

# Experimental Design

SI87

9: 009\_tBIDrev

8: 008\_tBIDrev

7: 007\_tBIDrev

SI2

6: 006\_tBIDrev

5: 005\_tBIDrev

4: 004\_tBIDrev

Uninfected

3: 003\_tBIDrev

2: 002\_tBIDrev

1: 001\_tBIDrev

# IDENTIFICATIONS OVERVIEW (FDR 0.01)

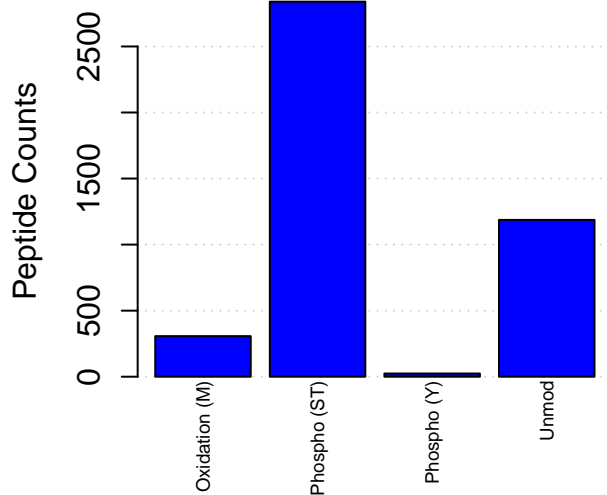
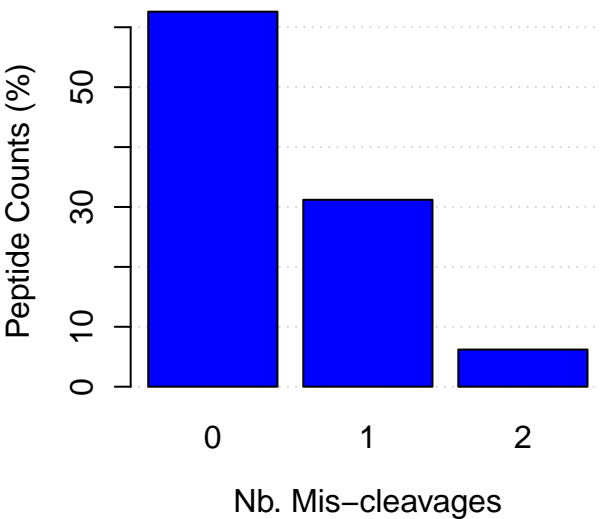
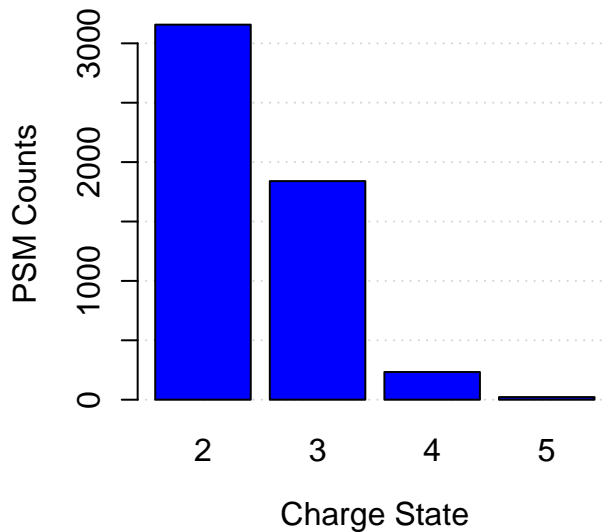
5252 PSM

3909 PEPTIDES

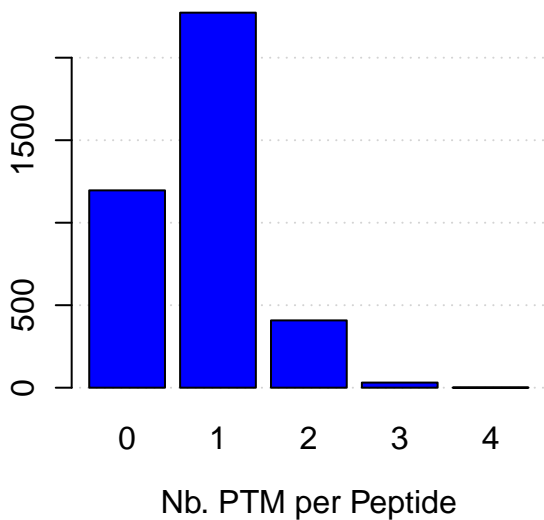
1196 UN-MOD.

