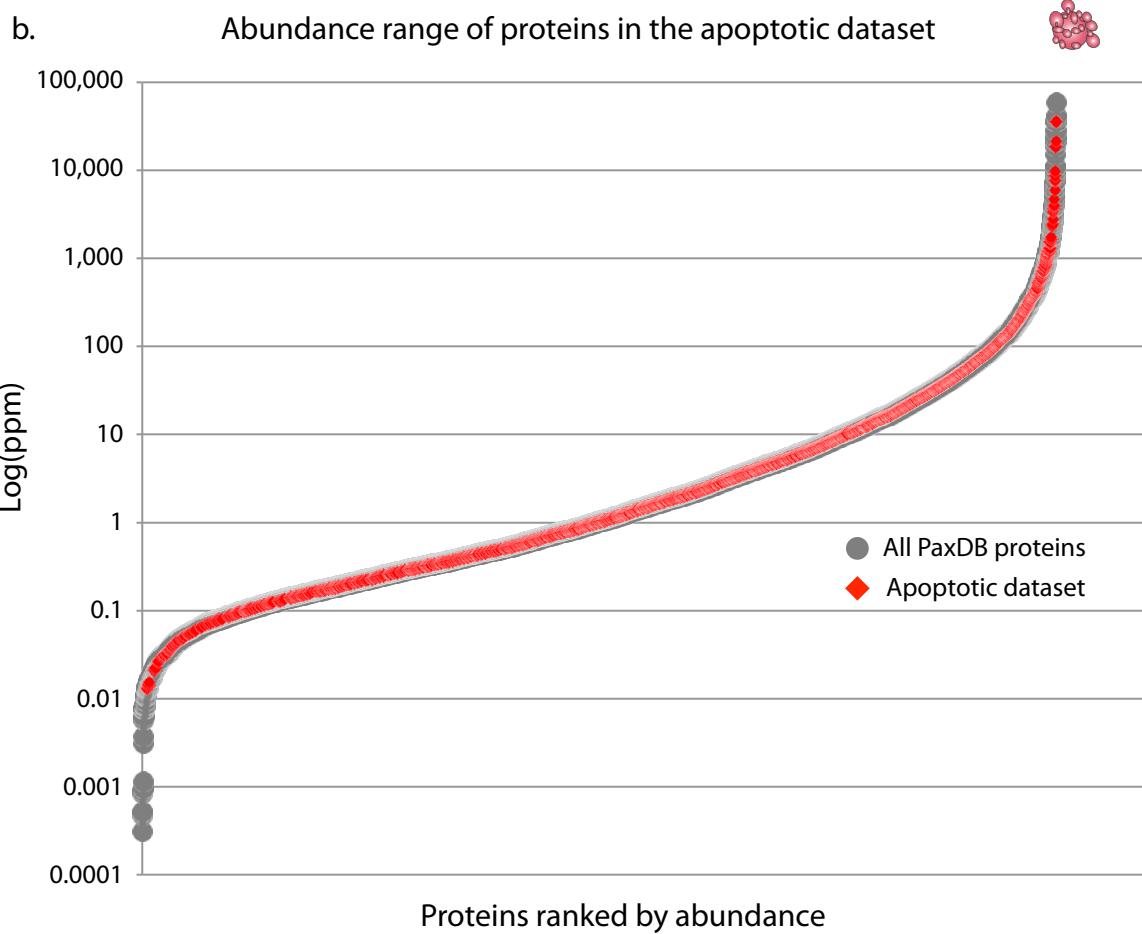
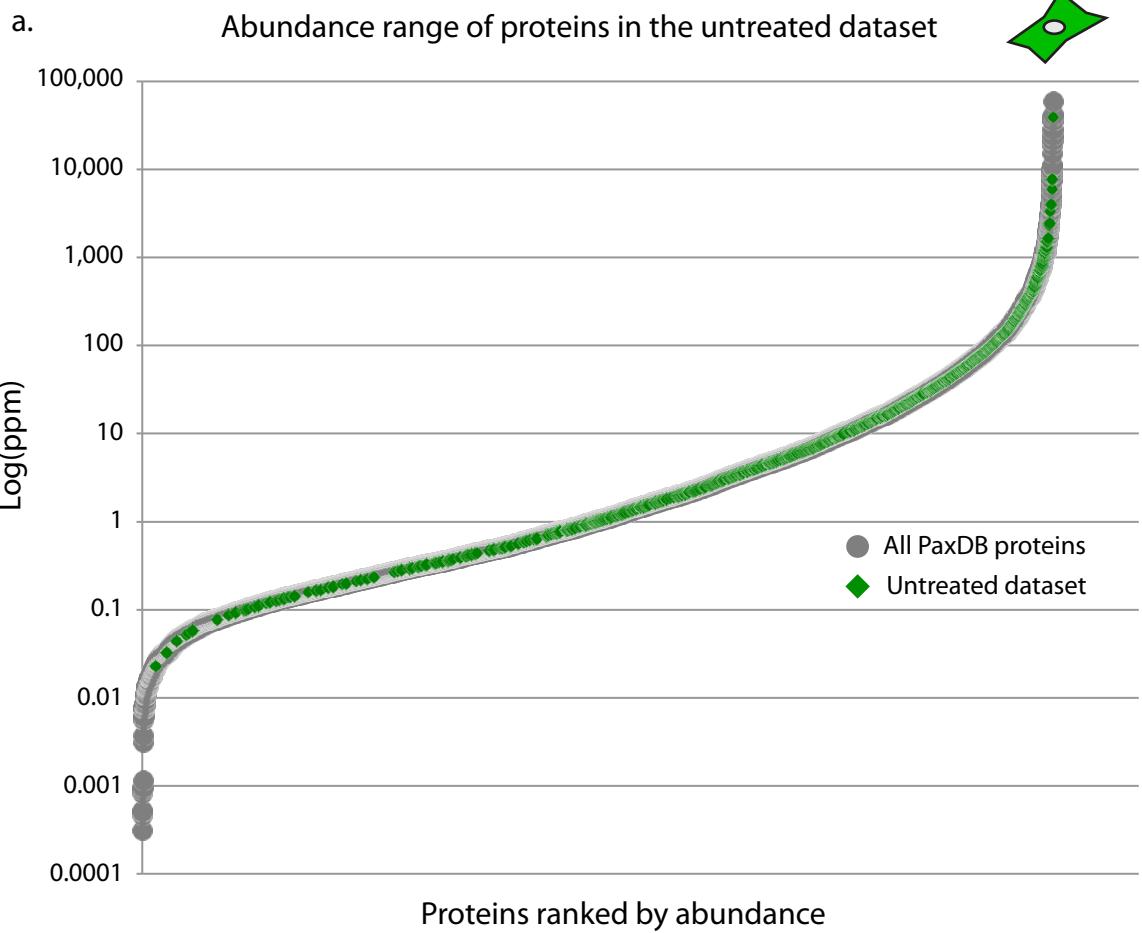
