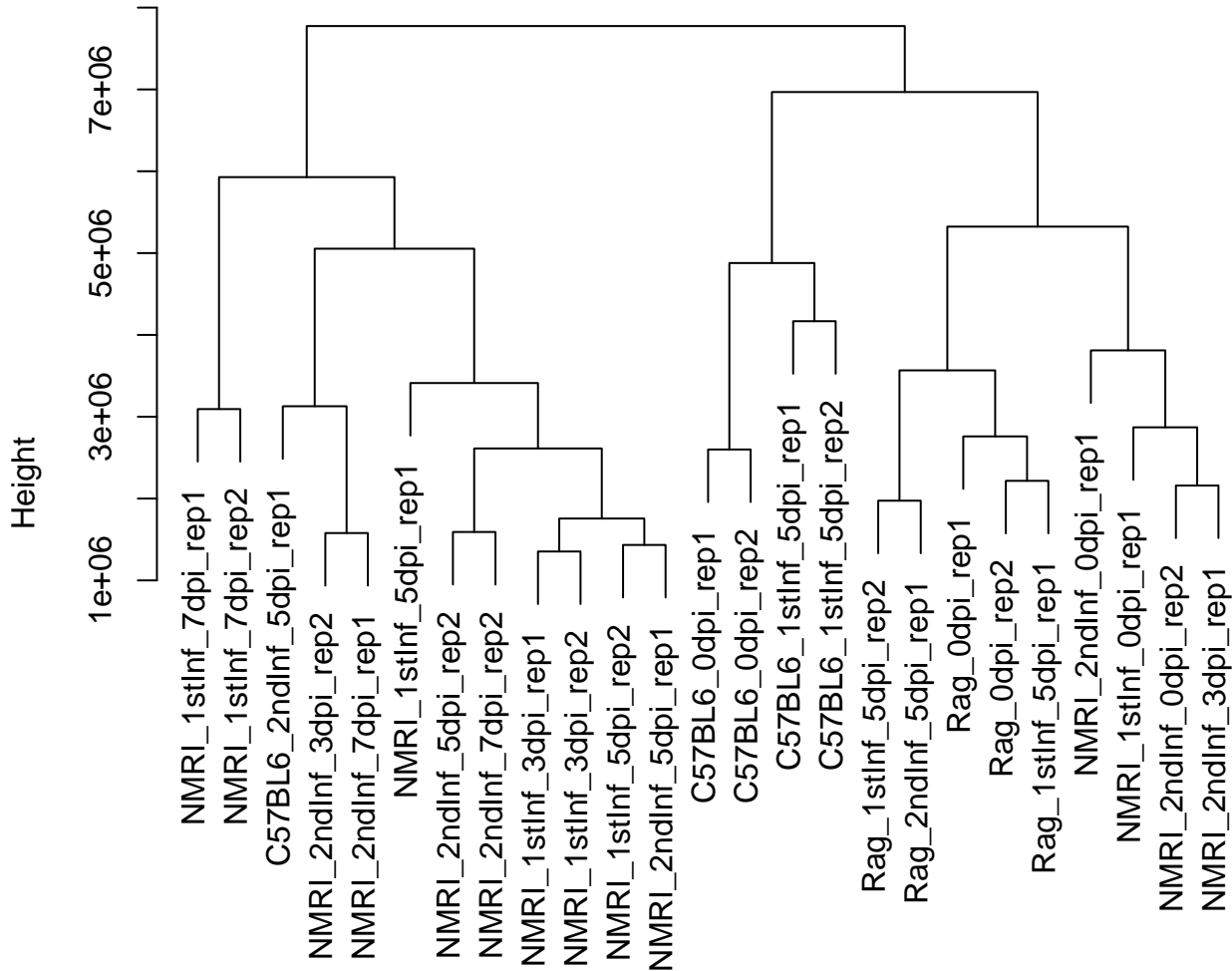


Cluster Dendrogram



dist(t(normCounts(Mm.RUVset.empirical)))
hclust (*, "complete")