ID rate Q-value < 0.01 HYE mix A HYE mix B 900 distinct_ions 600 300 0 750 distinct_modified_peptides 500 250 count 0 ion_ID_outlier no 750 distinct_peptides 500 250 75 distinct_proteins 50 25 0 20230314_TIMS1_SM...0ng_R3_S1-C5_1_6356 20230314_TIMS1_SM...0ng_R4_S1-H5_1_6365 20230314_TIMS1_SM...0ng_R2_S1-B5_1_6355 20230314_TIMS1_SM...0ng_R1_S1-A5_1_6354 20230314_TIMS1_SM...0ng_R4_S1-D5_1_6357 20230314_TIMS1_SM...0ng_R1_S1-E5_1_6362 20230314_TIMS1_SM...0ng_R3_S1-G5_1_6364 raw file name