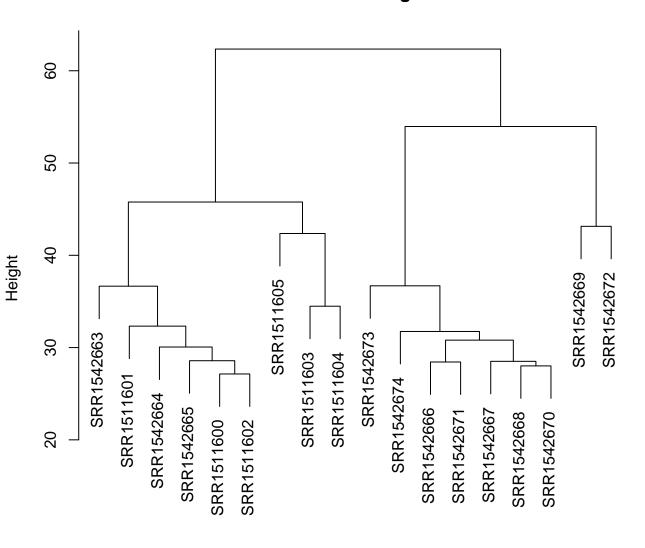
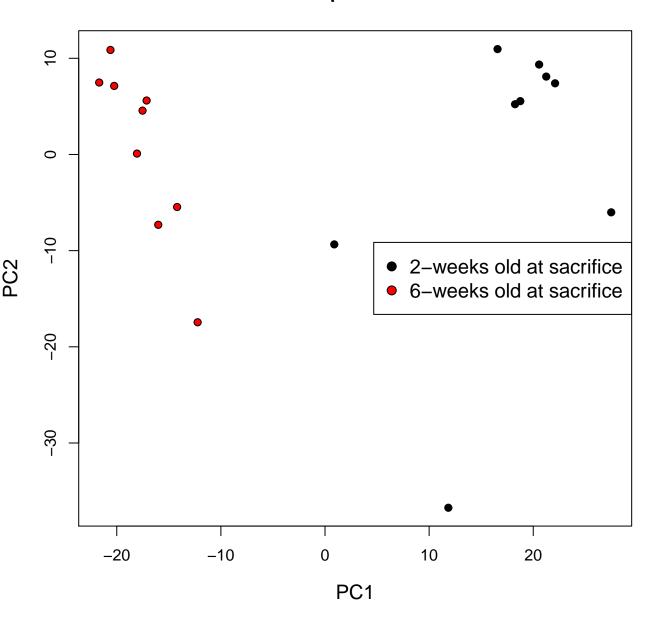
Cluster Dendrogram

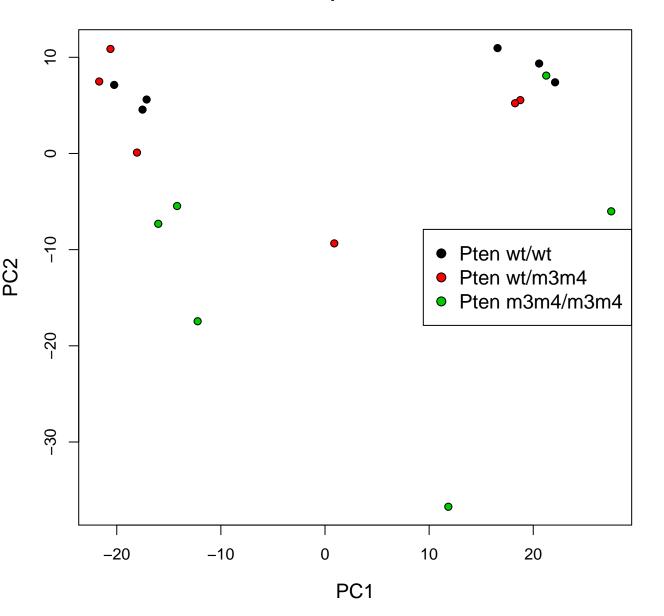


dist(t(yGene))
hclust (*, "complete")

Gene expression PCA



Gene expression PCA



PC2 v. Gene Assignment Rate

