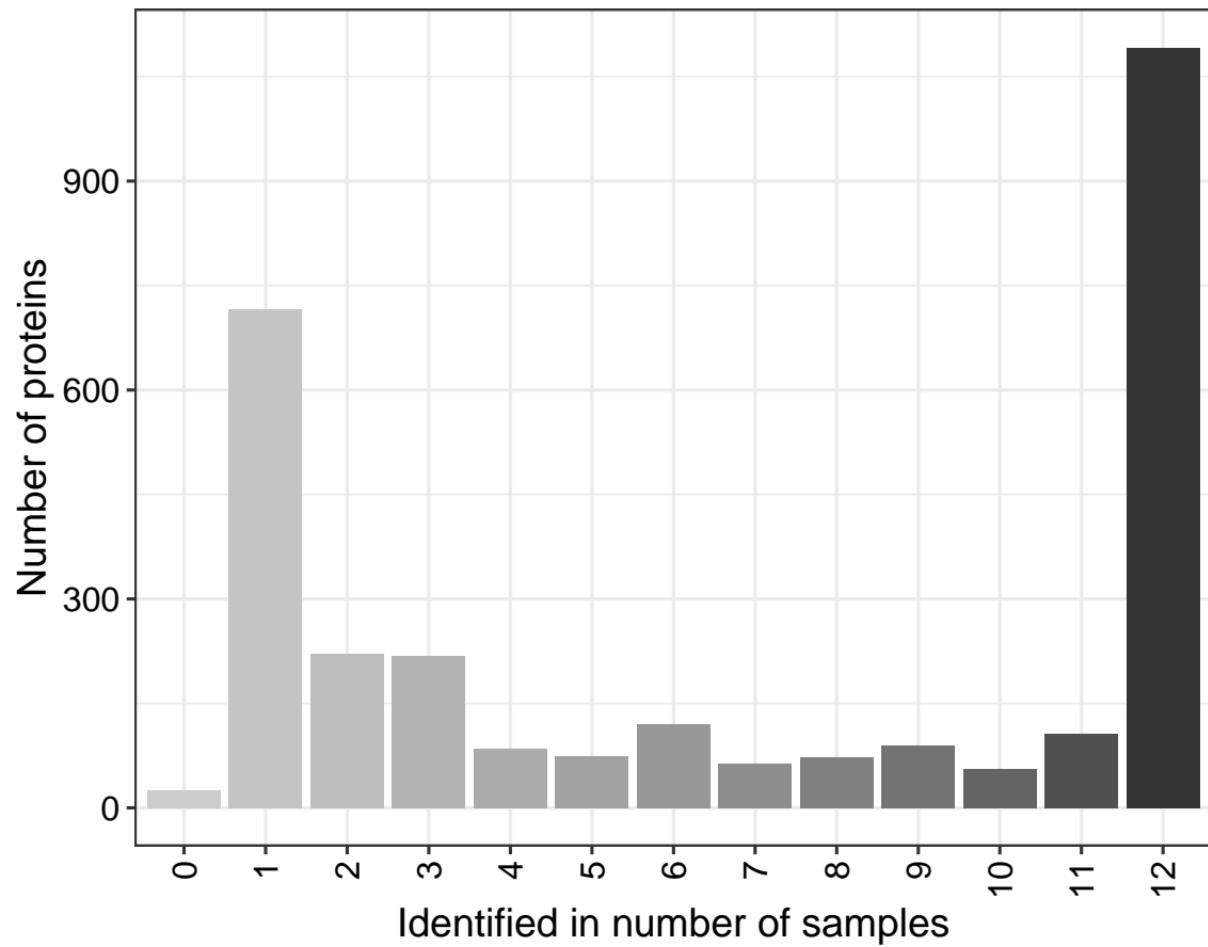
