

Experimental Design

SI87	9: 20150507–163002_009_tBIDrev
	8: 20150507–163002_008_tBIDrev
	7: 20150507–163002_007_tBIDrev
SI2	6: 20150507–163002_006_tBIDrev
	5: 20150507–163002_005_tBIDrev
	4: 20150507–163002_004_tBIDrev
<u>Uninfected</u>	3: 20150507–163002_003_tBIDrev
	2: 20150507–163002_002_tBIDrev
	1: 20150507–163002_001_tBIDrev

IDENTIFICATIONS OVERVIEW (FDR 0.01)

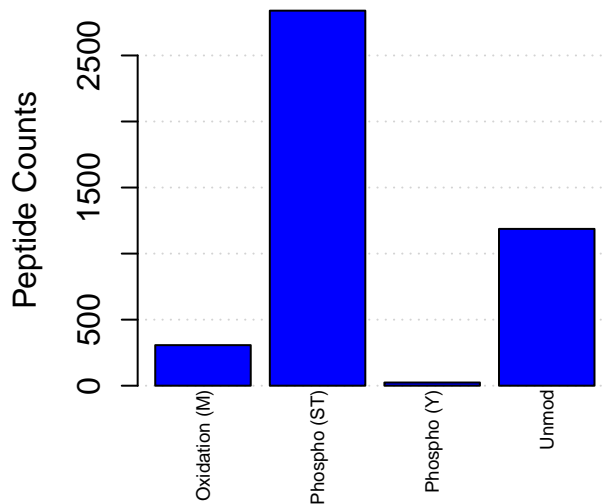
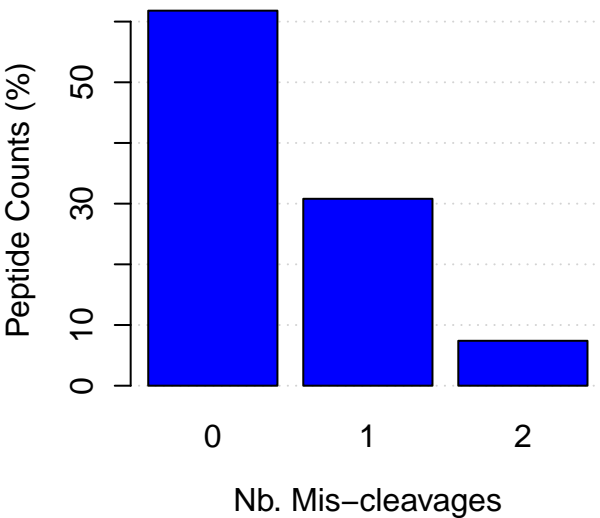
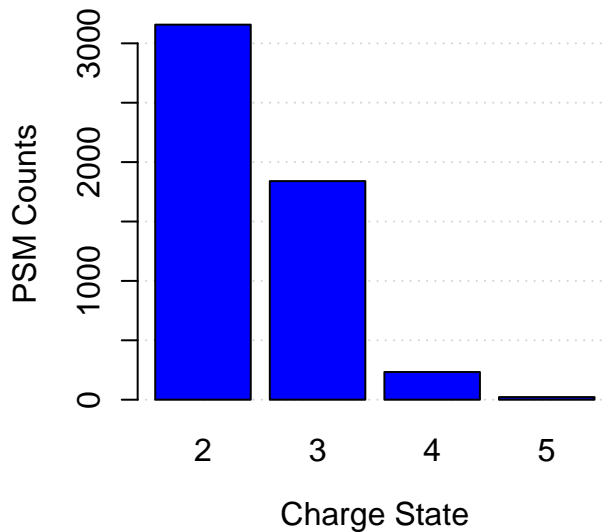
5252 PSM

3909 PEPTIDES

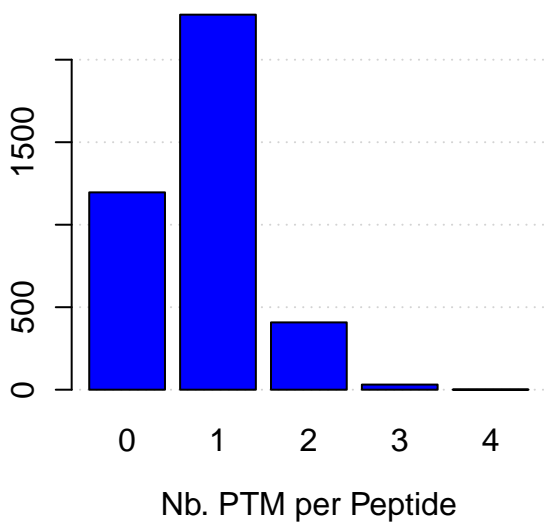
1196 UN-MOD.

