

# Experimental Design

Doc_TSC	8: C15-05098
	7: C15-05097
Doc_Ctrl	6: C15-05095
	5: C15-05094
Urea_TSC	4: C15-05092
	3: C15-05091
<u>Urea_Ctrl</u>	2: C15-05089
	1: C15-05088

# IDENTIFICATIONS OVERVIEW (FDR 0.01)

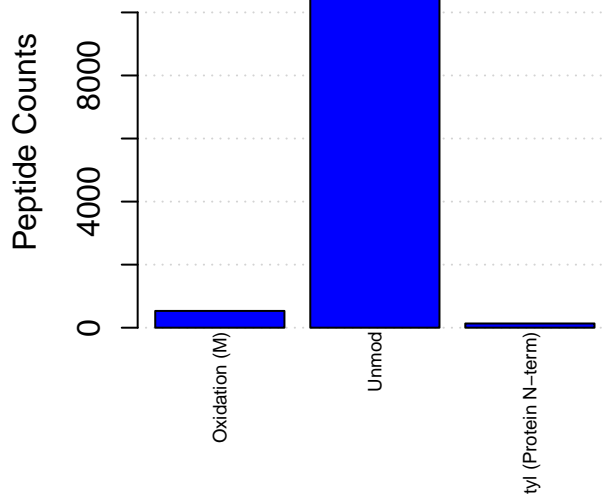
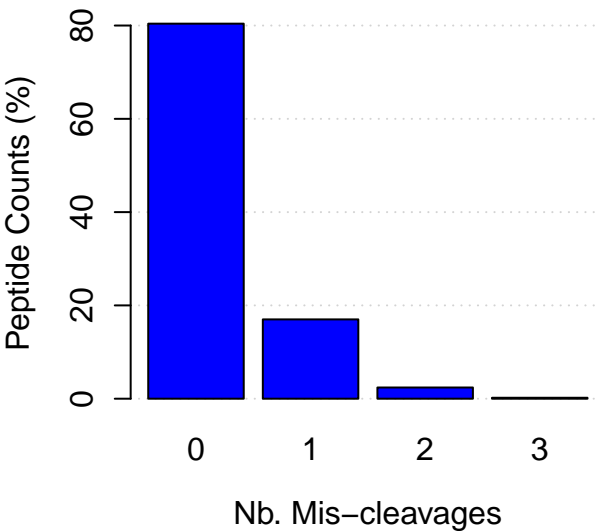
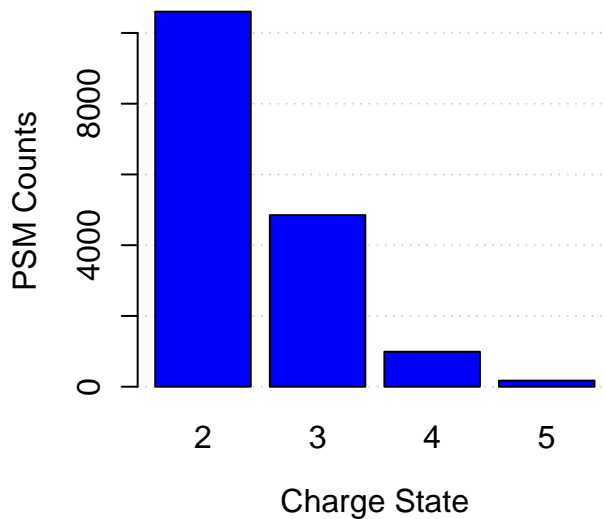
16611 PSM

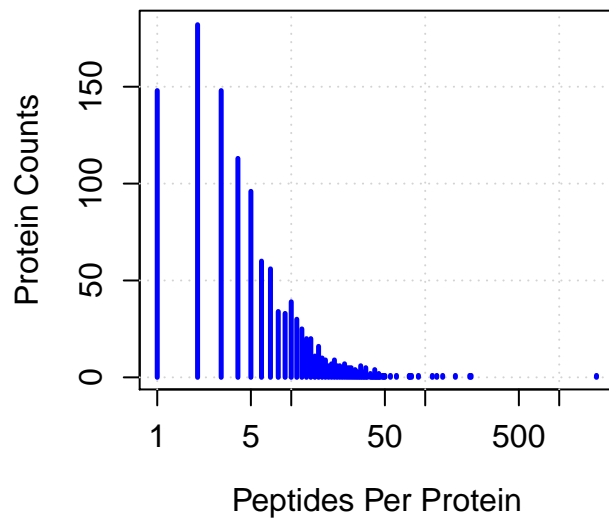
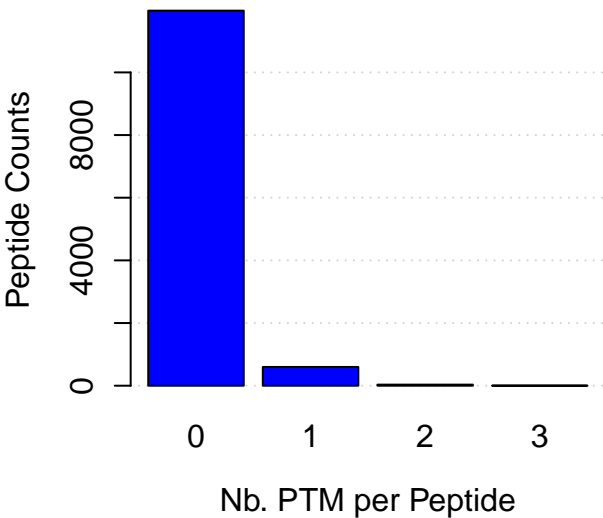
12604 PEPTIDES

