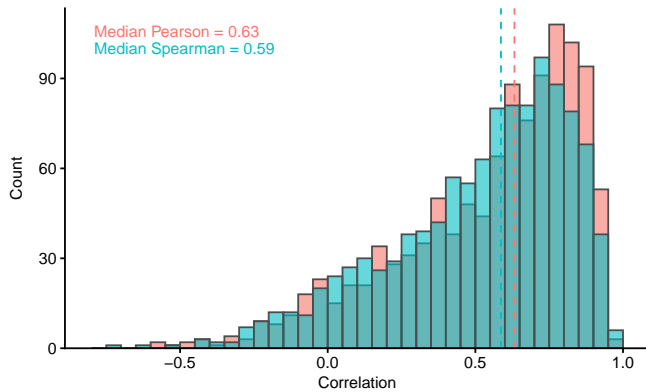
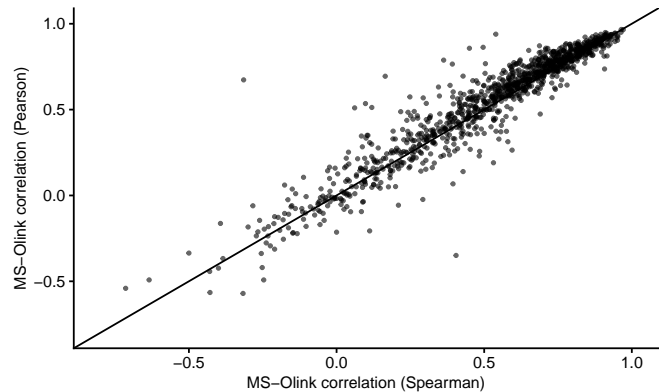


Correlation method Pearson Spearman

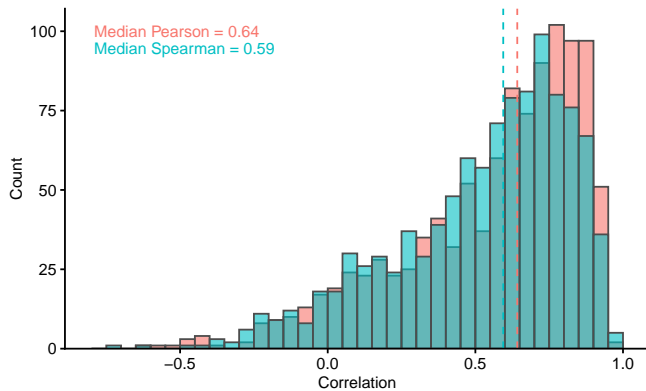
Unprocessed data, all overlapping proteins (N = 1129)



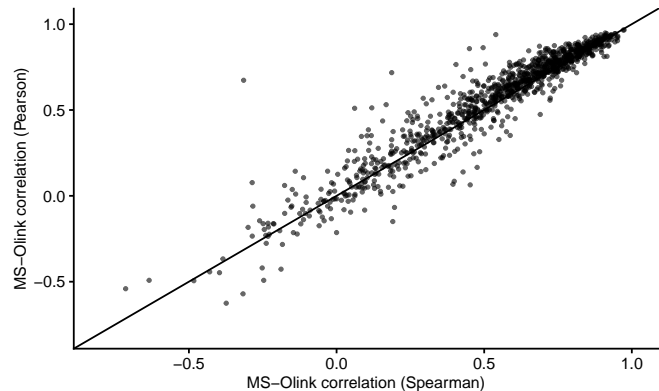
Unprocessed data, all overlapping proteins (N = 1129)



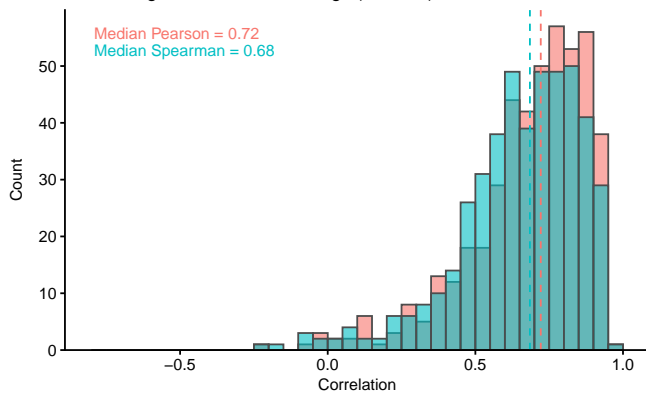
Values <LOD and QC warnings set to missing (N = 1064)



Values <LOD and QC warnings set to missing (N = 1064)



No missing values or QC warnings (N = 463)



No missing values or QC warnings (N = 463)