2713 MOD.

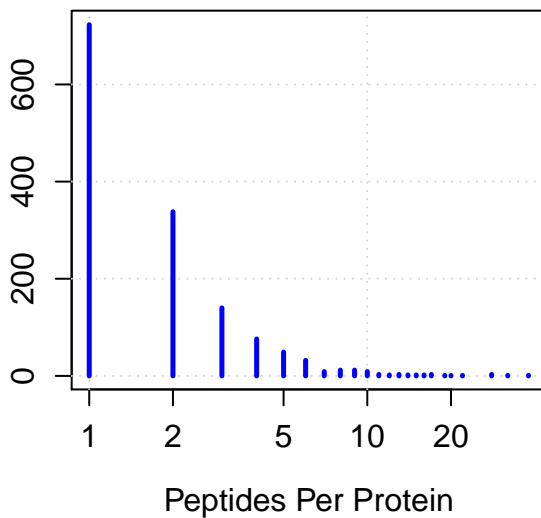
1425 PROTEINS



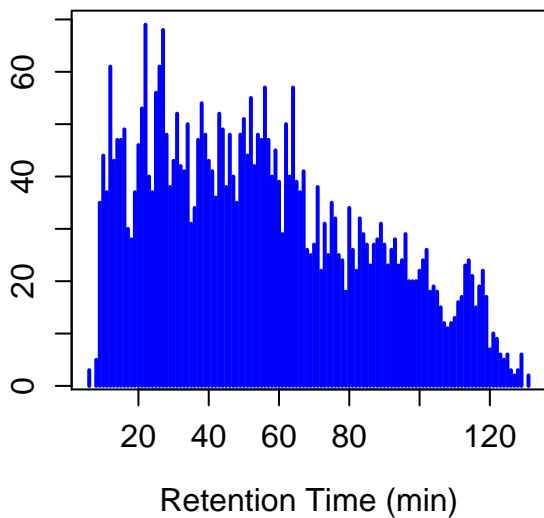
Peptide Counts



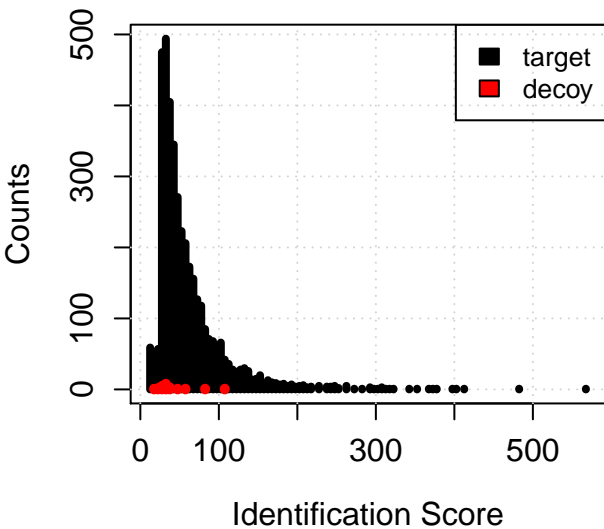
Protein Counts



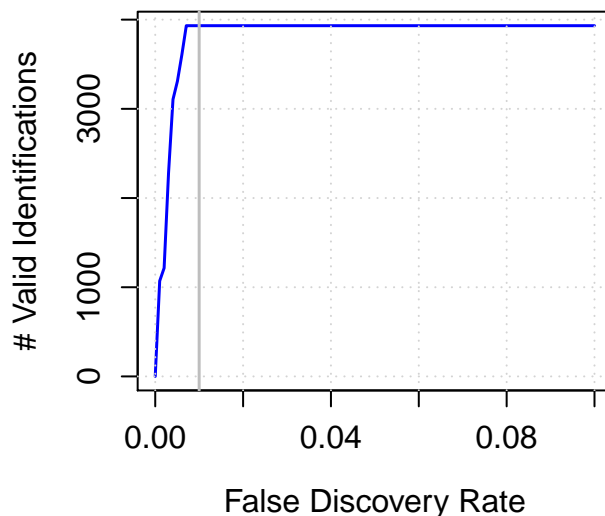
