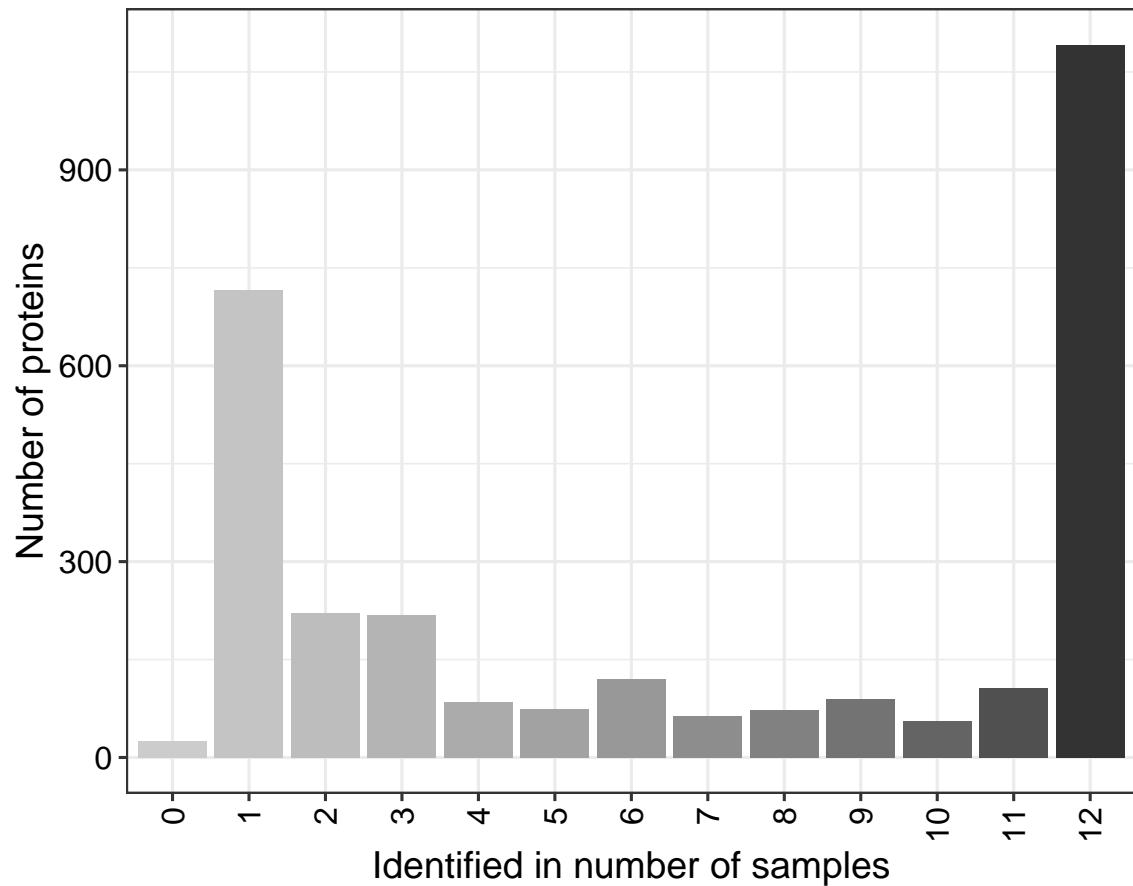