11970 UN-MOD.

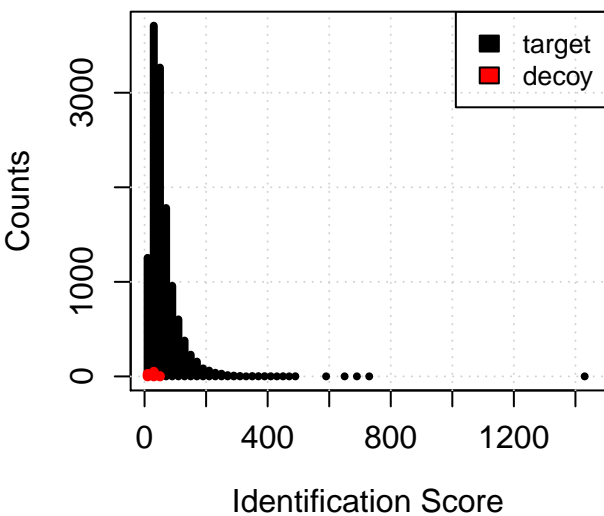
634 MOD.

1169 PROTEINS

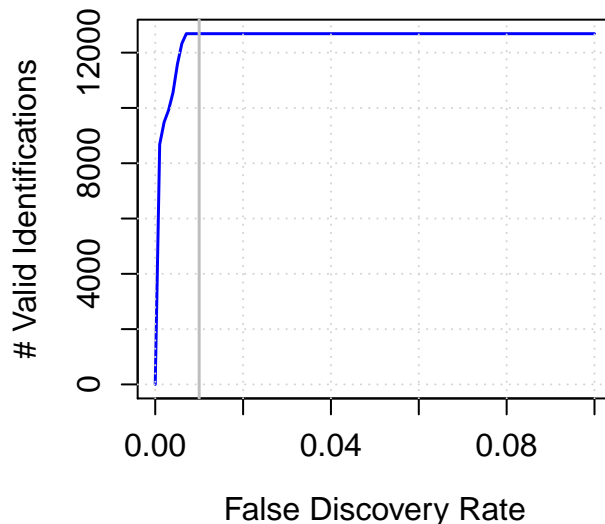




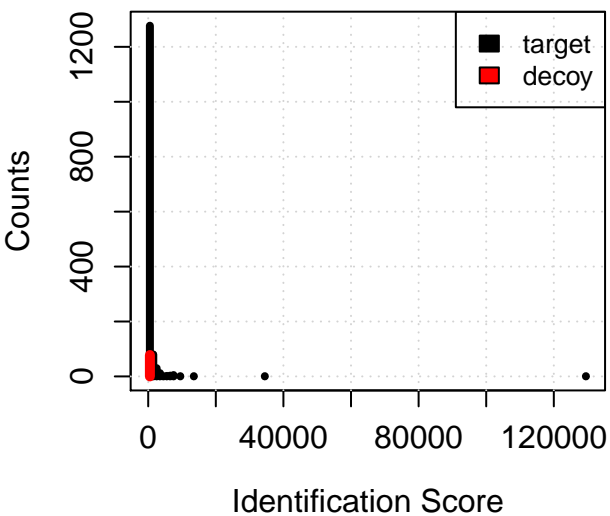
**Peptide Level**



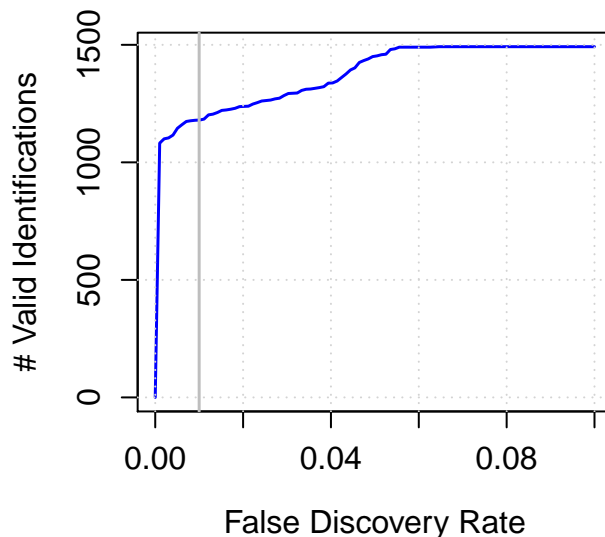
**Peptide Level**

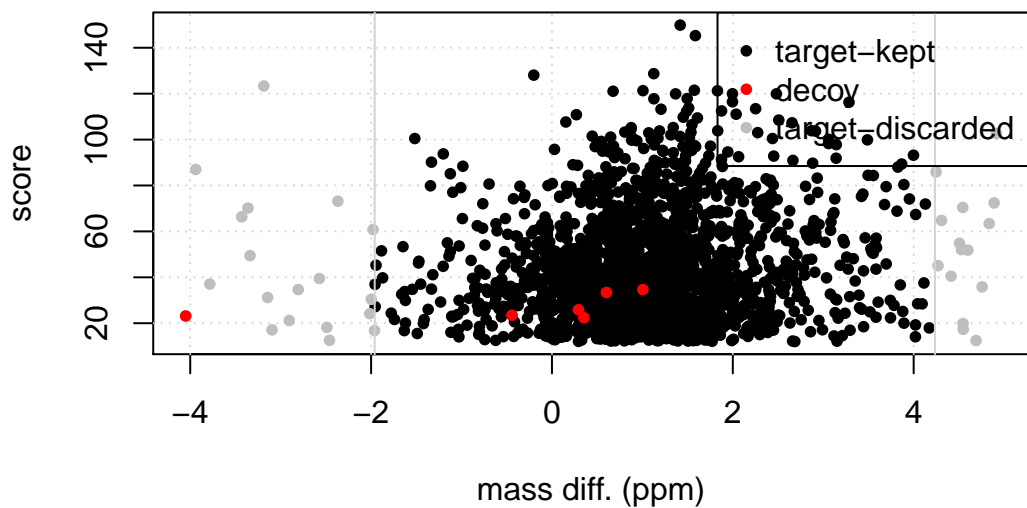
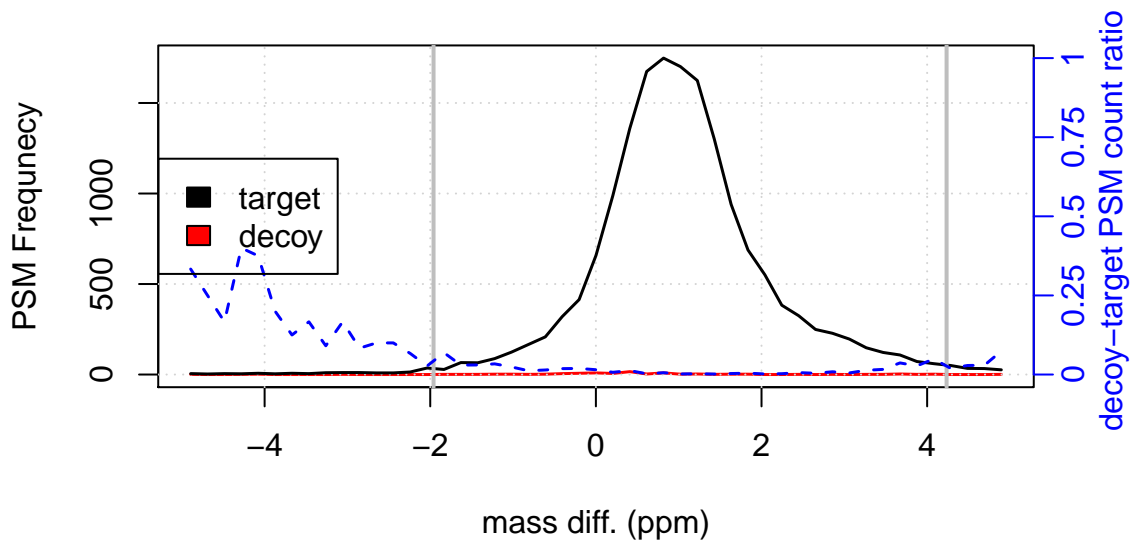