2713 MOD.

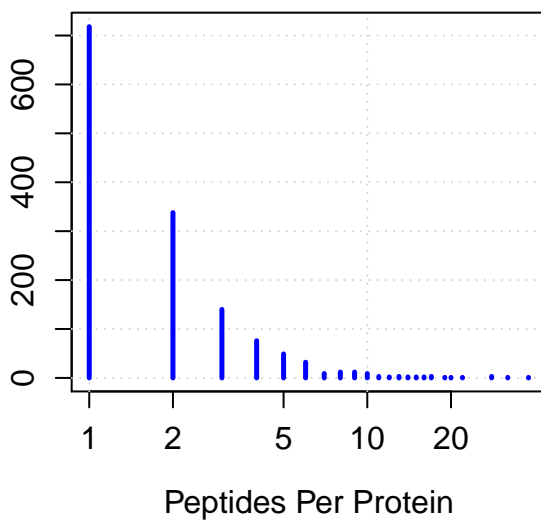
1420 PROTEINS



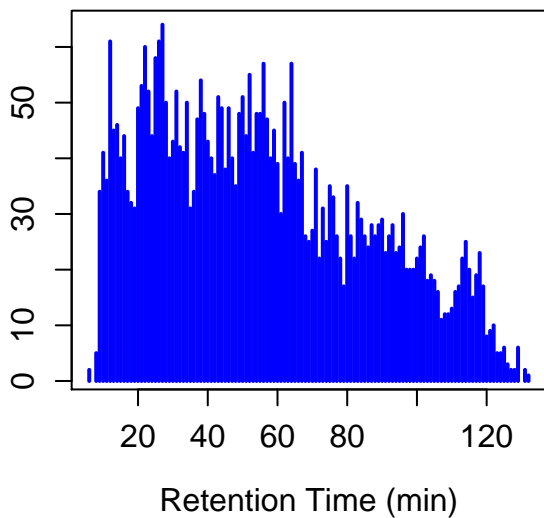
Peptide Counts



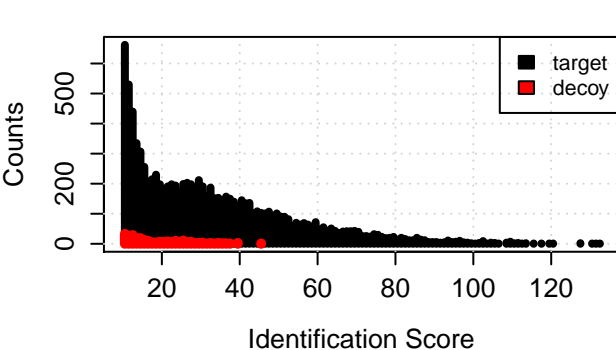
Protein Counts