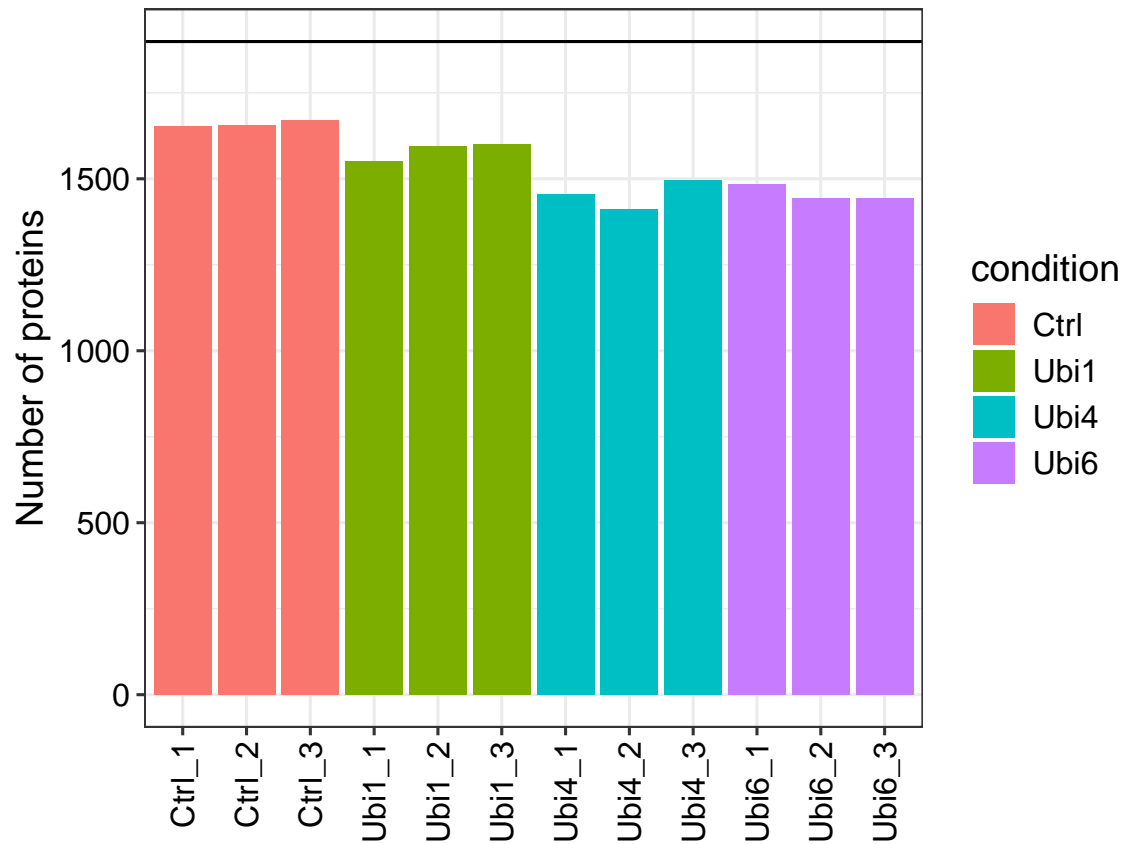


