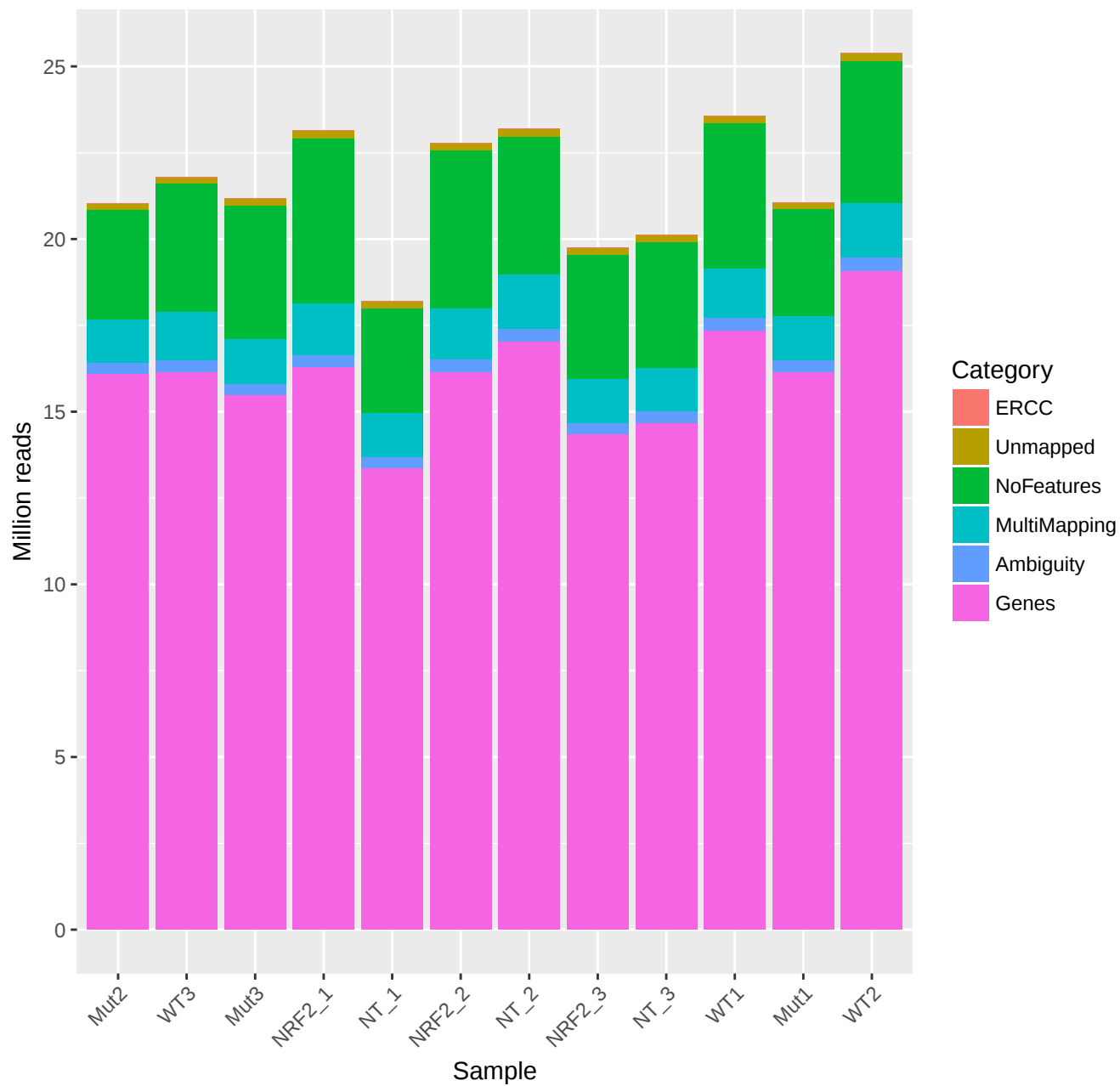