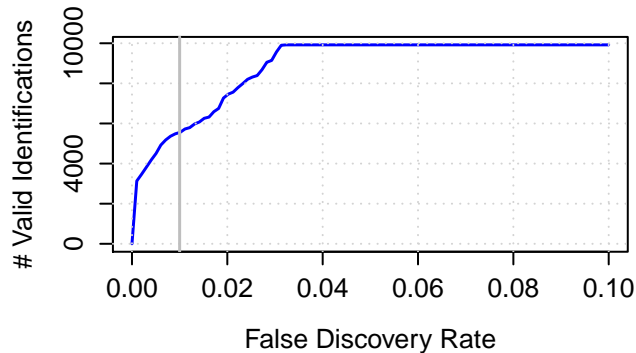
# Identified Features



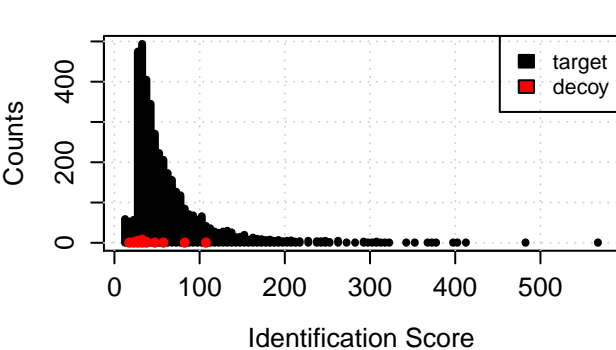
**Feature Level**



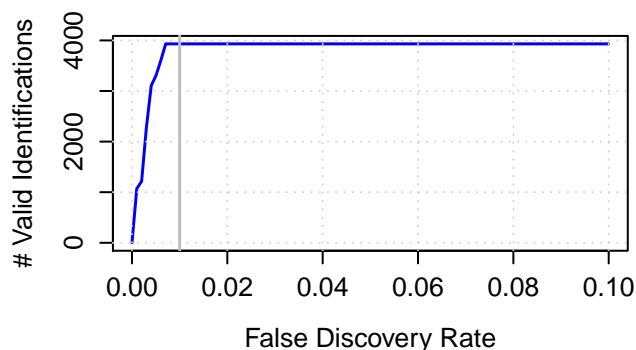
**Feature Level**



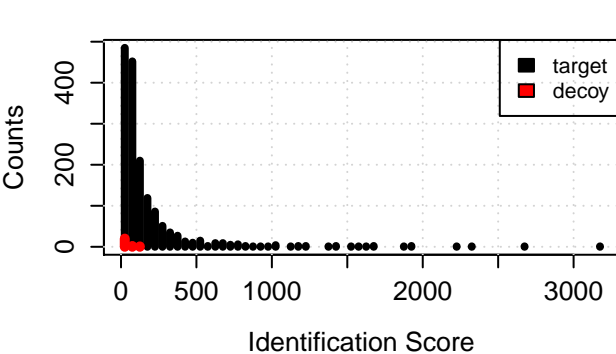