Identified Features



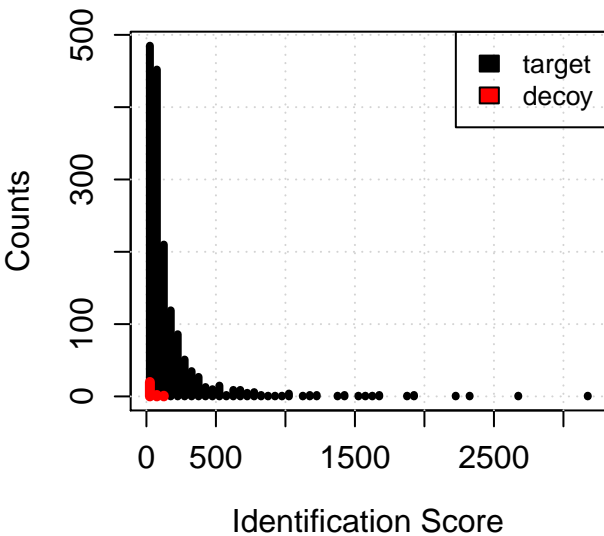
Peptide Level



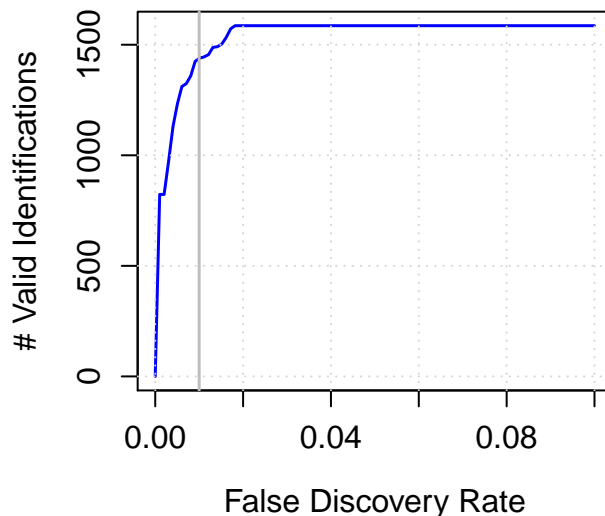
Peptide Level

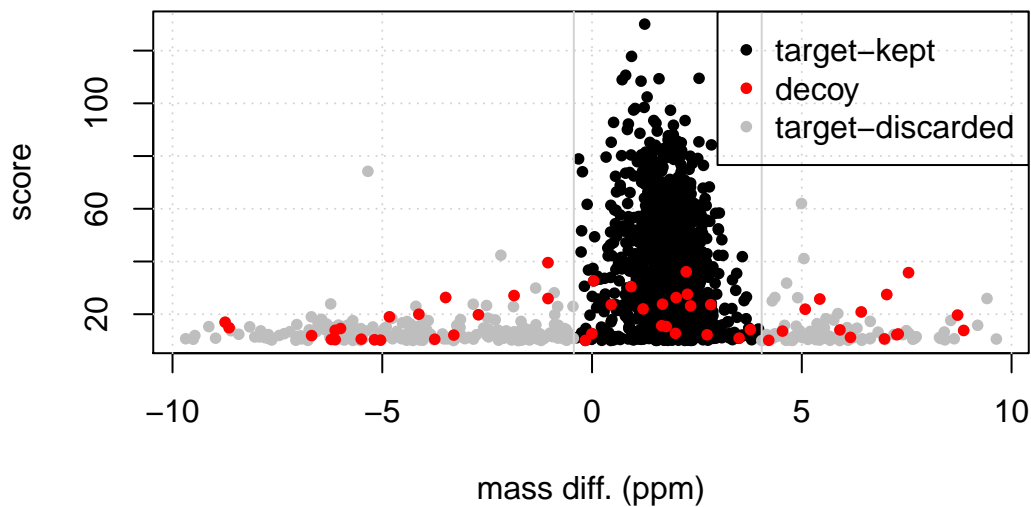
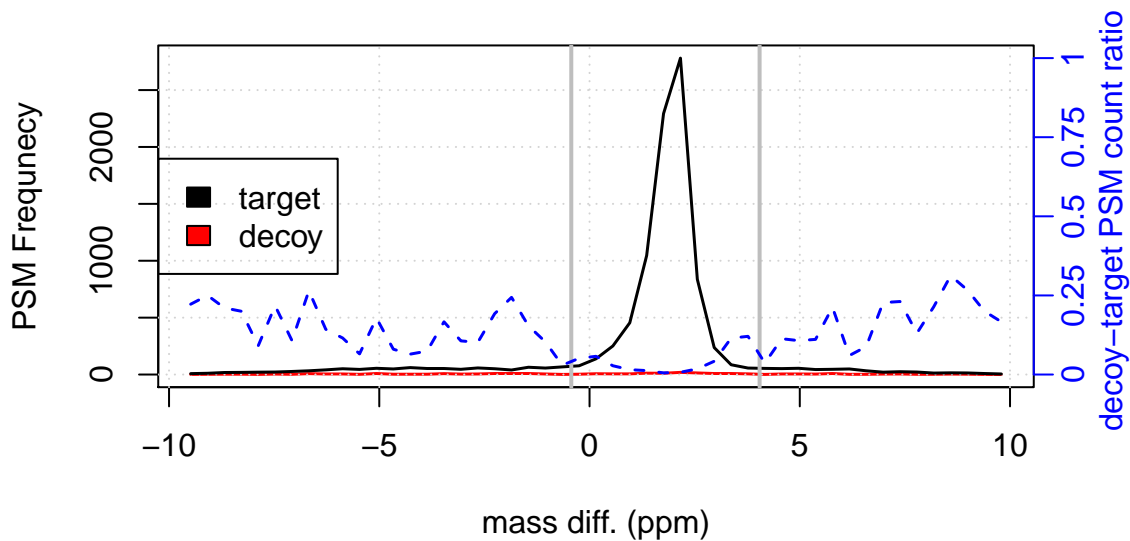