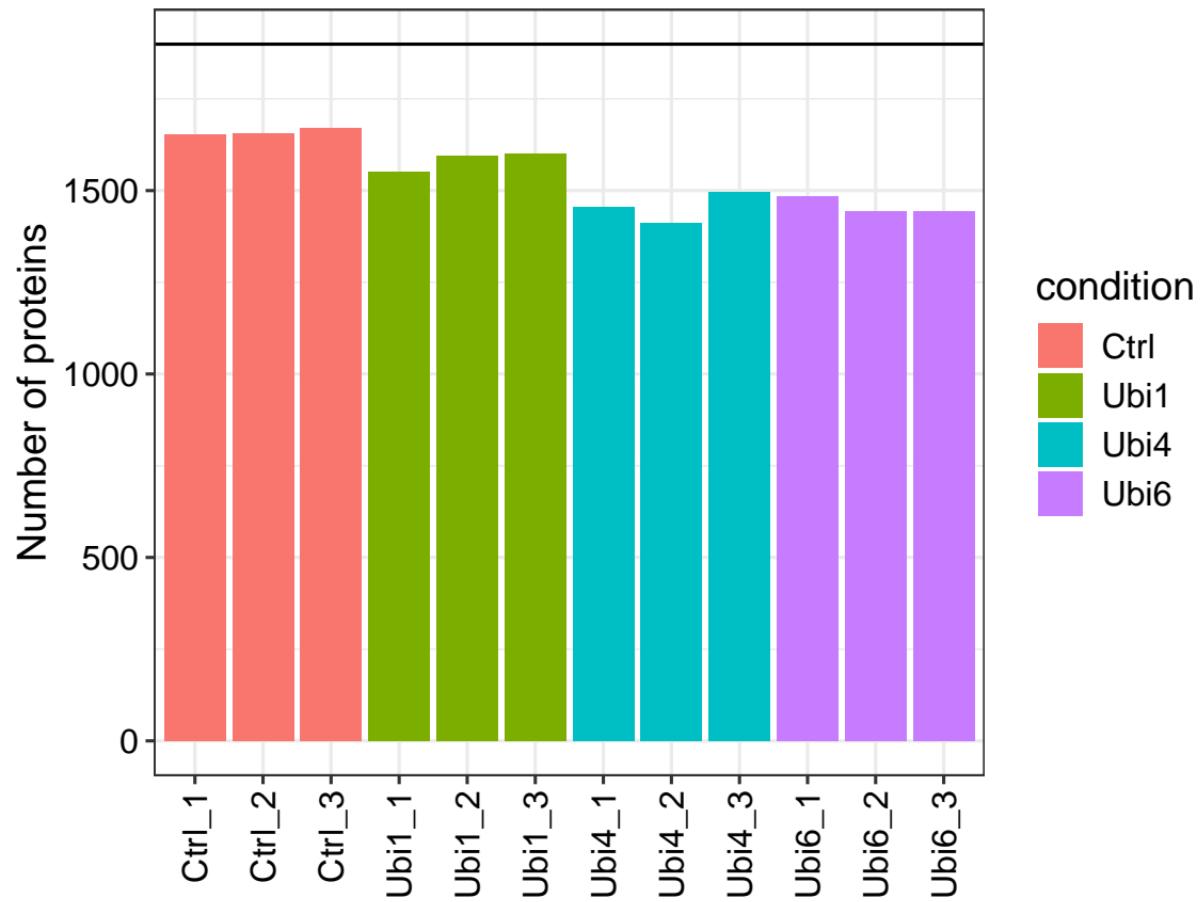