**Protein Level**

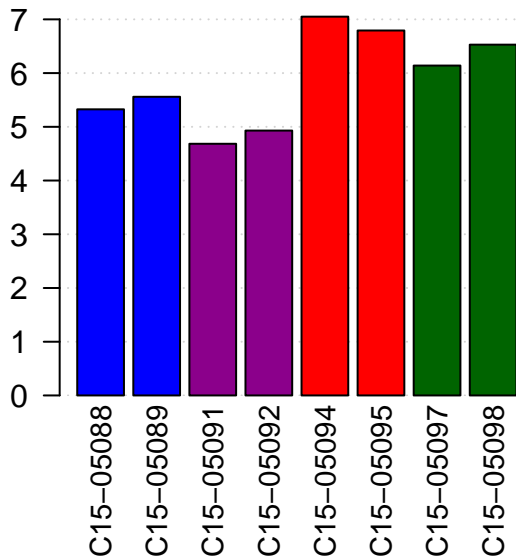


**Protein Level**

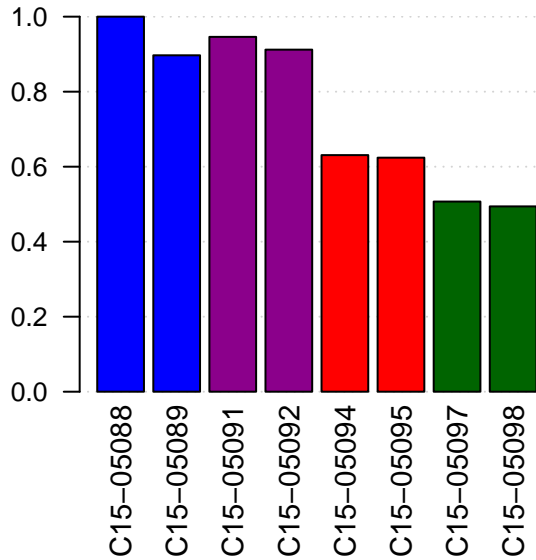




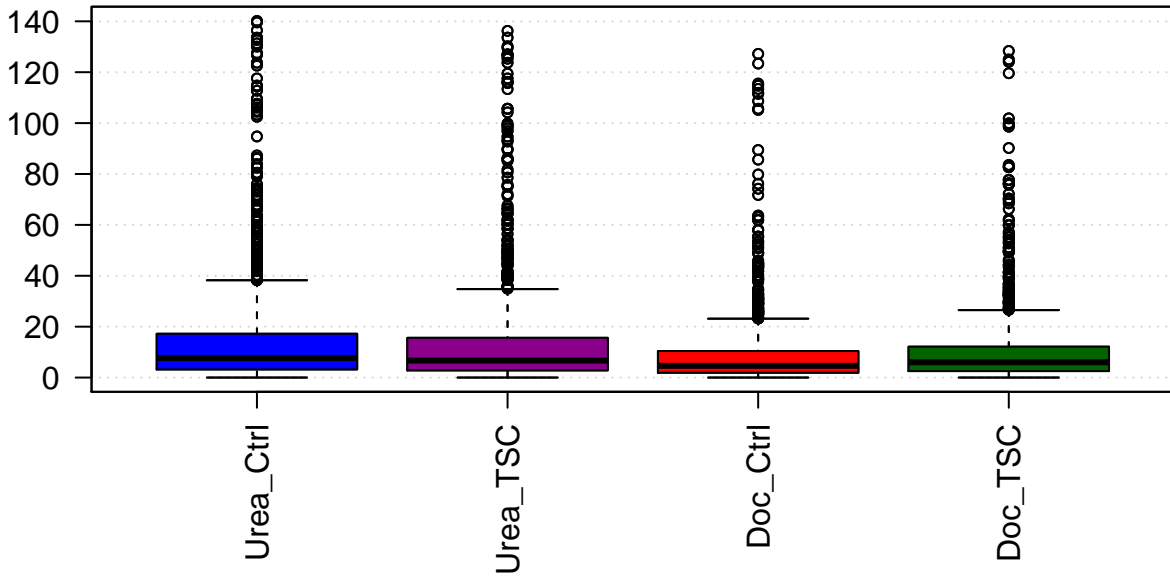