Protein Level



Protein Level

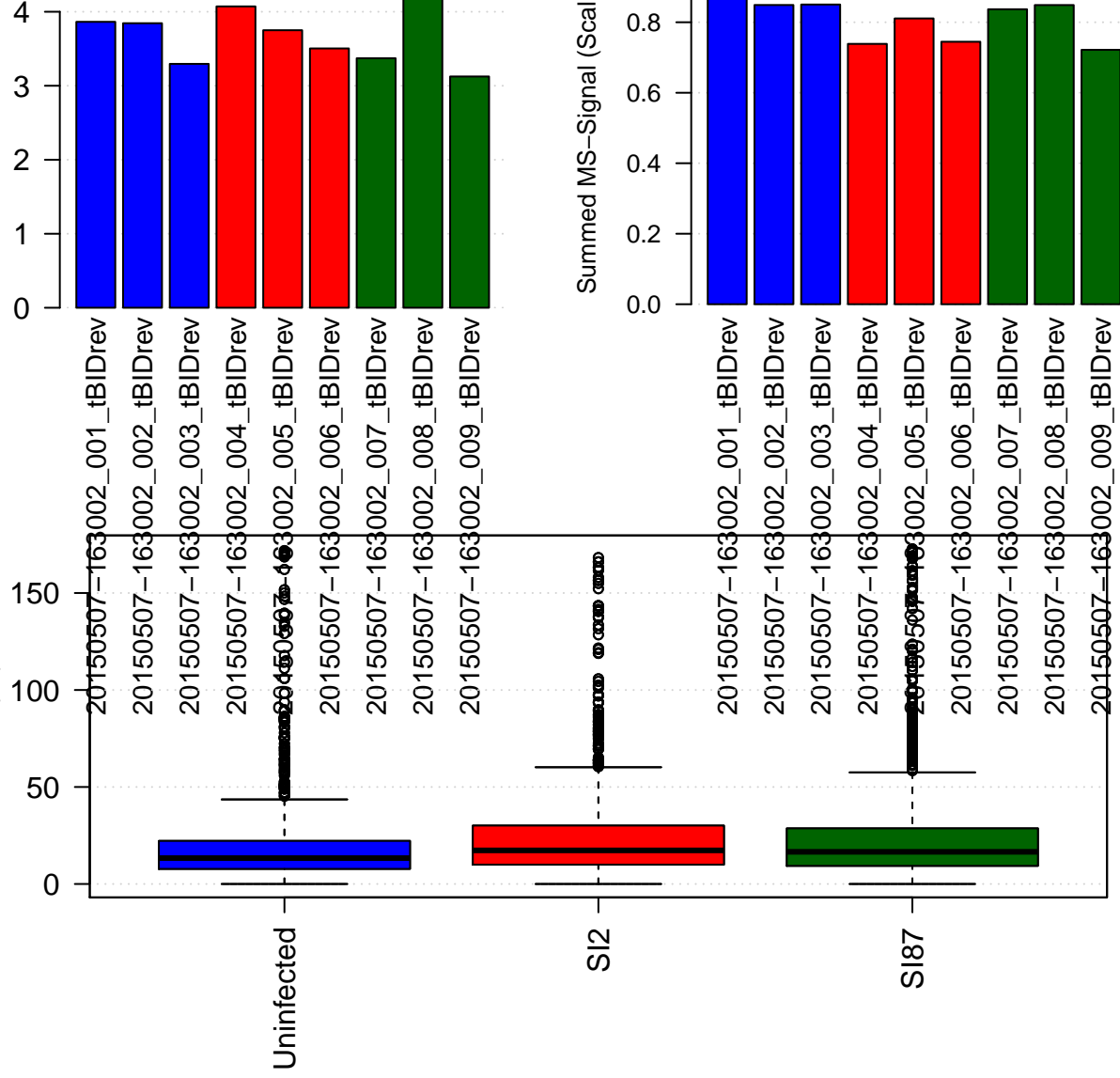


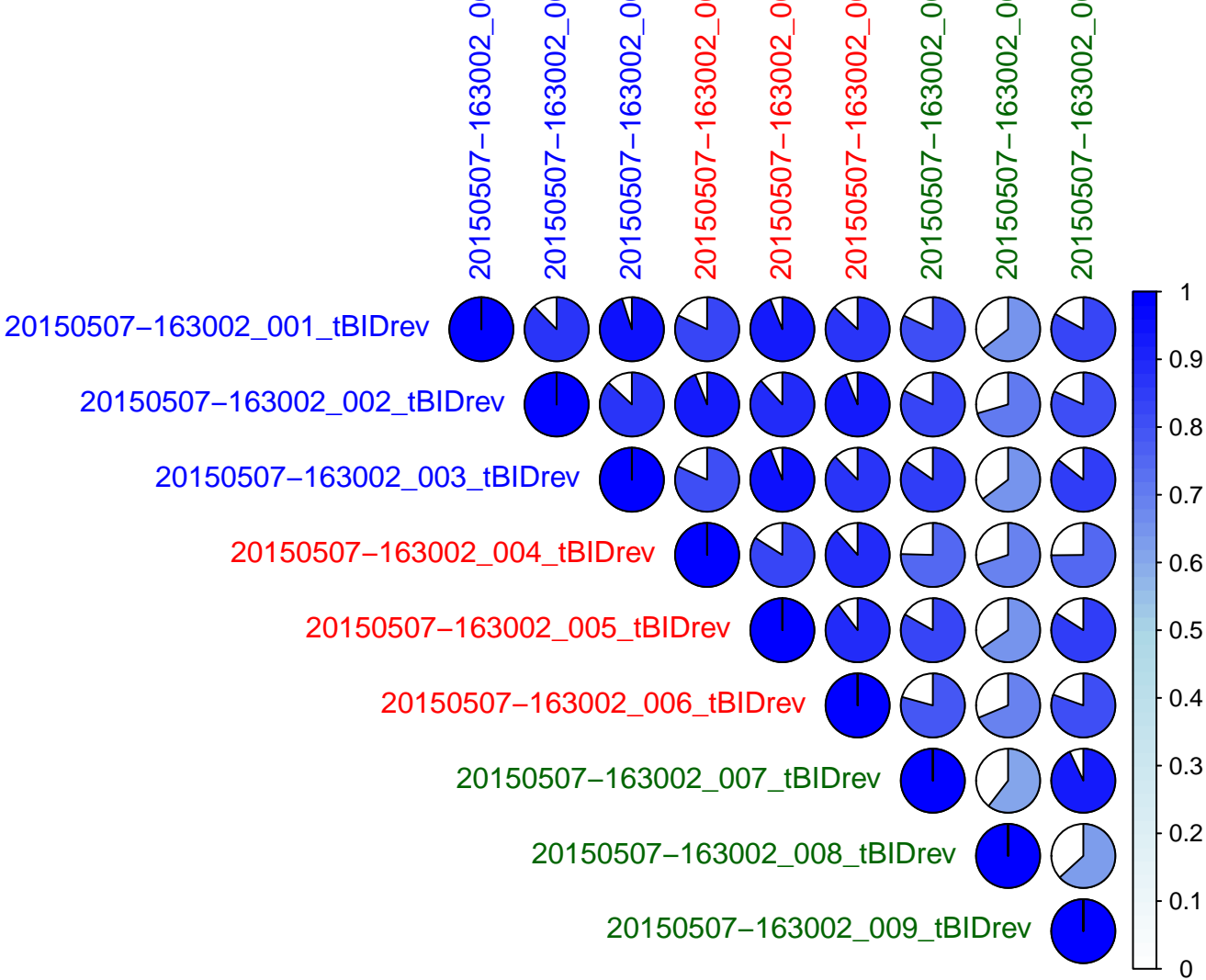