**Peptide Level**



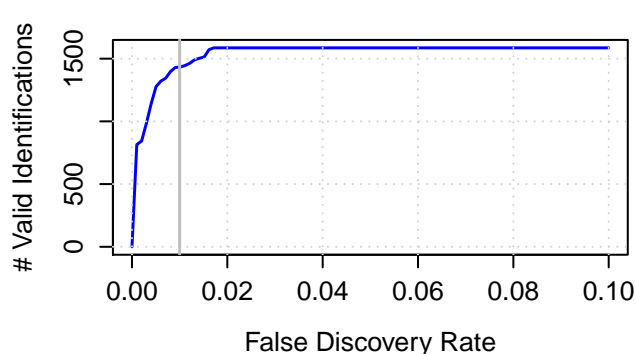
**Peptide Level**



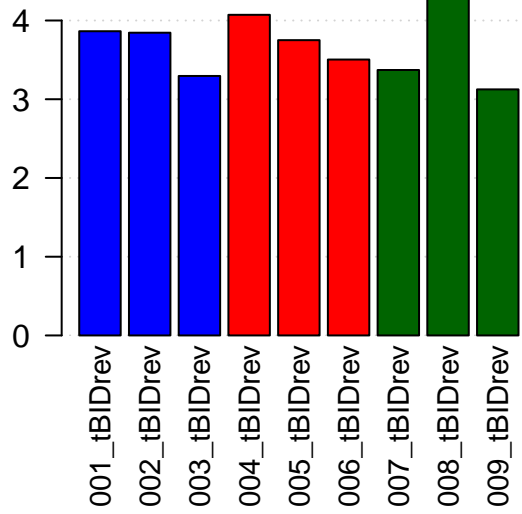
**Protein Level**



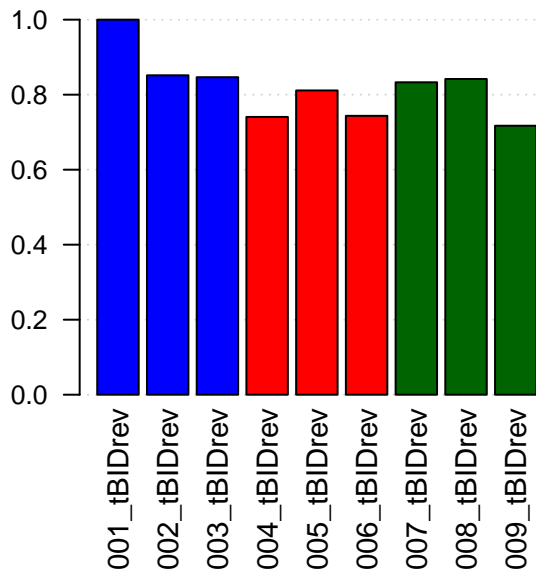