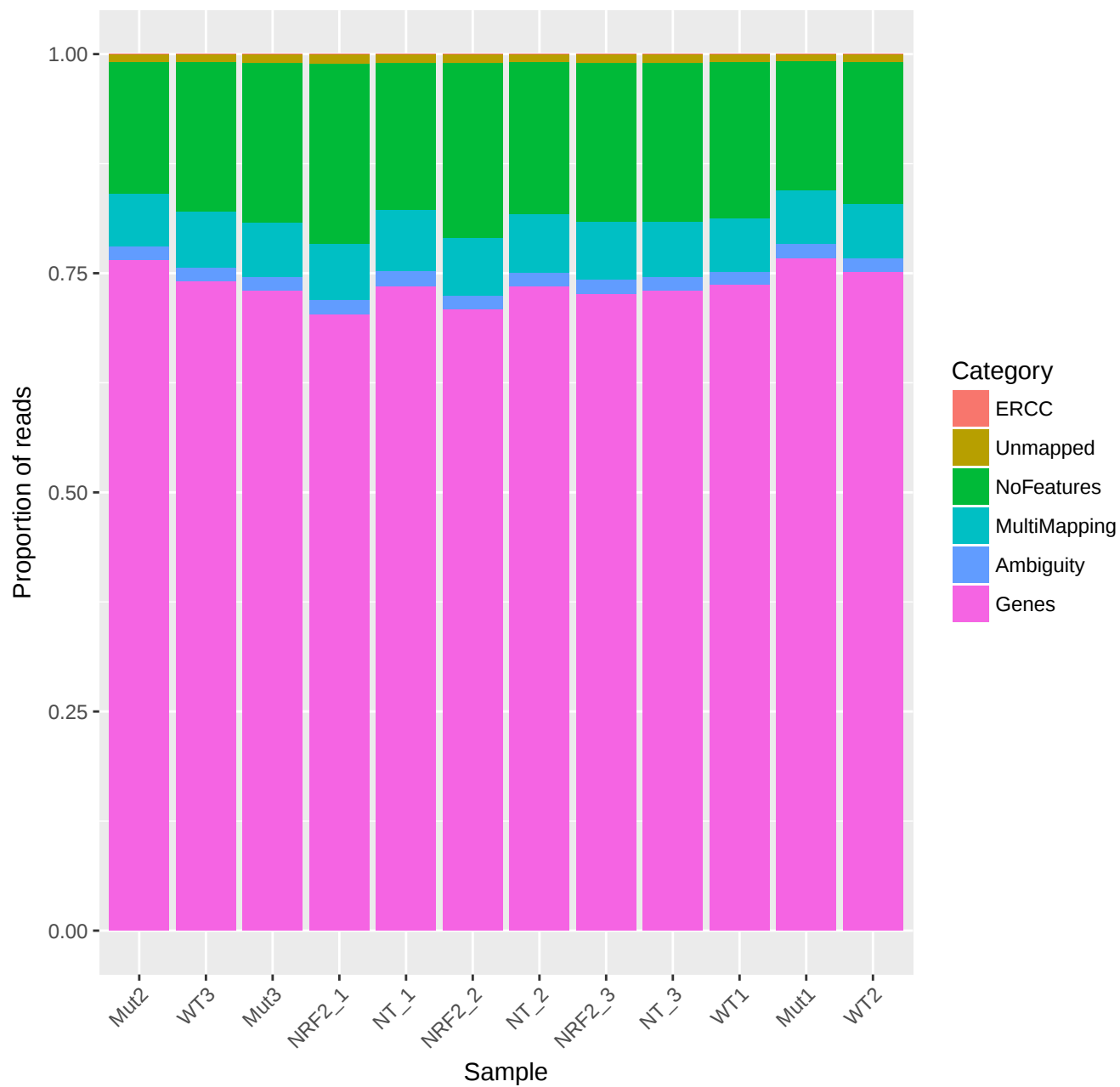