% Missing Values

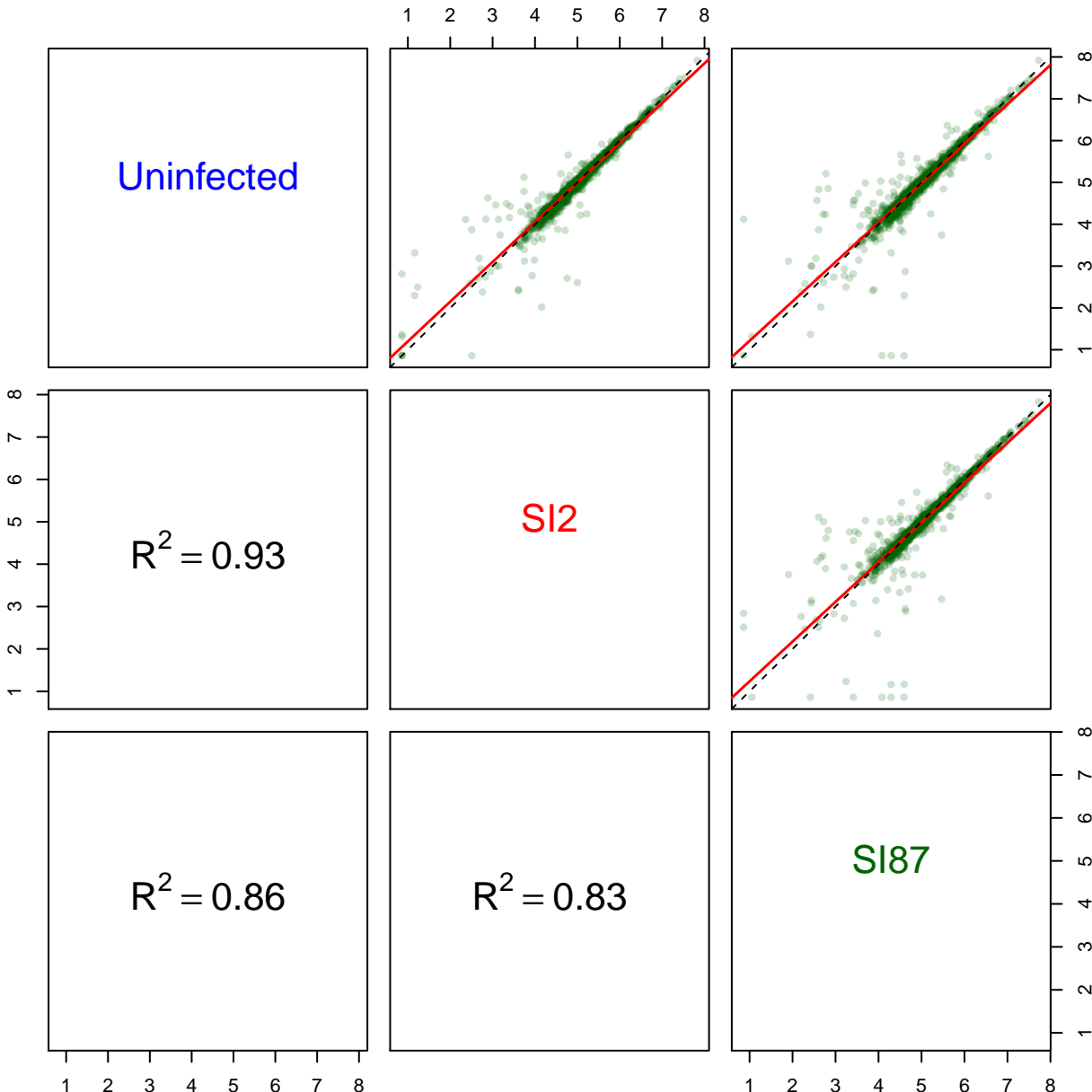
Summed MS-Signal (Scaled)

C.V. (%)