**Protein Level**



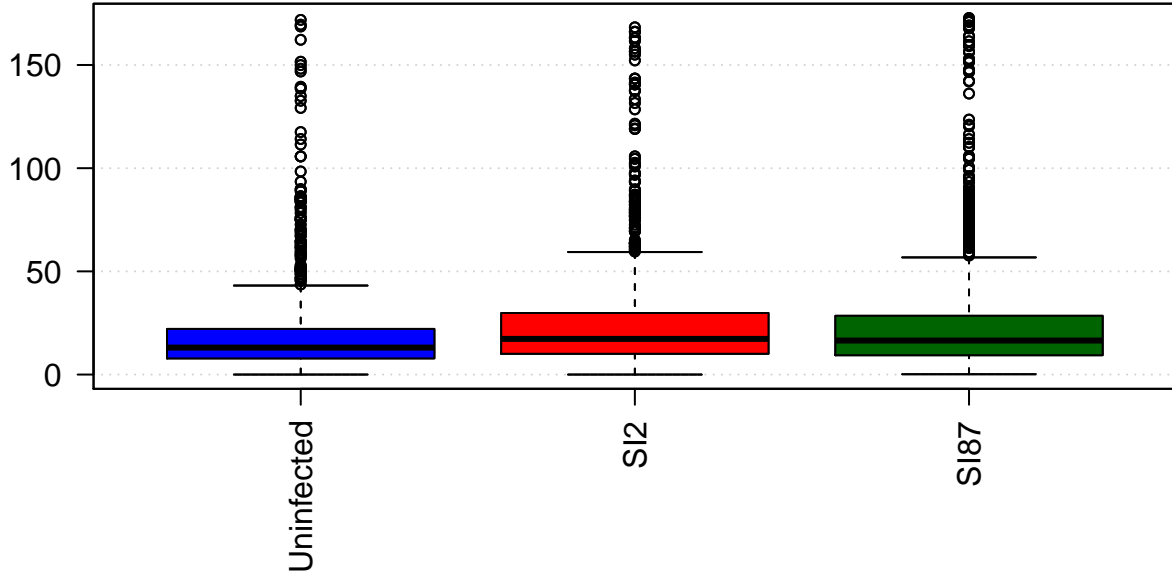
% Missing Values

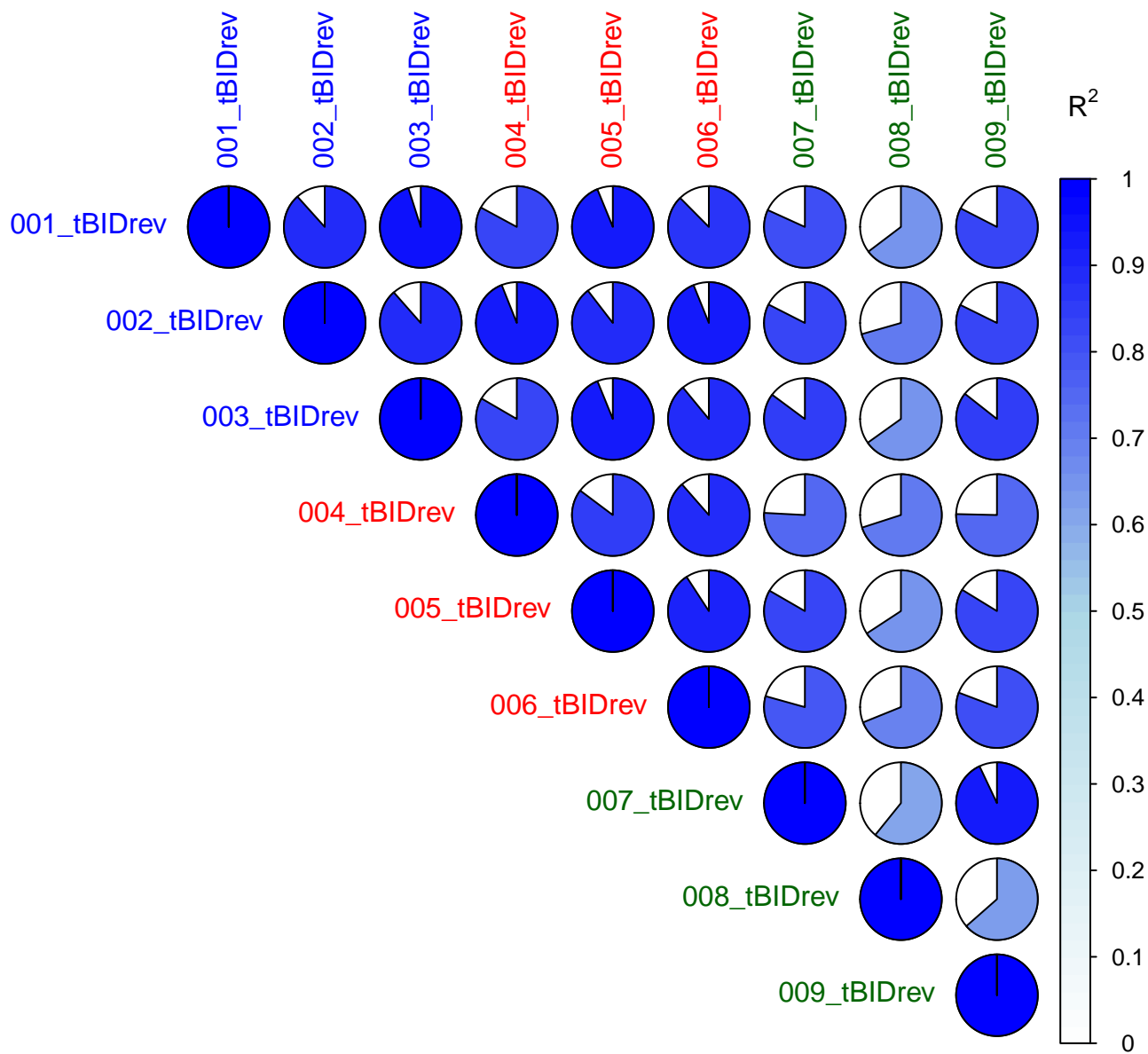


Summed MS-Signal (Scaled)

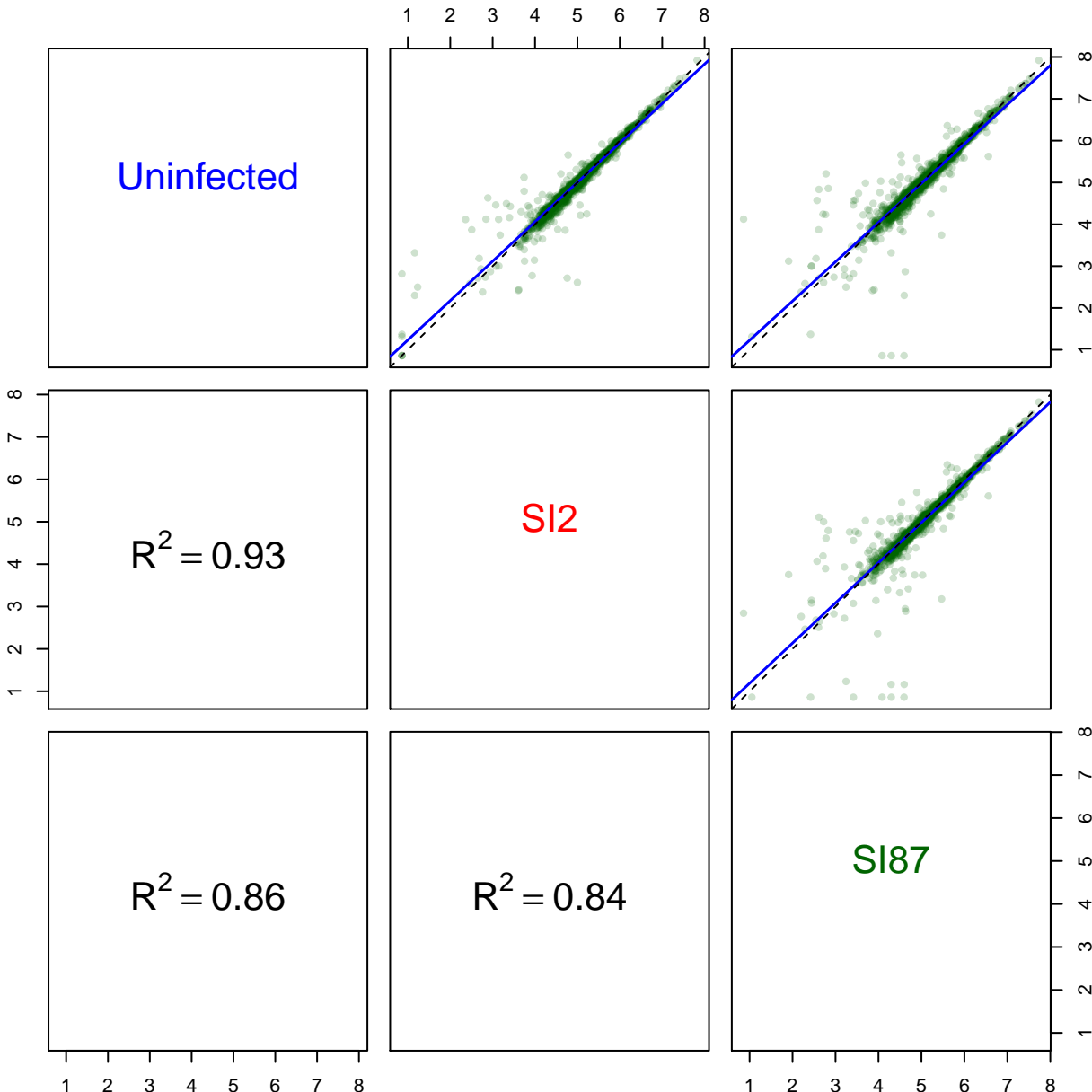


C.V. (%)