% Missing Values

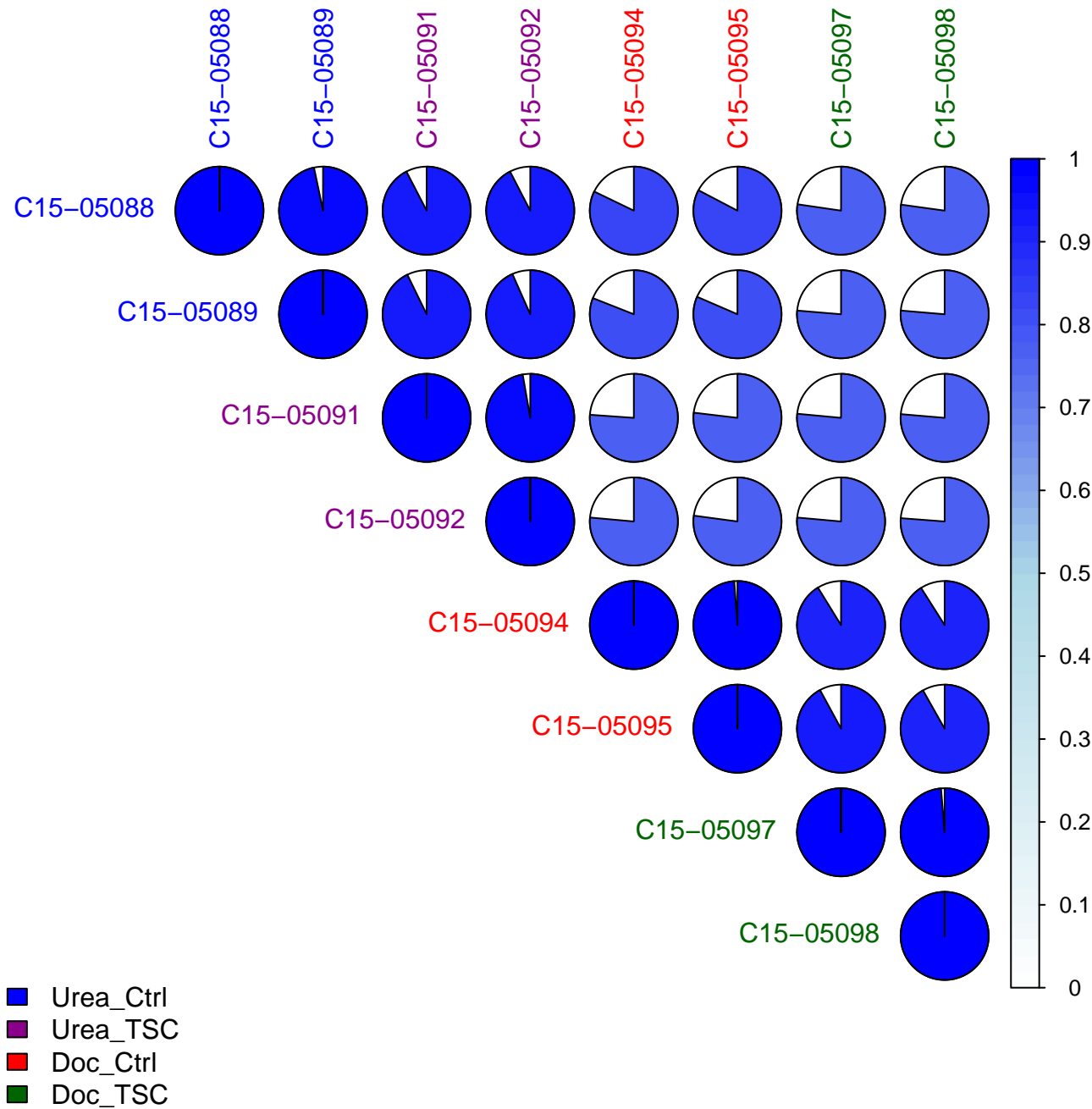


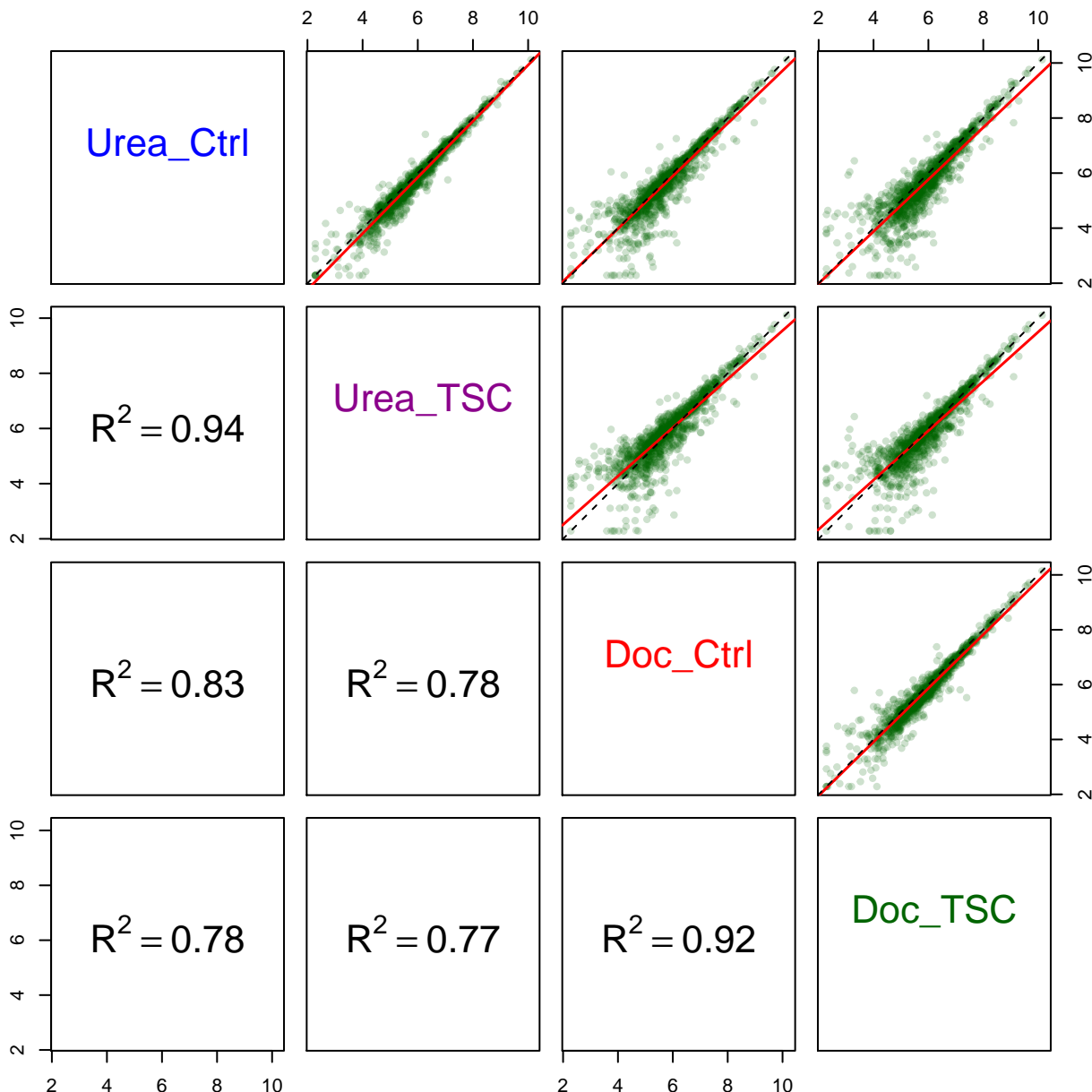
Summed MS-Signal (Scaled)



C.V. (%)



