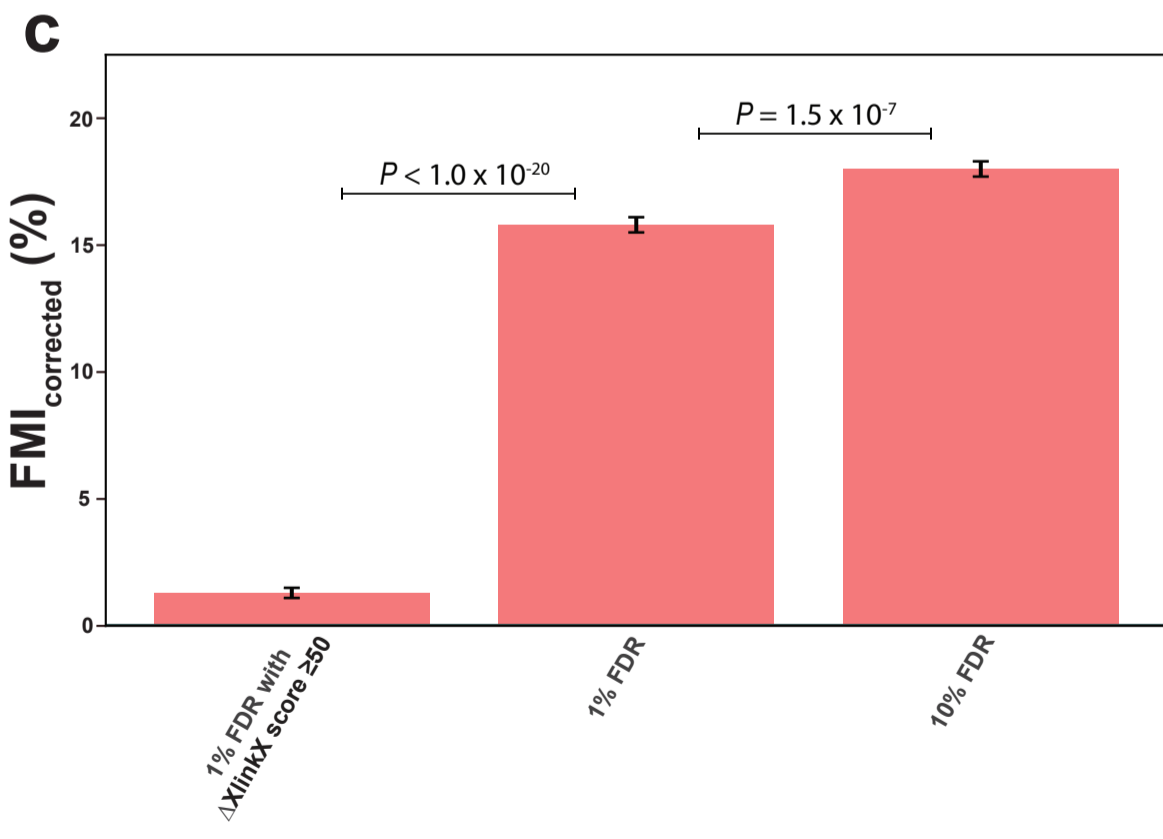
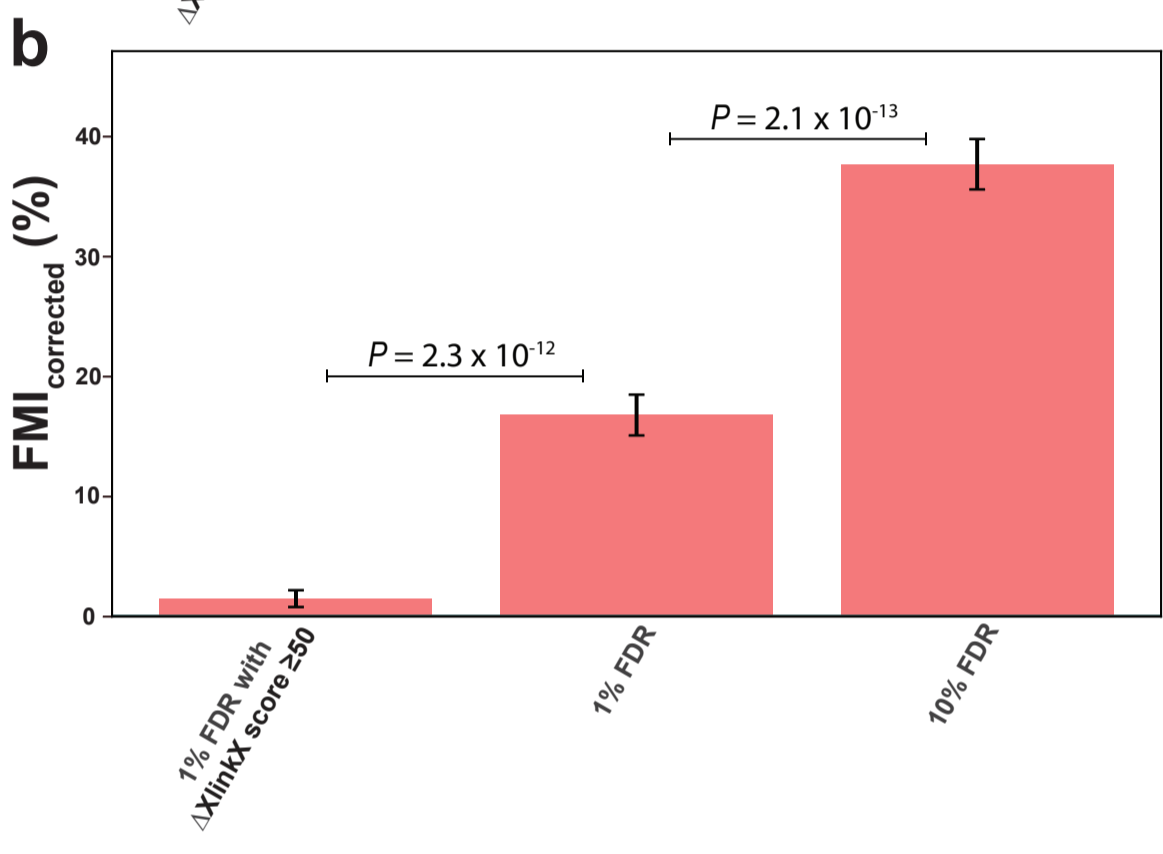