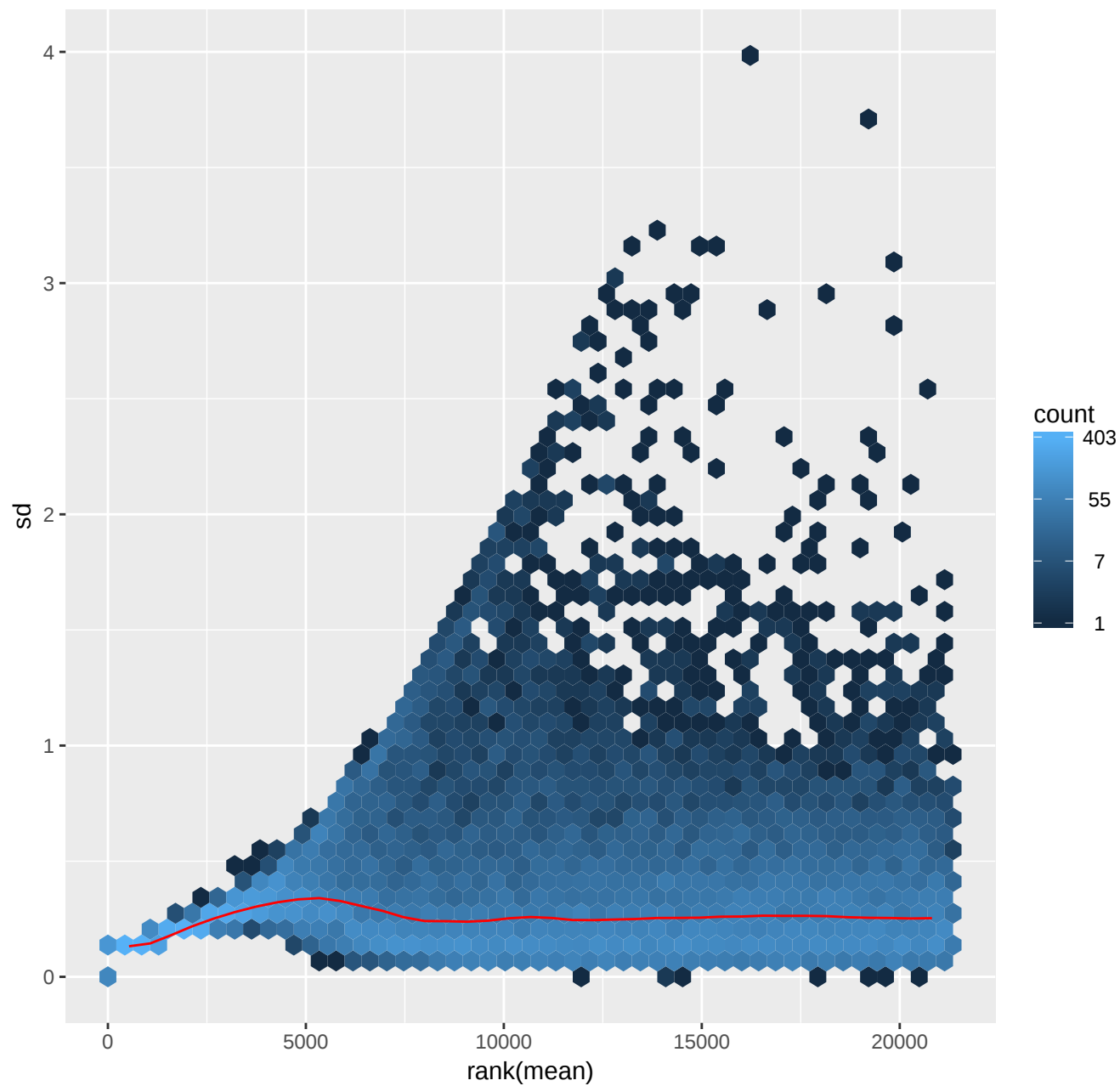
Read assignment by category



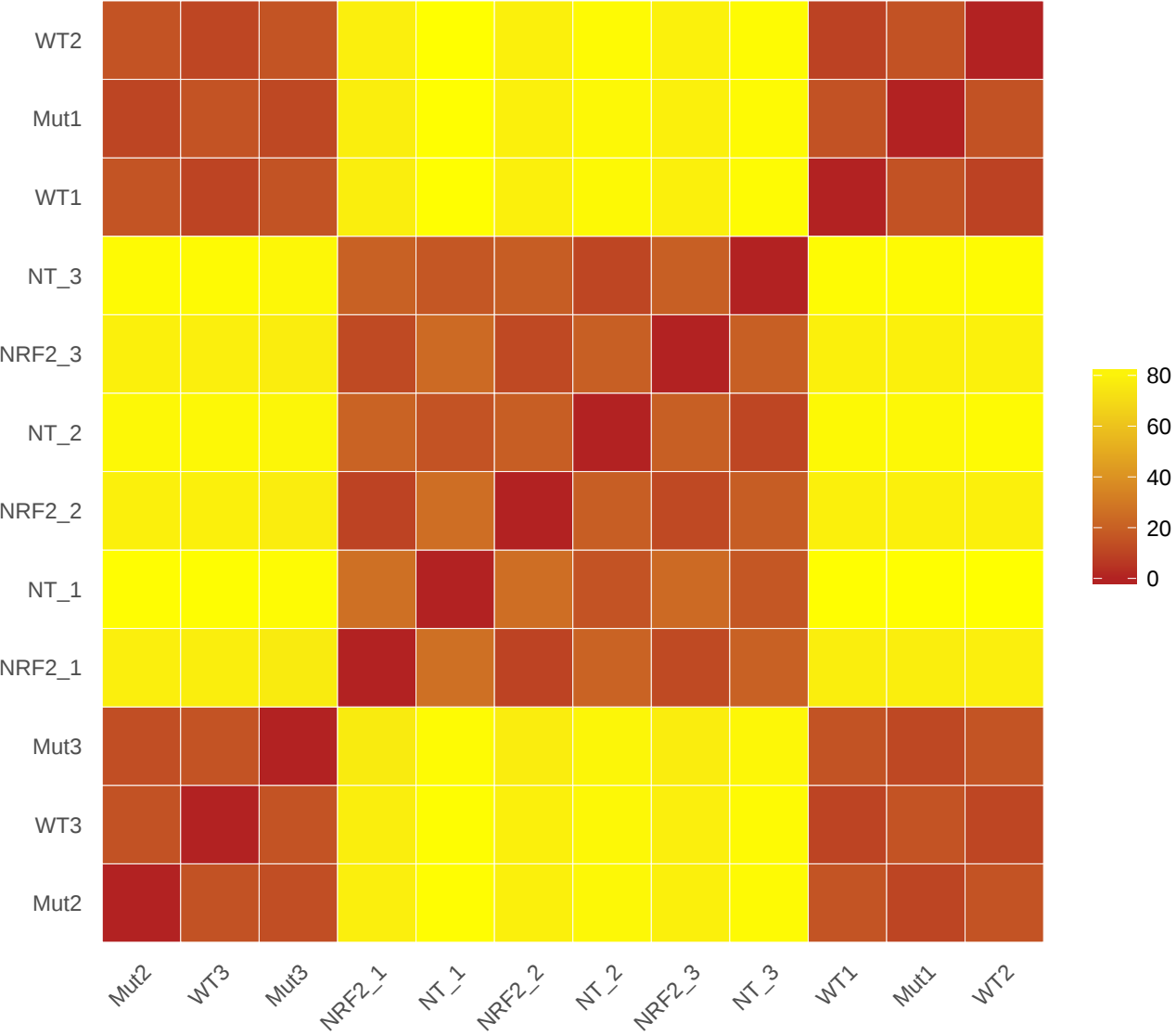