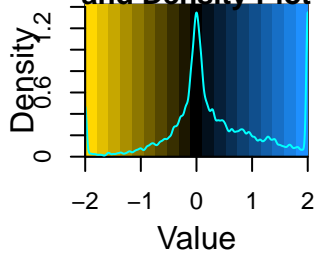






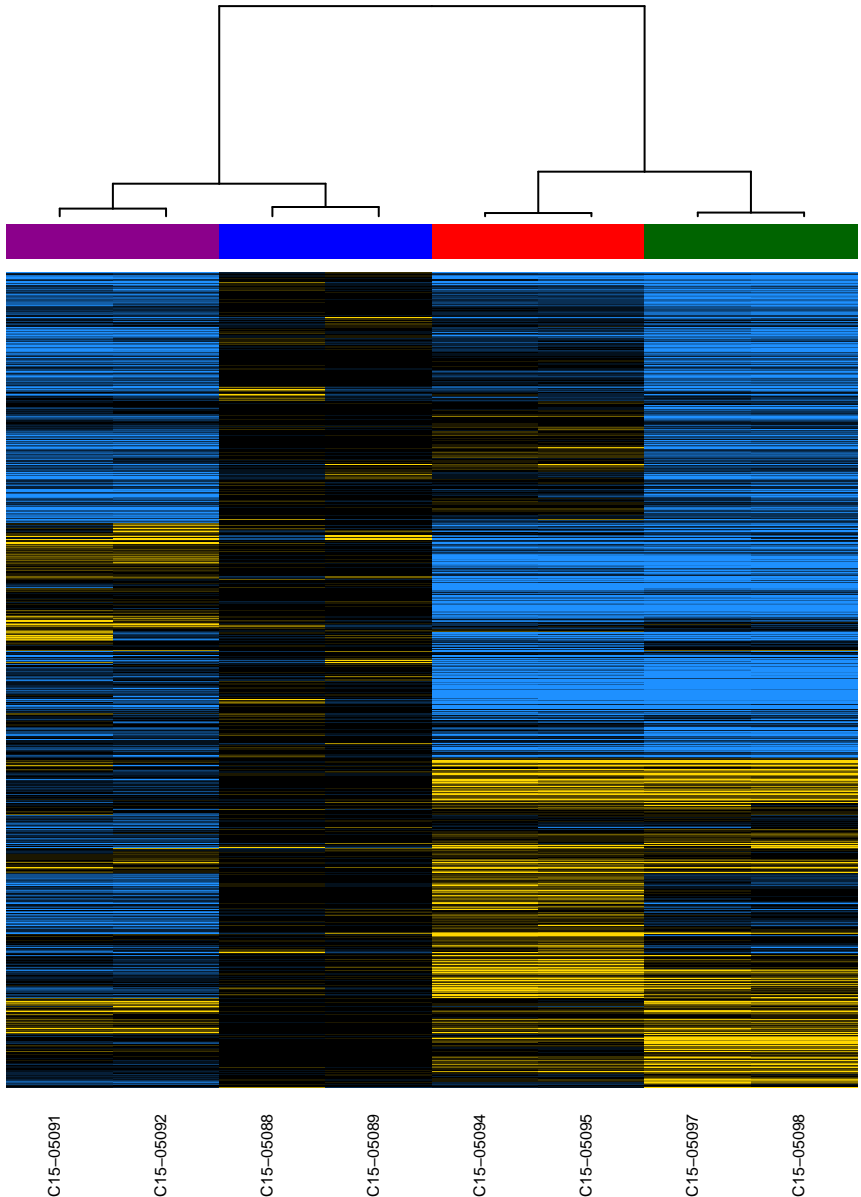
# Color Key

## and Density Plot