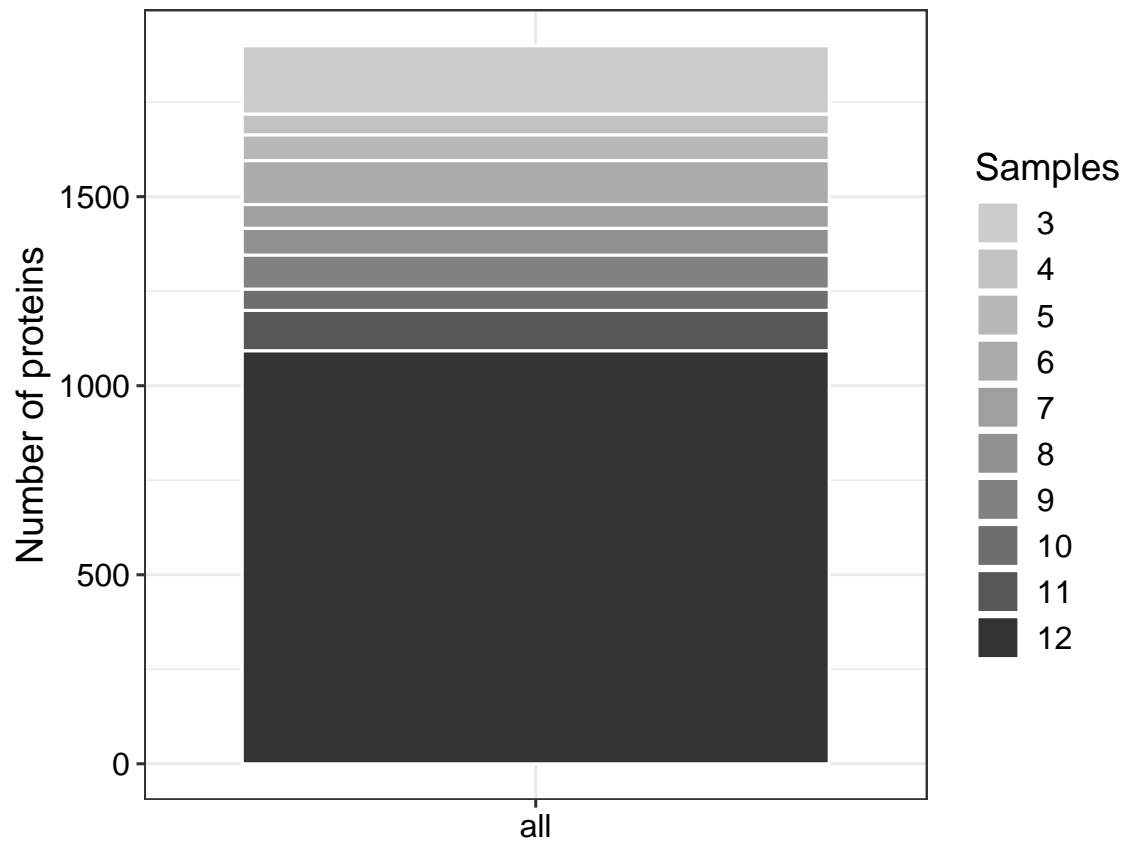
Protein identifications overlap



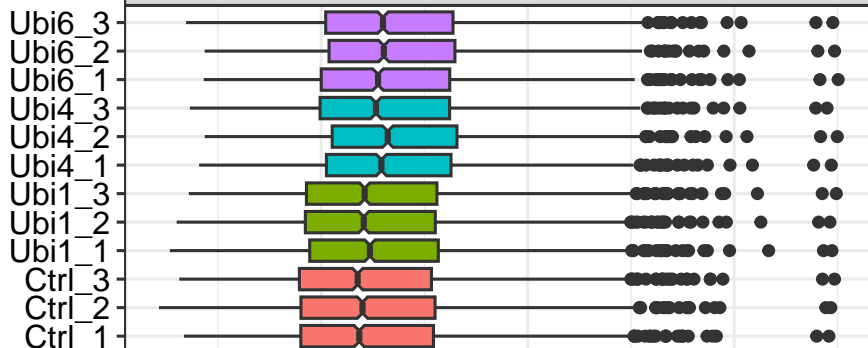
Proteins per sample



Protein coverage



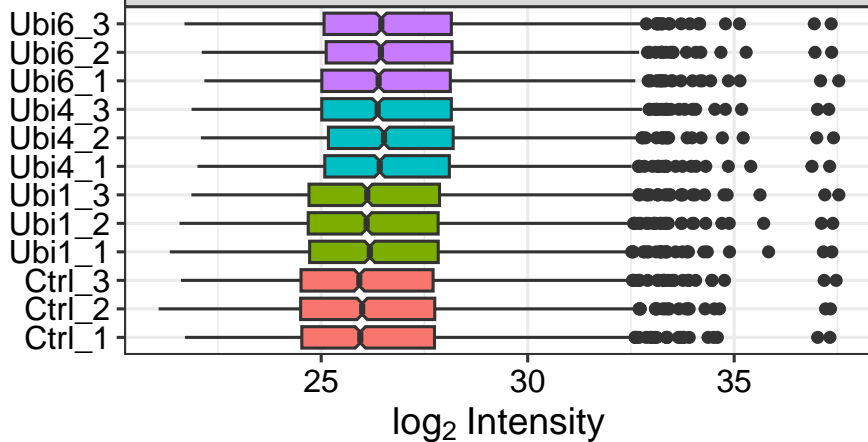
data_filt



condition



data_norm



Missing values pattern

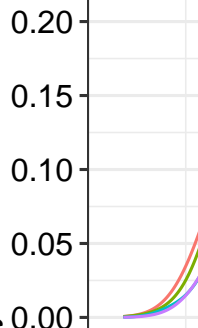
Missing value

Valid value

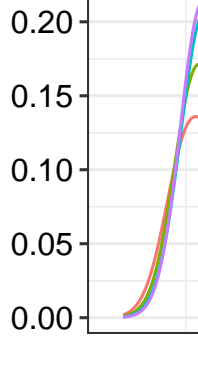
Ctrl_3
Ctrl_1
Ctrl_2
Ubi1_3
Ubi1_2
Ubi1_1
Ubi6_1
Ubi6_2
Ubi6_3
Ubi4_3
Ubi4_1
Ubi4_2



data_norm



data_imp



condition



log₂ Intensity