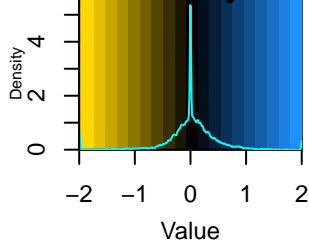


■ Uninfected  
■ SI2  
■ SI87

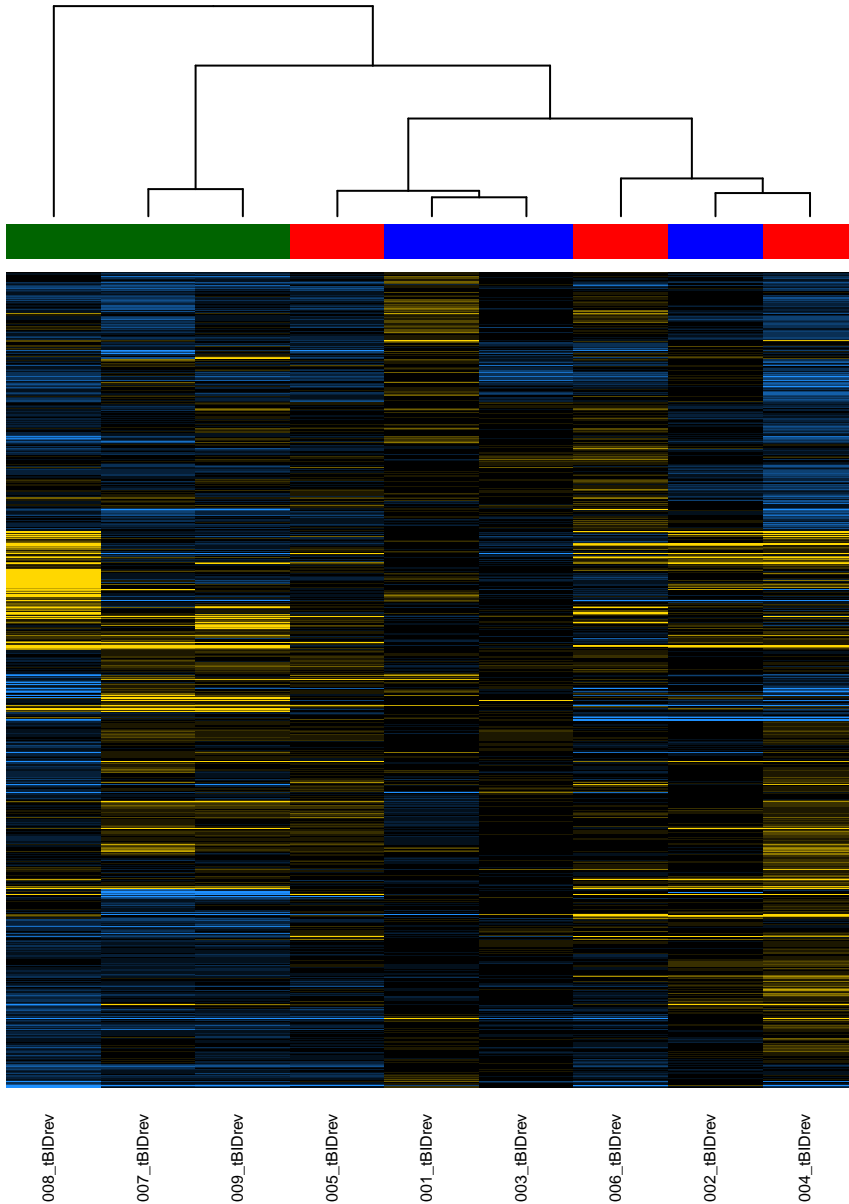


# Color Key

## and