Read assignment by category



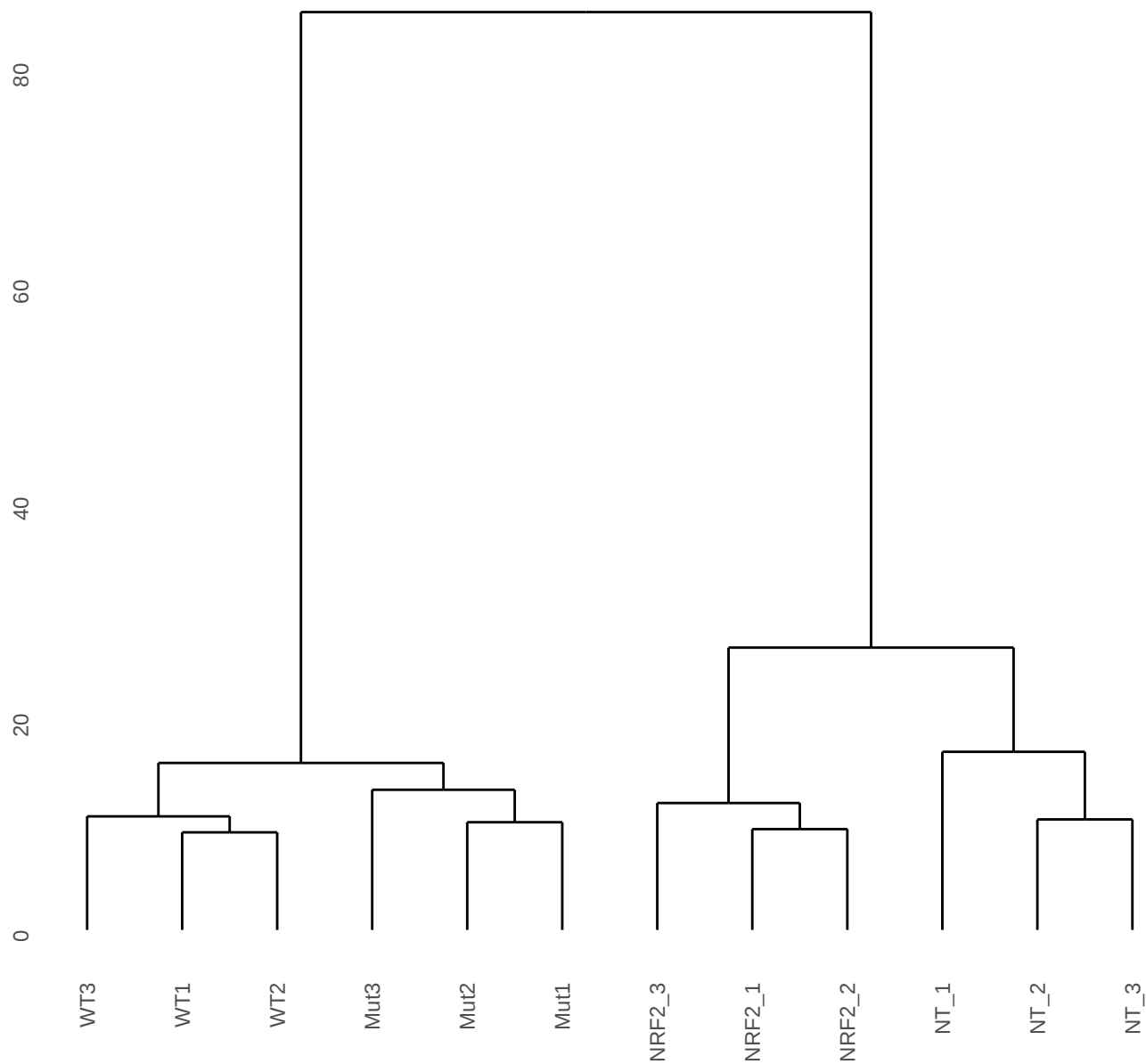