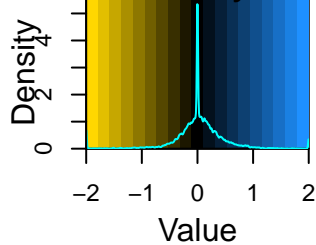


■ Uninfected
■ SI2
■ SI87

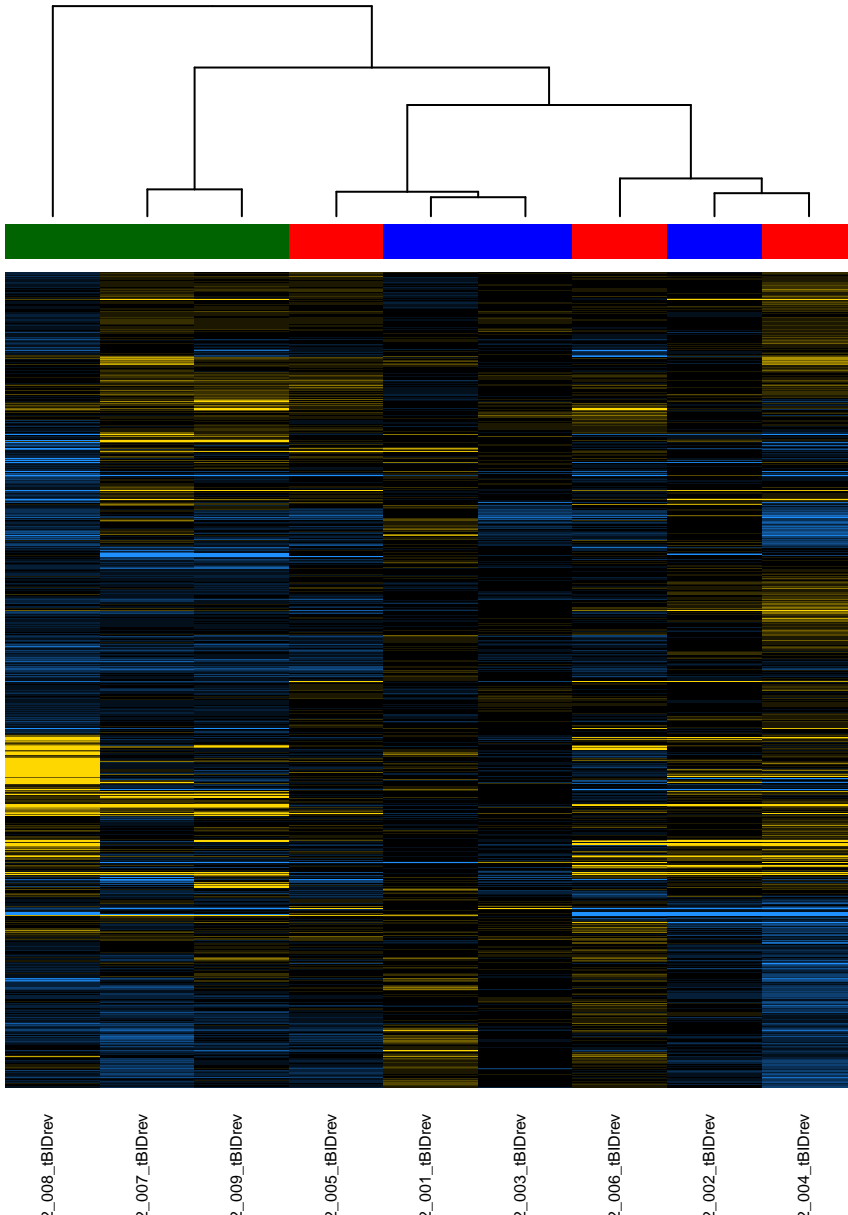


Color Key