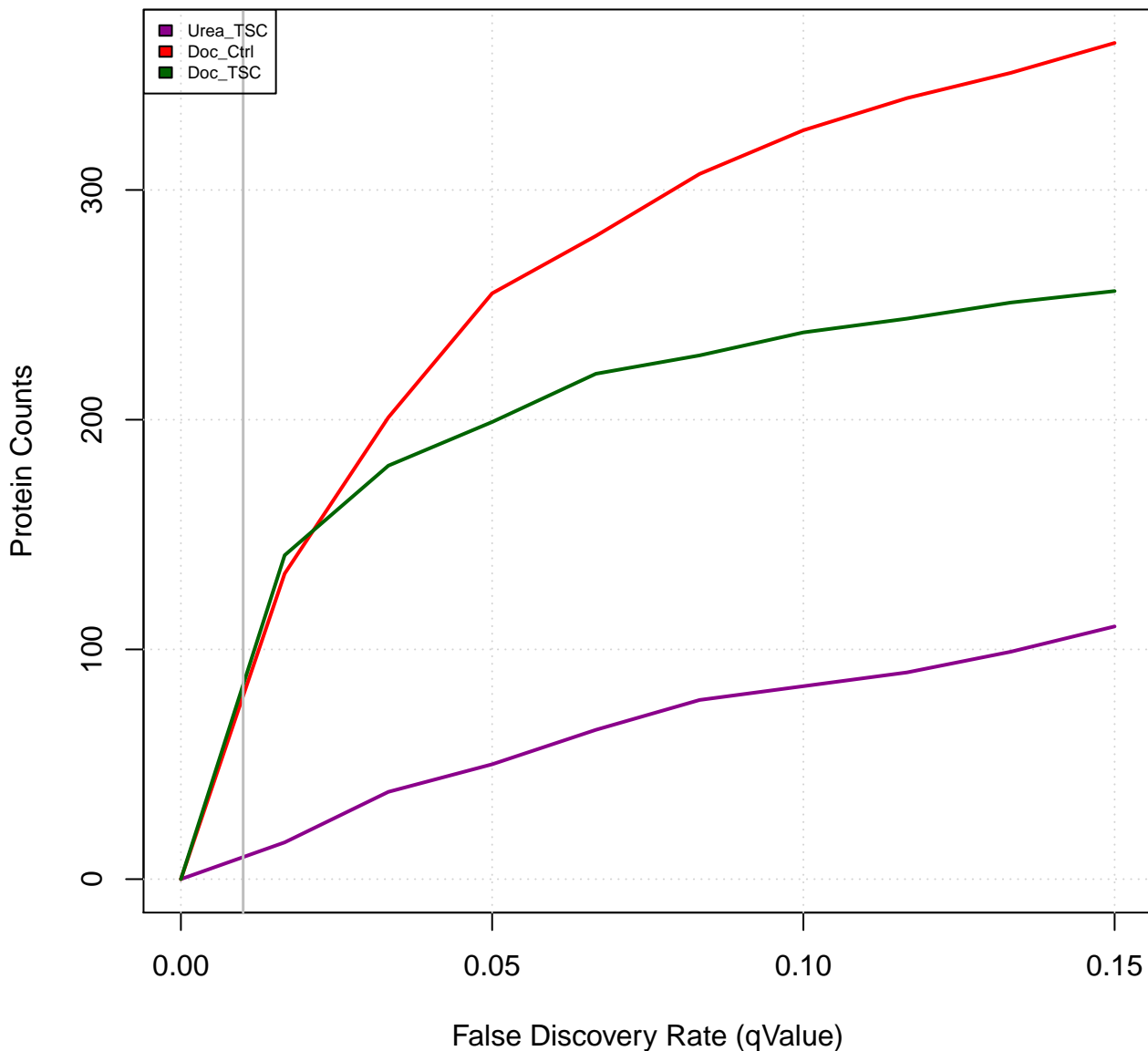


## Protein Level

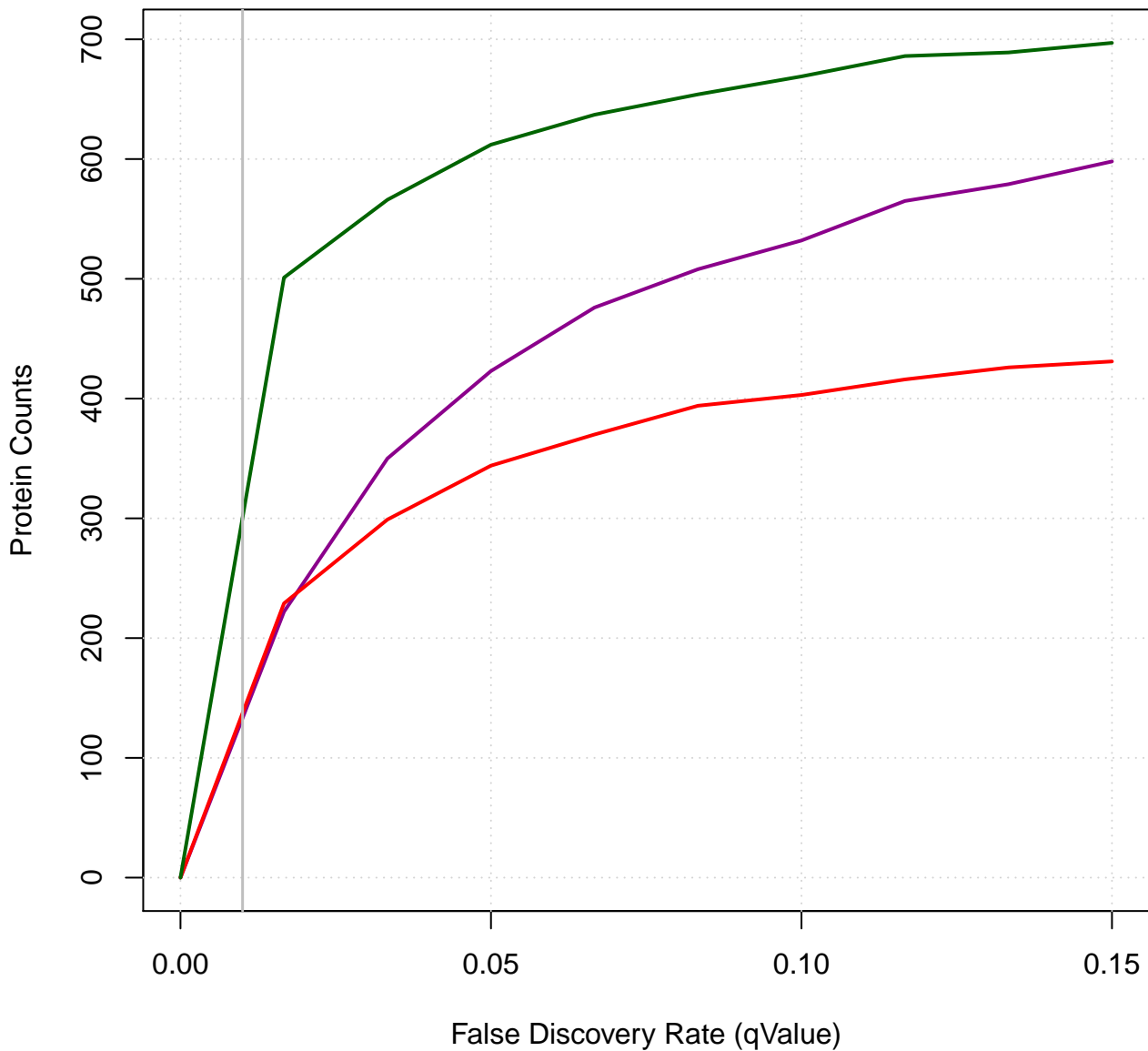
- Doc\_Ctrl
- Doc\_TSC
- Urea\_Ctrl
- Urea\_TSC