# Variance-stabilising normalisation



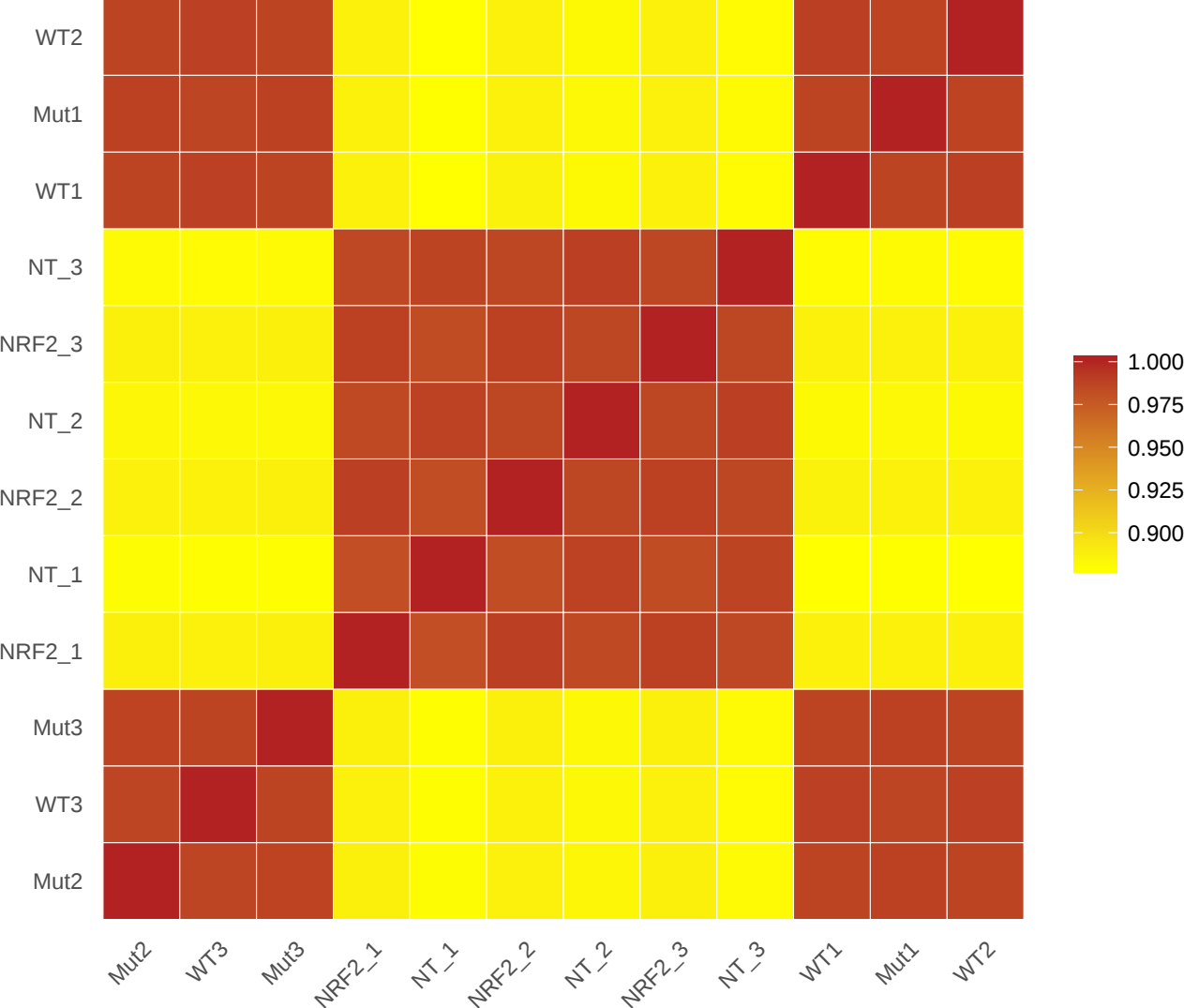
Distance-based heatmap



