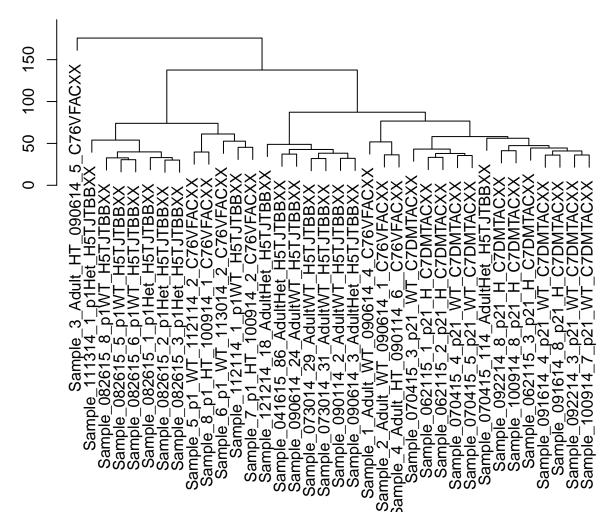
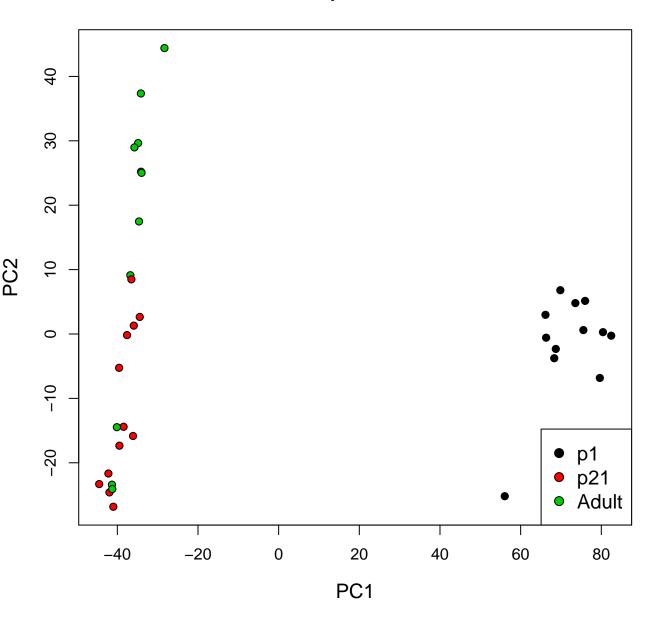
Cluster Dendrogram

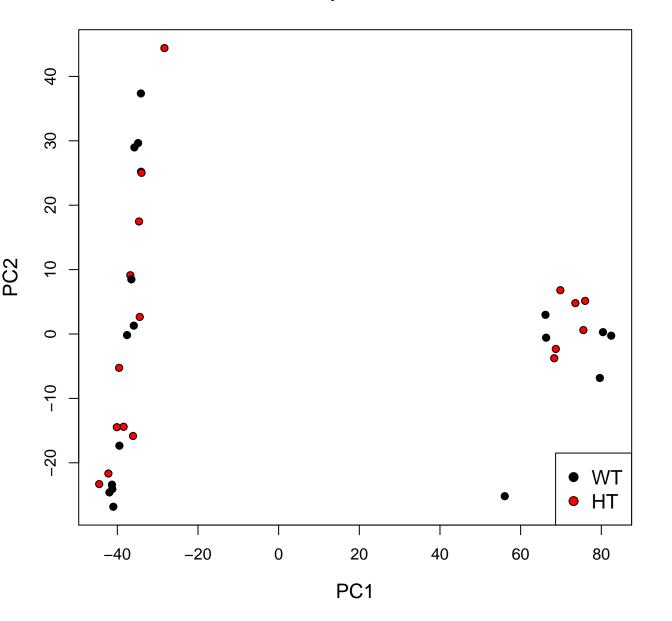


dist(t(yGene))
hclust (*, "complete")

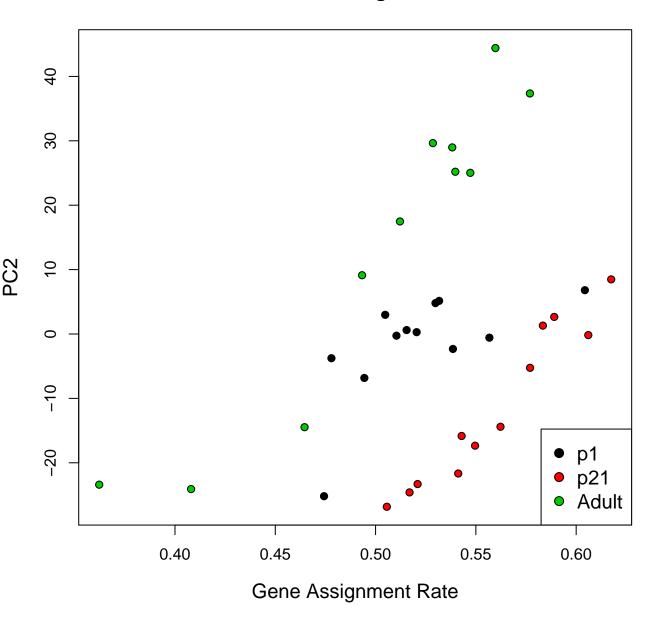
Gene expression PCA



Gene expression PCA



PC2 v. Gene Assignment Rate



PC2 v. MitoRate