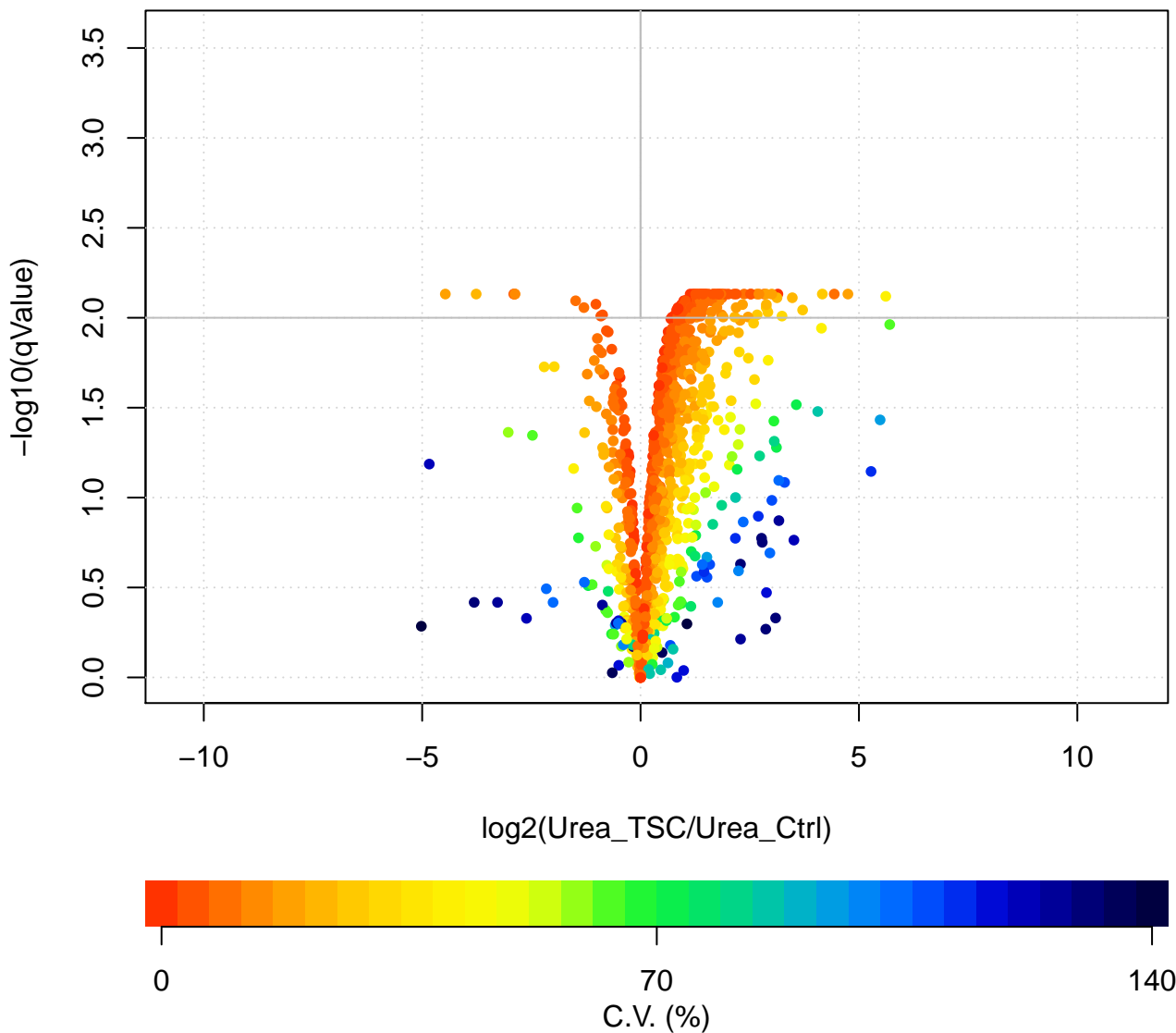
# DOWN REGULATION



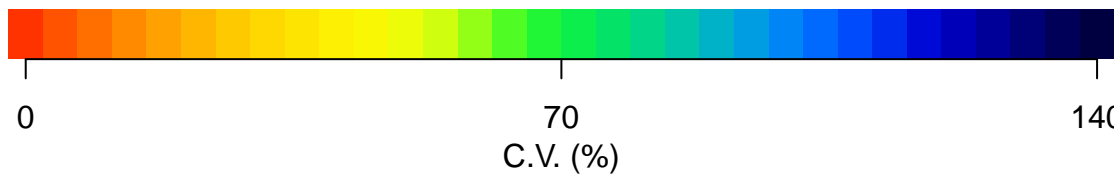