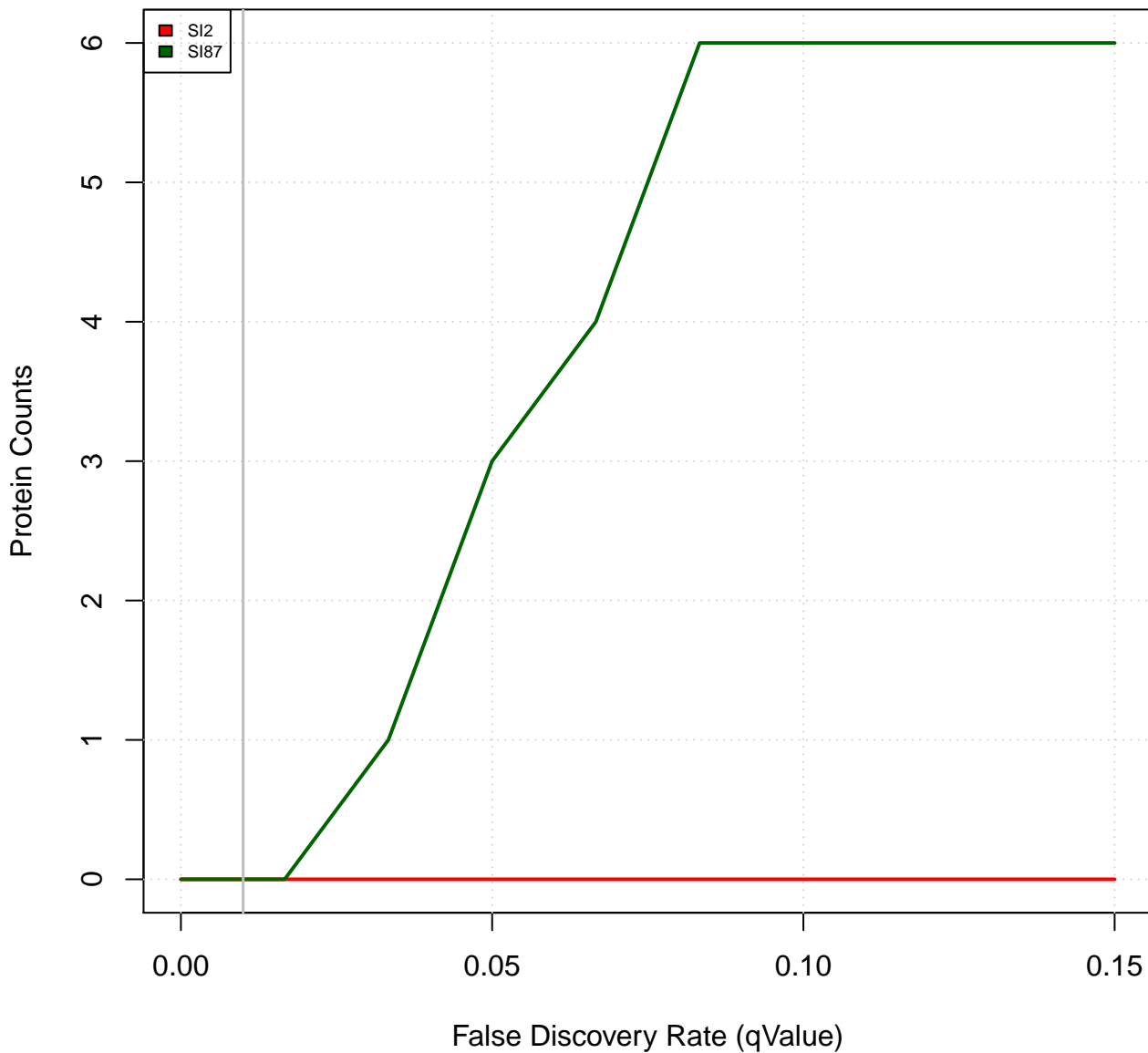
and Density Plot



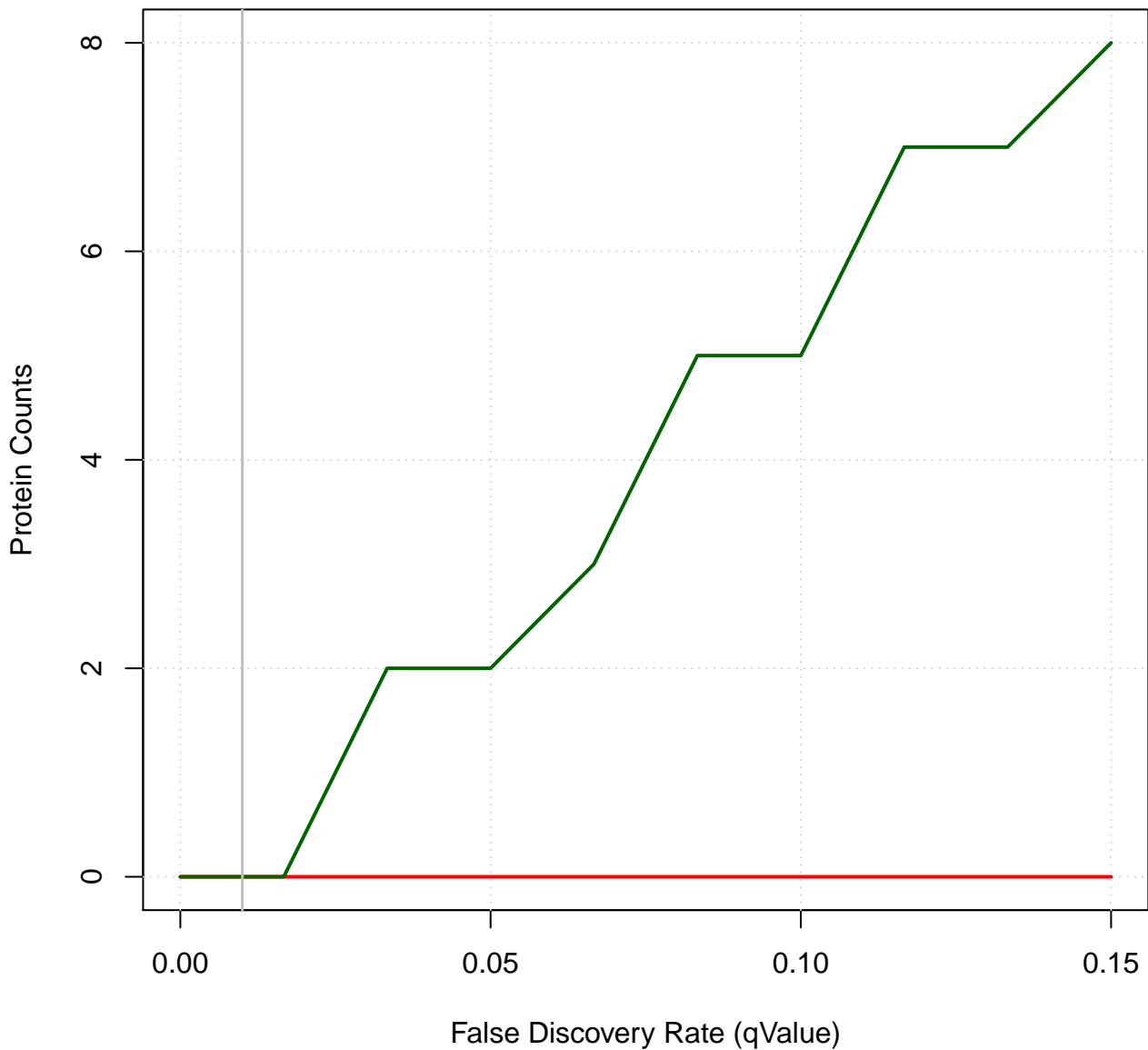
Protein Level