Density Plot

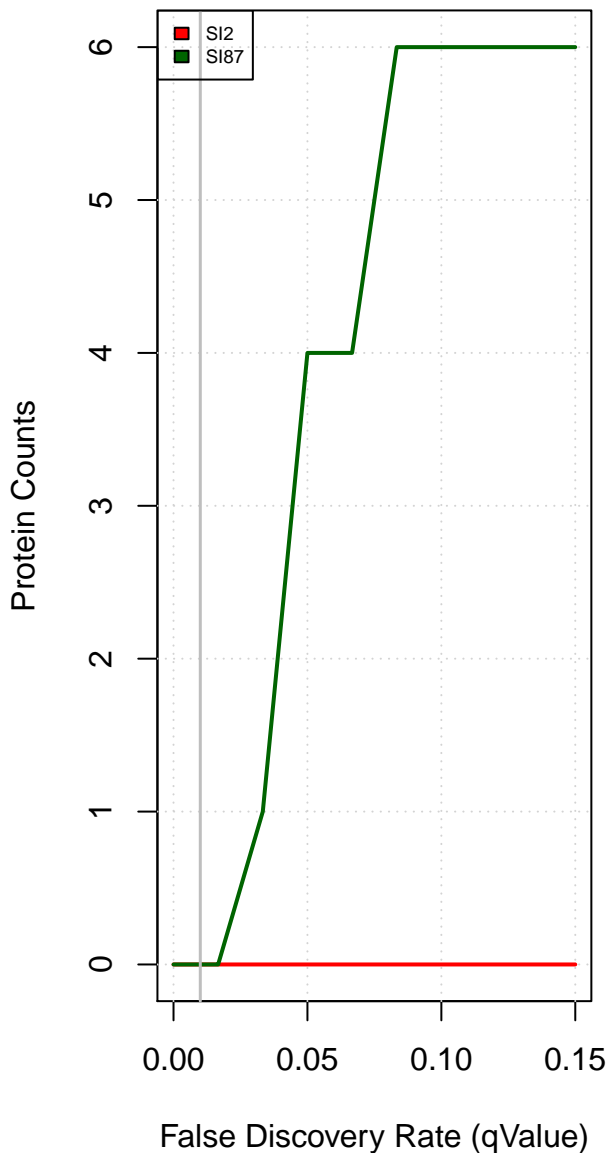


# Protein Level

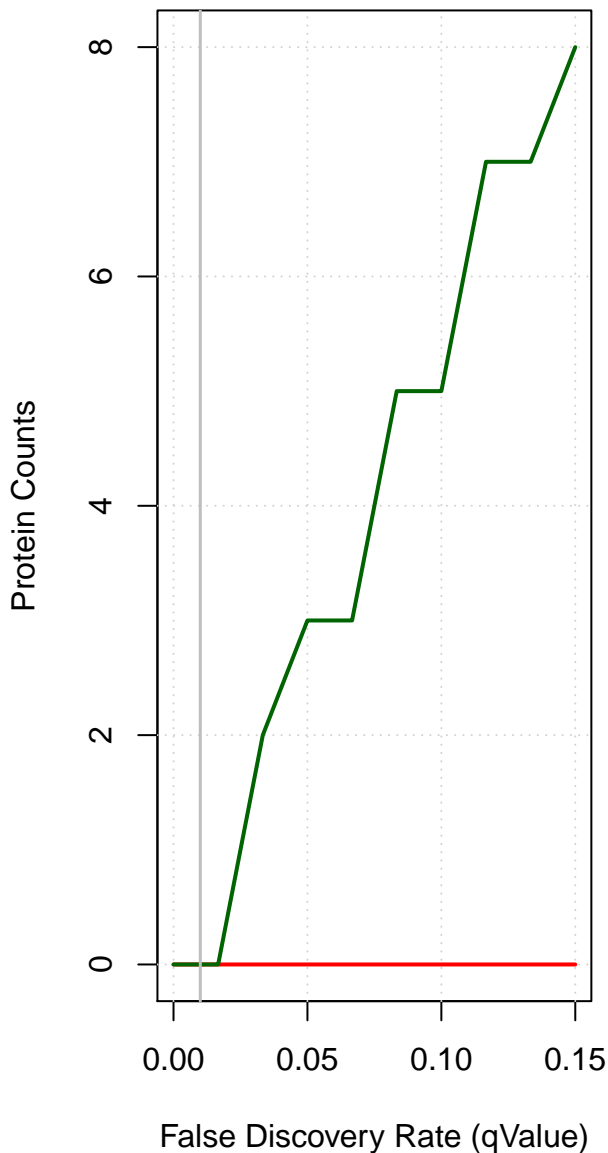




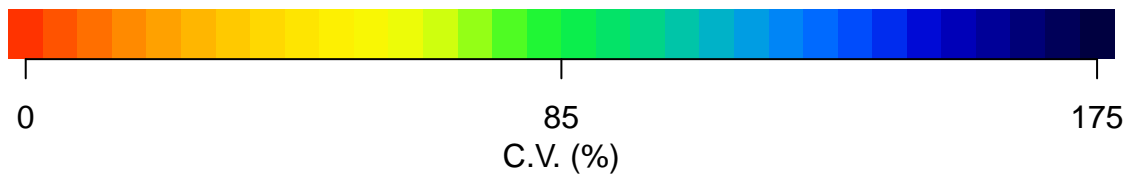