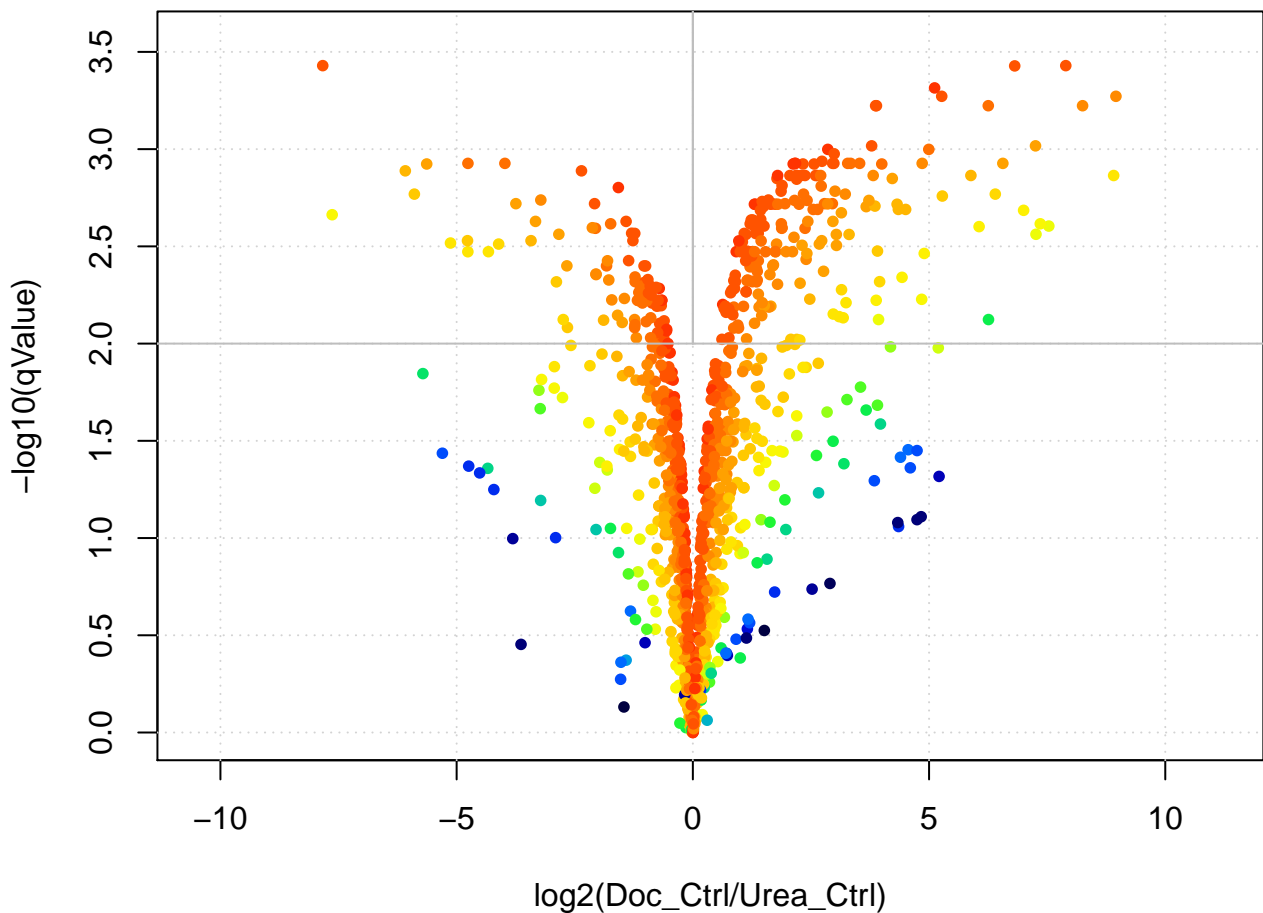
## UP REGULATION



# Protein Level



# Protein Level



# Protein Level