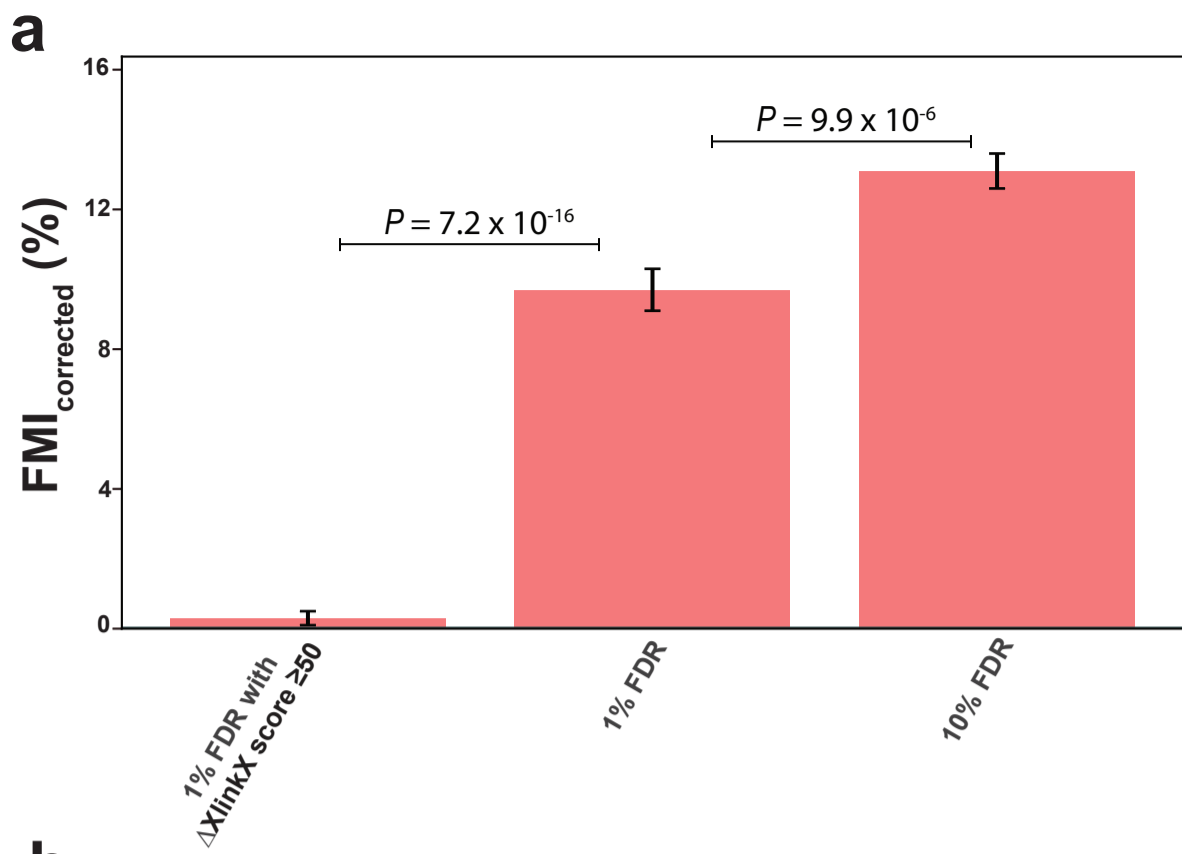