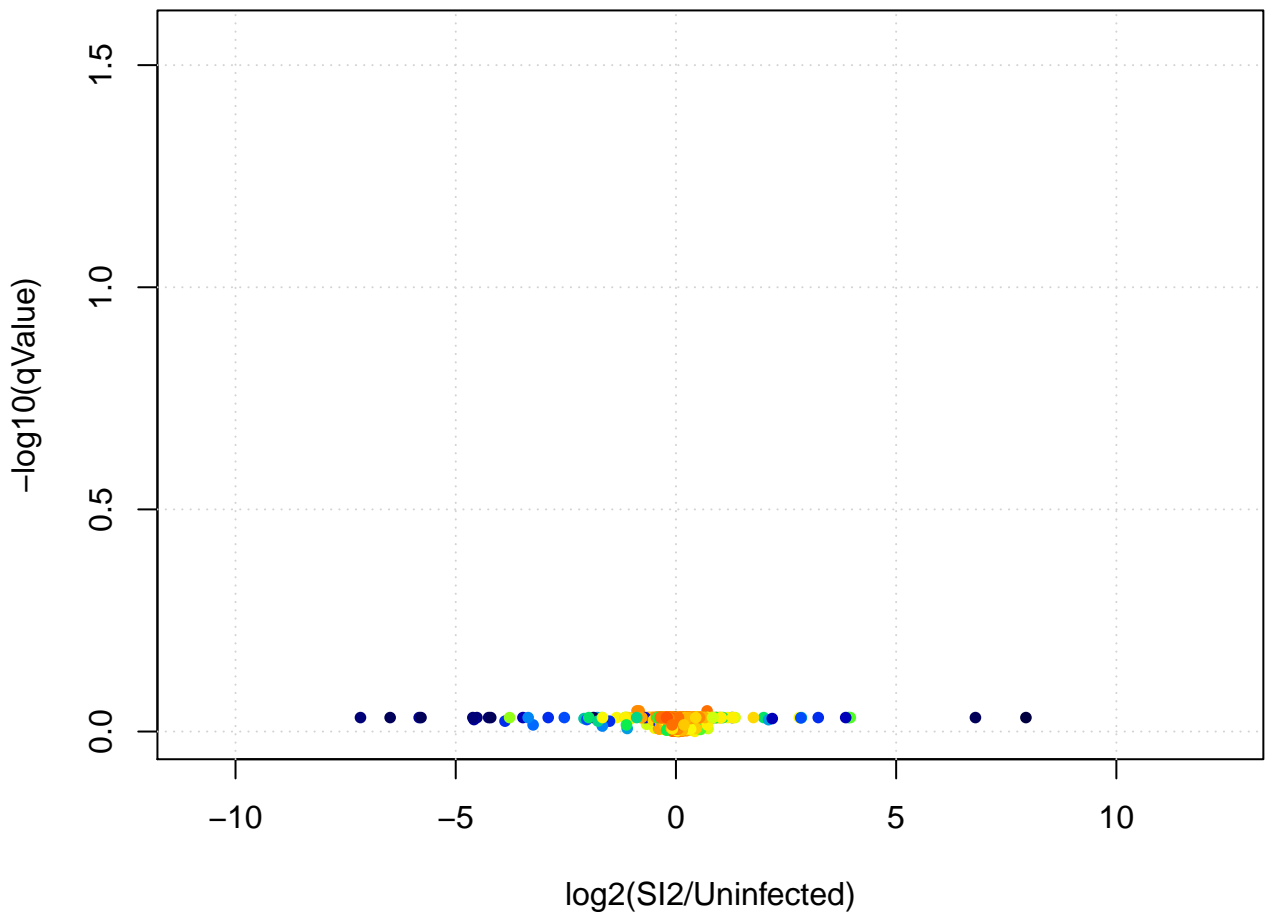
## DOWN REGULATION



## UP REGULATION



# Protein Level



# Protein Level