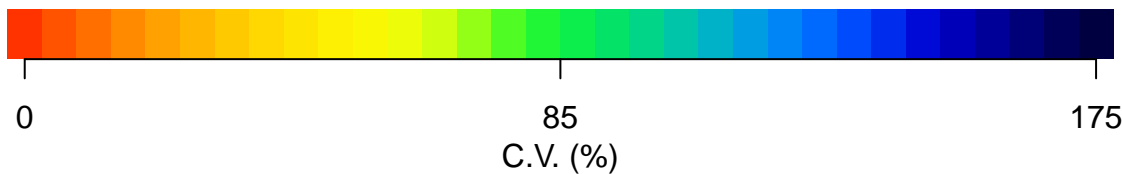
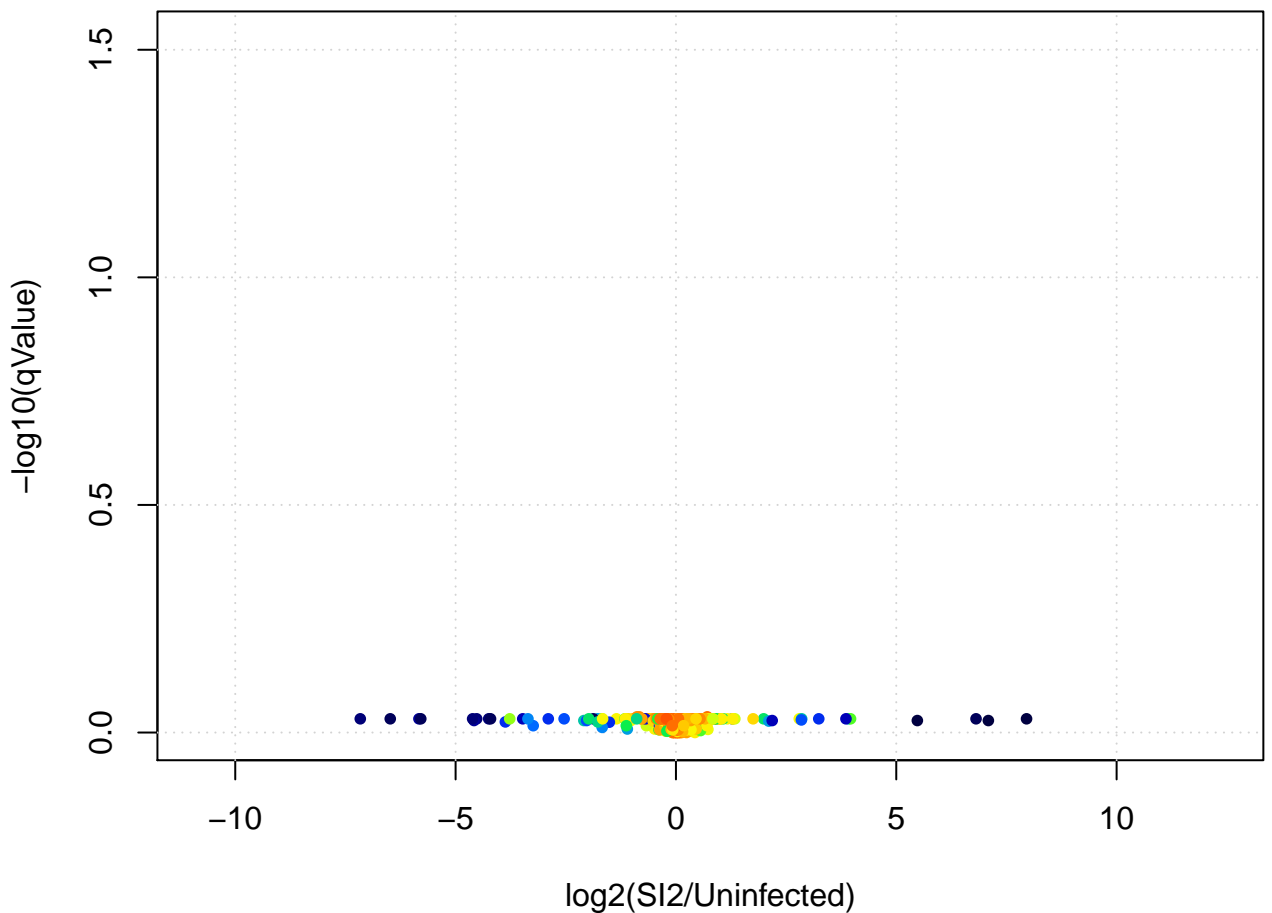
DOWN REGULATION



UP REGULATION



Protein Level



Protein Level