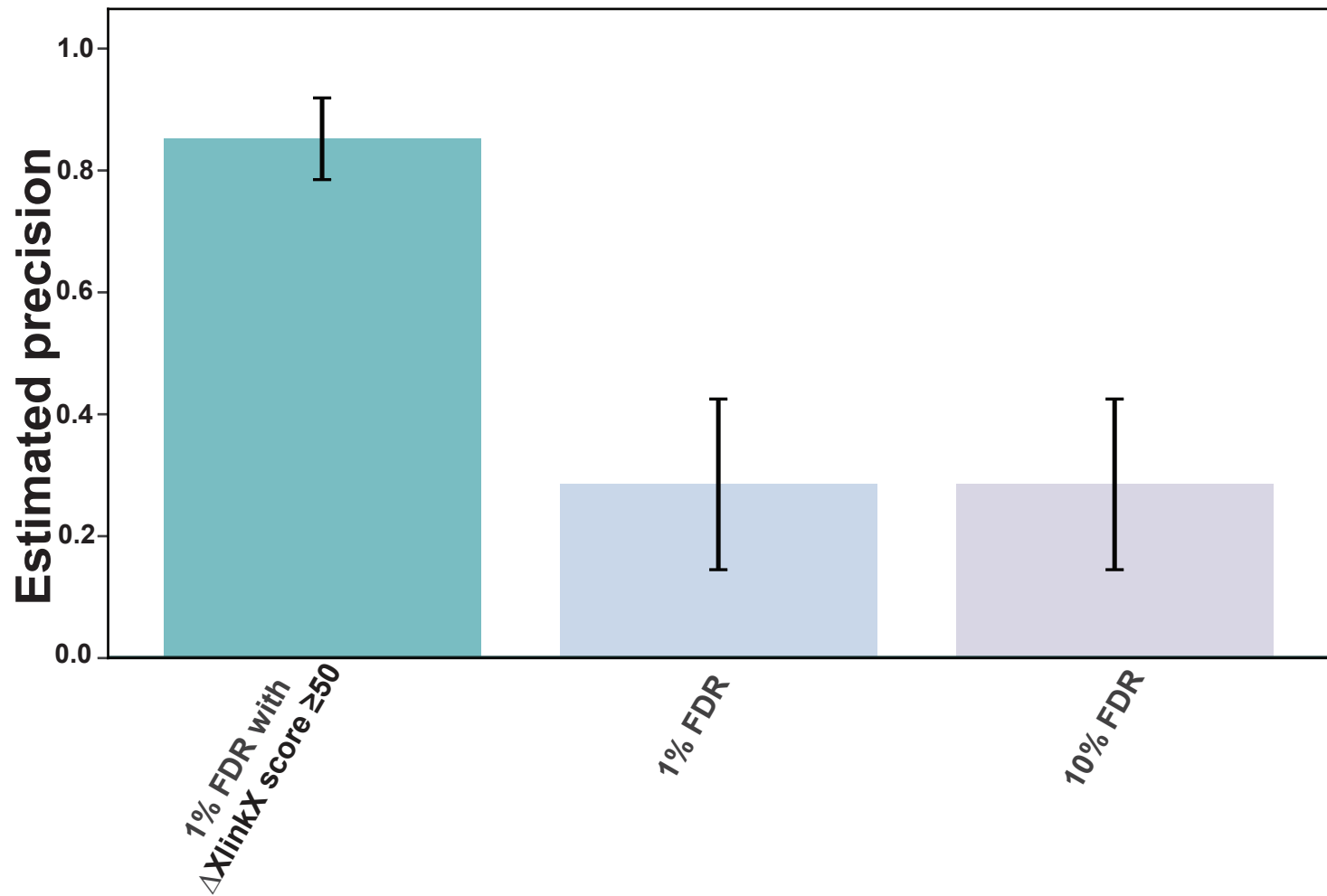
Supplementary Figures

Supplementary Figure 1. Corrected FMI for the three datasets analyzed in the study (Utilizing equation 3 in Methods section). (a) Human proteome XL-MS, (b) *E. coli* proteome XL-MS and (c) Mouse mitochondrial XL-MS.

Supplementary Figure 2. Estimated precision using PCA experiments for the three datasets of different quality from the human K562 proteome-wide XL-MS experiment. See Supplementary Note 1 for a detailed description of the methodology.



Supplementary Figure 1



Supplementary Figure 2