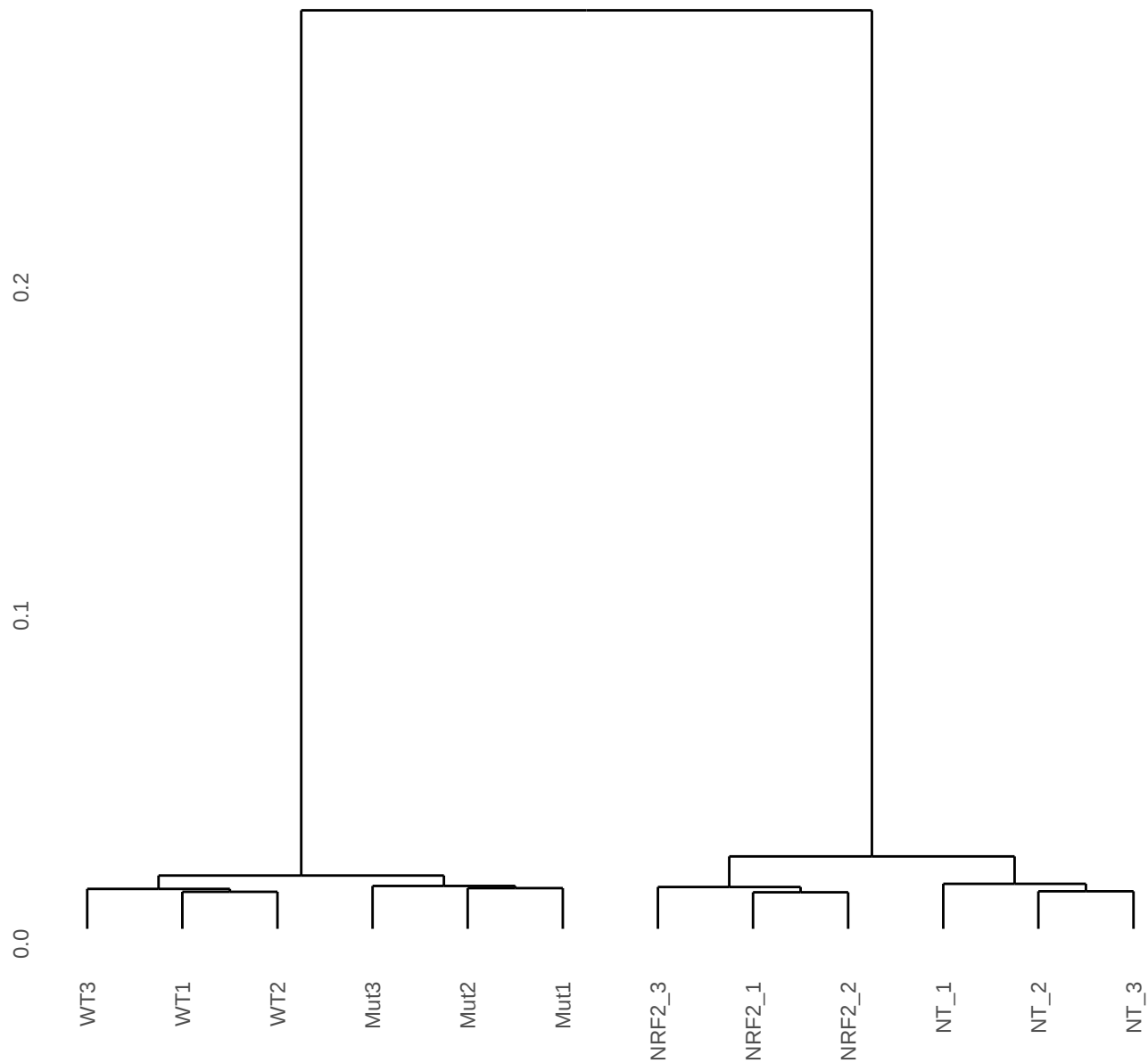
## Distance-based hierarchical clustering