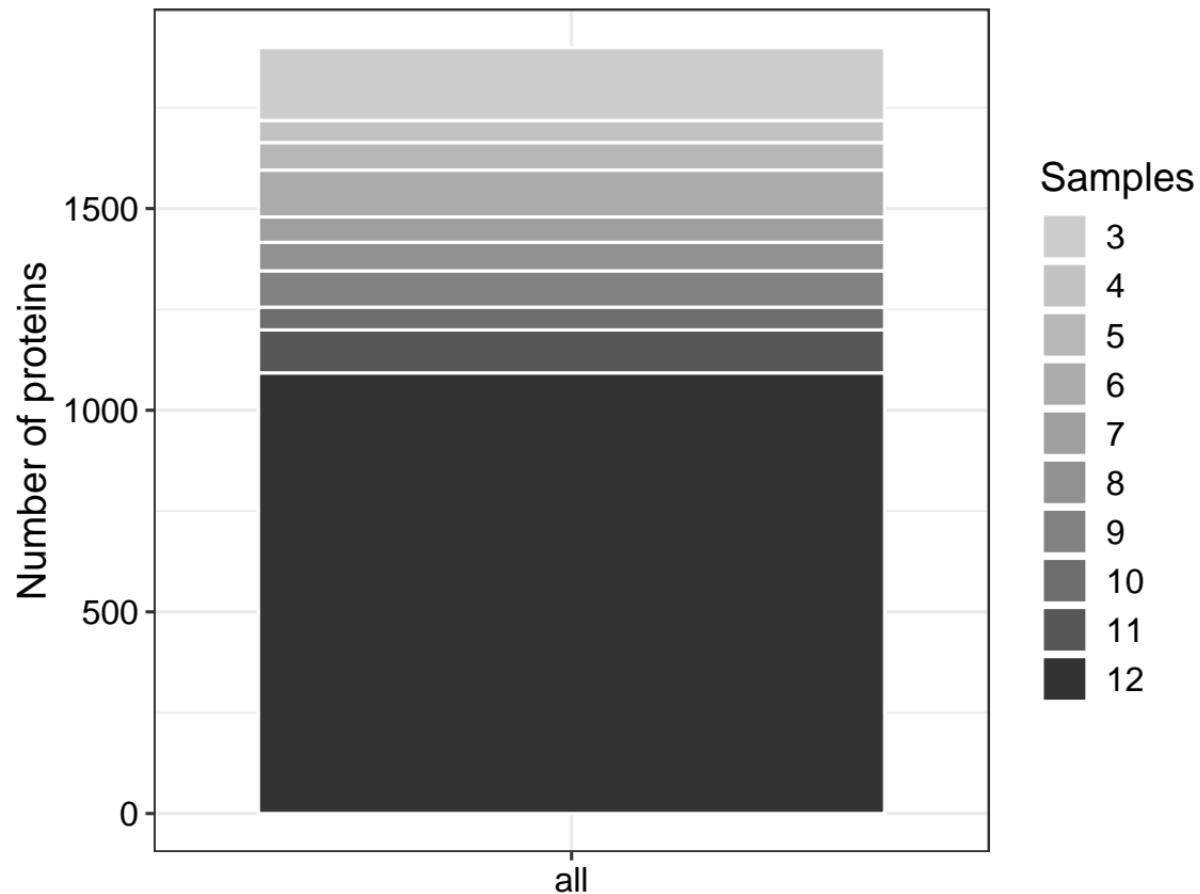
Protein identifications overlap



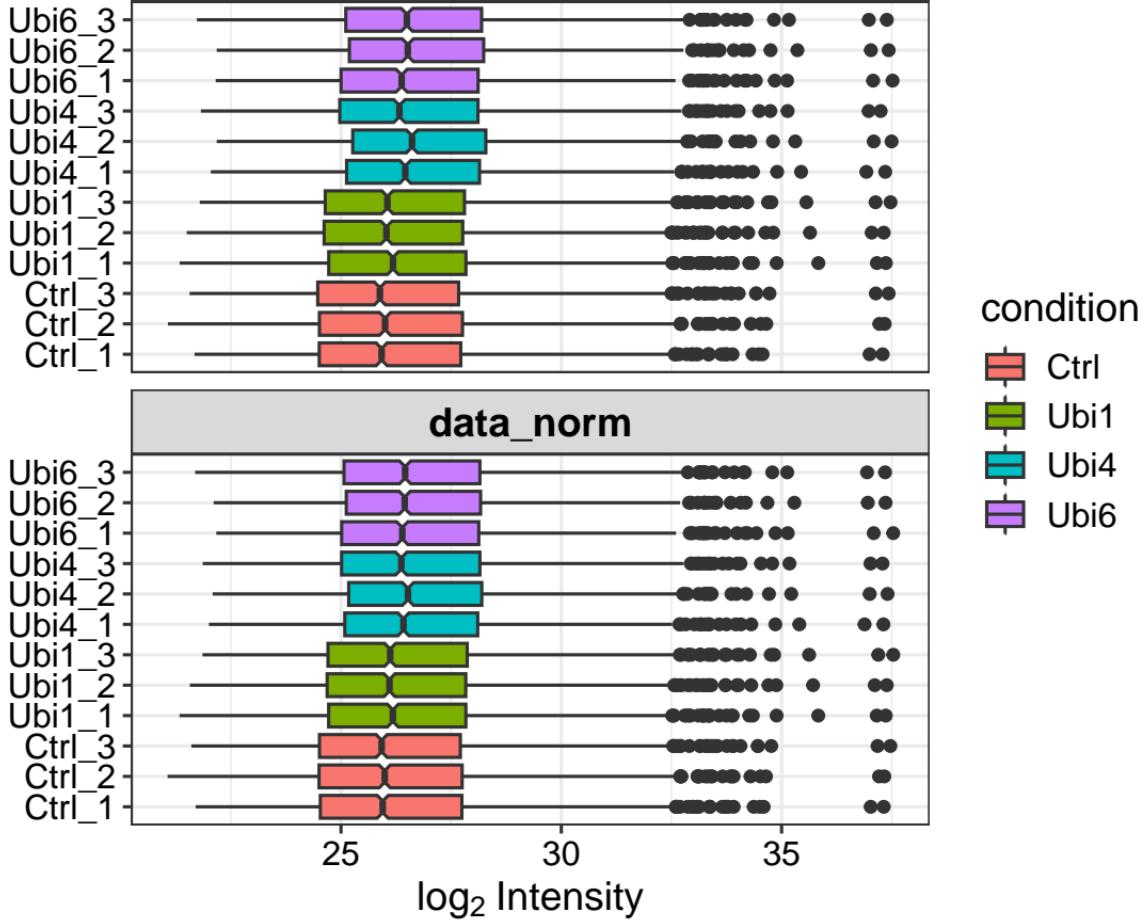
Proteins per sample



Protein coverage



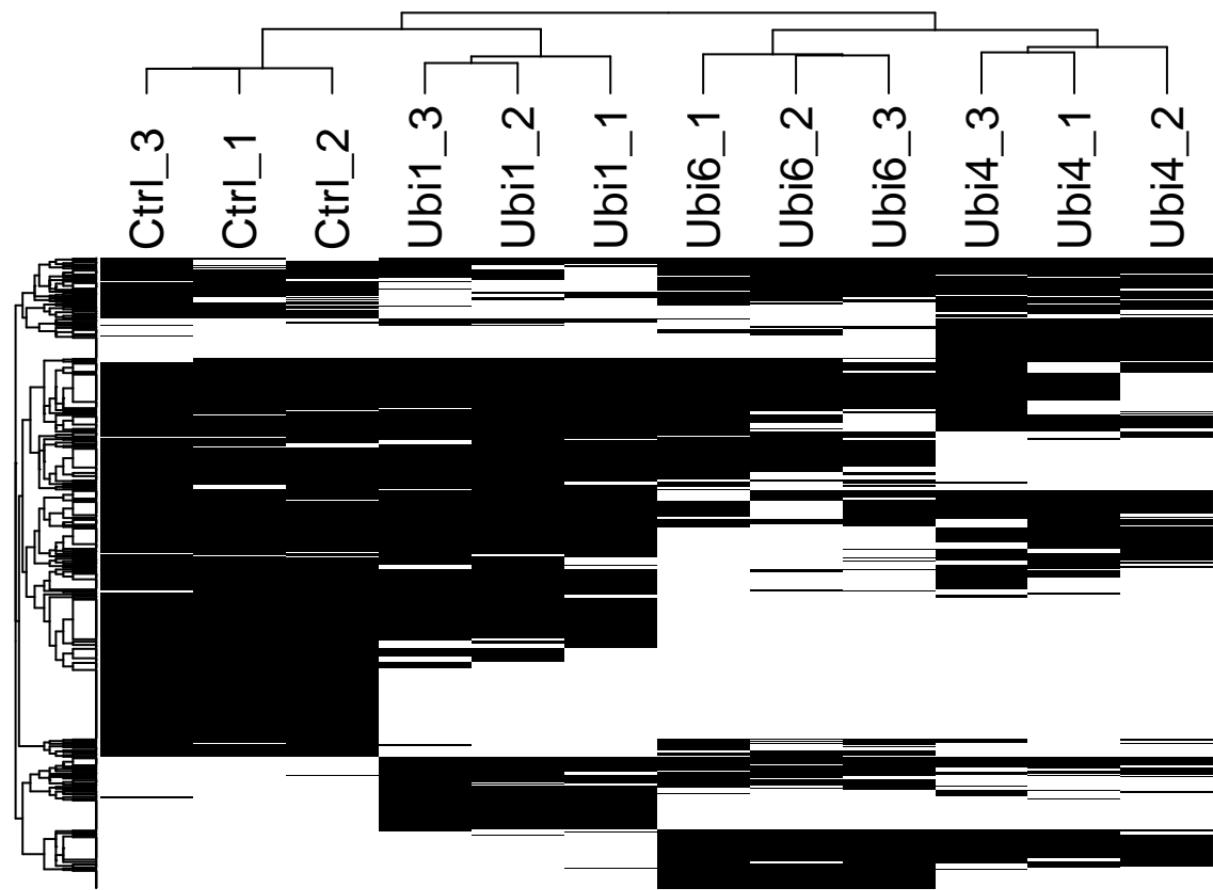
data_filt



Missing values pattern

Missing value

■ Valid value



data_norm

Density

0.20

0.15

0.10

0.05

0.00

data_imp

condition

Ctrl

Ubi1

Ubi4

Ubi6

25

30

35

\log_2 Intensity