



Supplemental Figure S7 (related to Figure 5): Reproducibility between AP-MS replicates.

(A) Distribution of rank orders of AP-MS replicate Pearson correlation compared to all other replicates based on the raw spectral counts from AP-MS results, using a matrix of 236 baits (each of the 118 in duplicate) x 344 prey proteins (the same set described in the text for PANTHER analysis). For example, a rank order of 1 for EGR2 indicates that EGR2 replicate 1 was more similar to EGR2 replicate 2 than all other 235 individual AP-MS experiments. (B) Distribution of Pearson correlations for each pair of AP-MS replicates for the same C2H2-ZFP, as in A. The correlation between replicates is above 90% for 93/118 C2H2-ZFs.