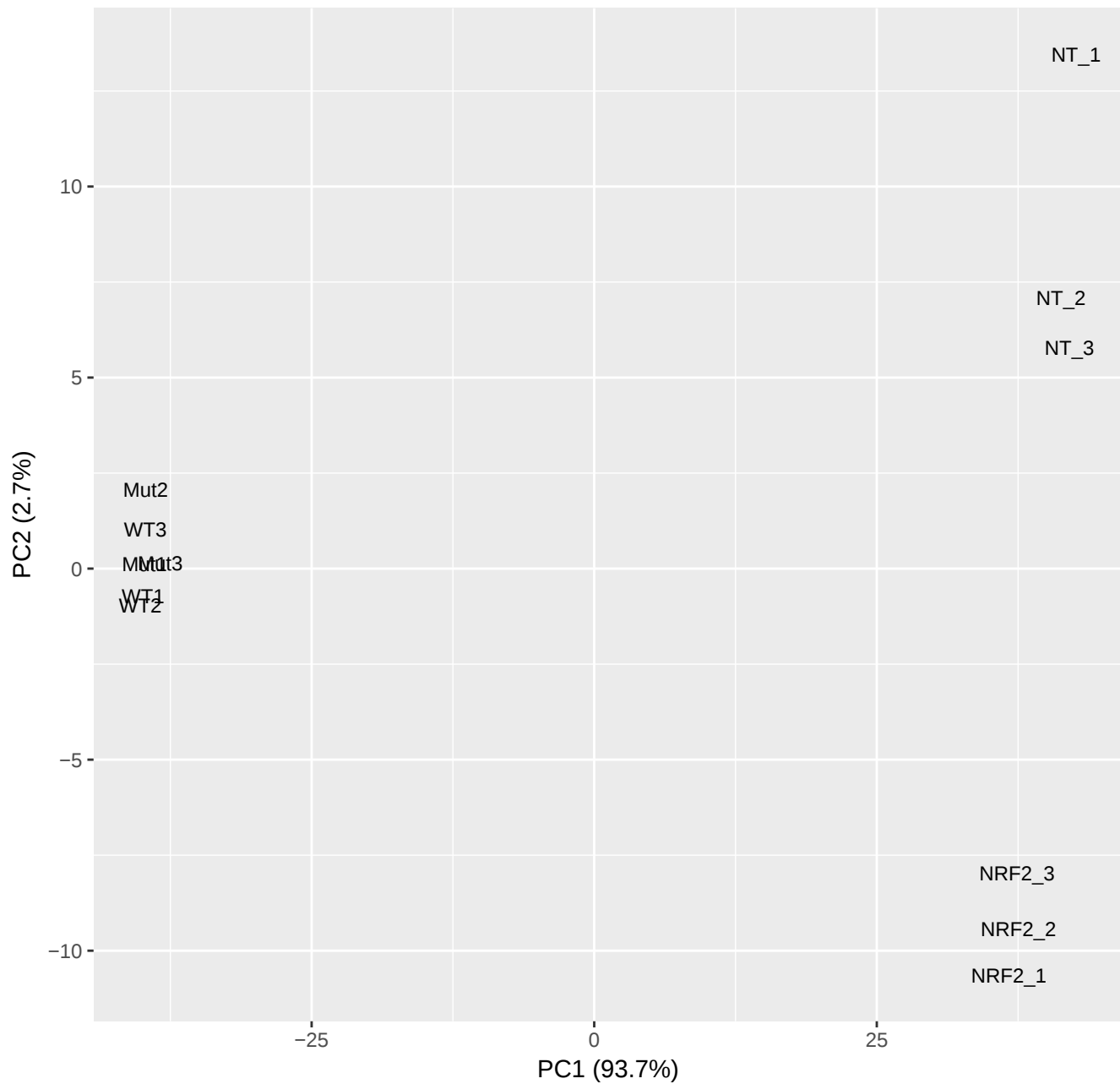
Spearman correlation heatmap



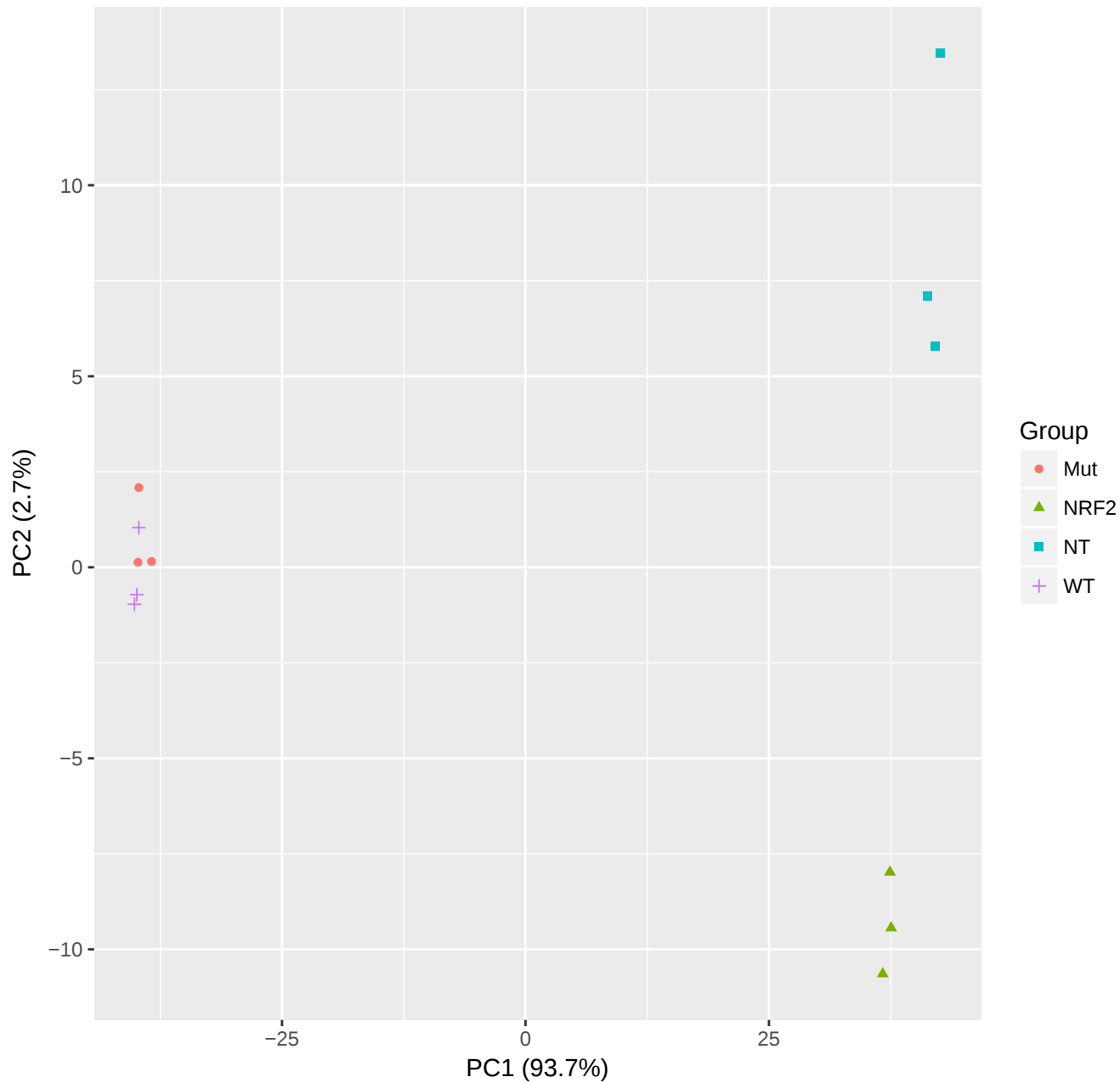
# Spearman's rank-based hierarchical clustering



PCA



PCA for group Group



Complexity curves

Percentage of reads for those genes

100

10

1

10

1000

Top n genes

Sample

Mut2  
WT3  
Mut3  
NRF2\_1  
NT\_1  
NRF2\_2  
NT\_2  
NRF2\_3  
NT\_3  
WT1  
Mut1  
WT2