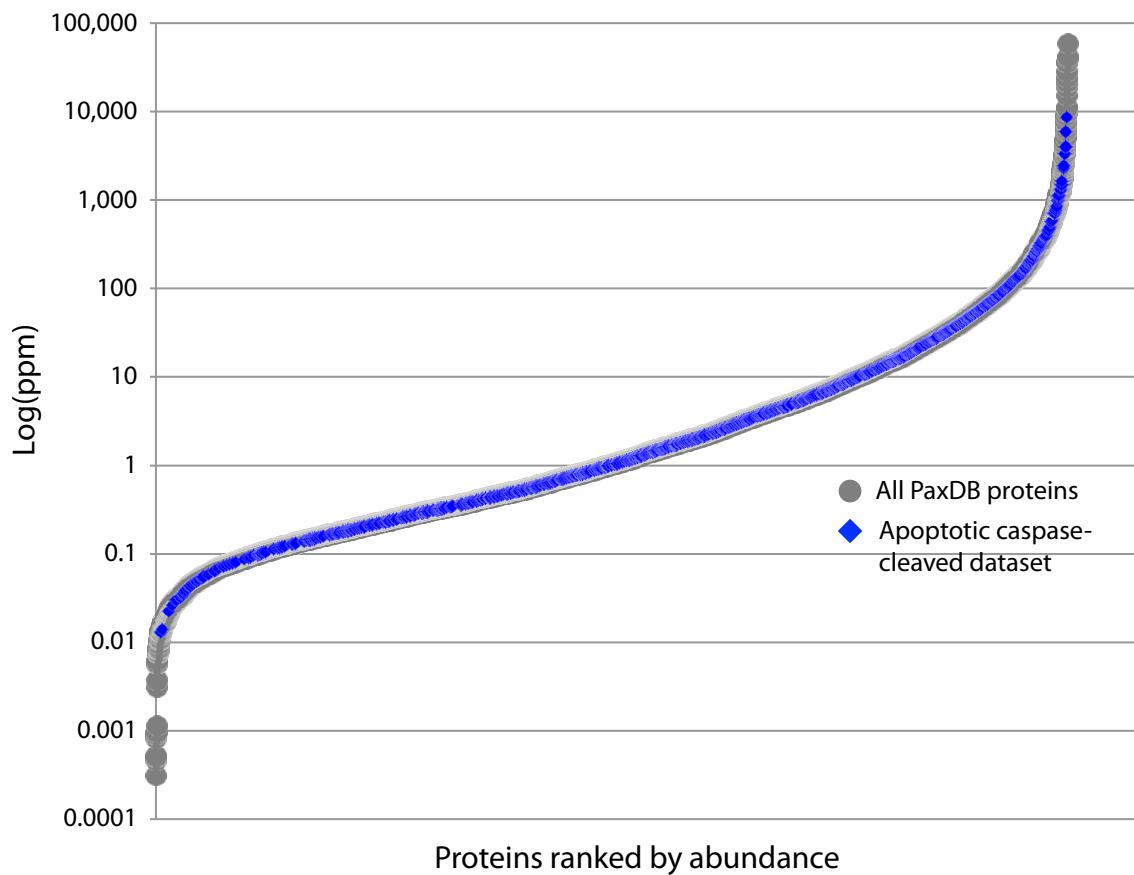


Supplemental Figure 1: Protein abundance

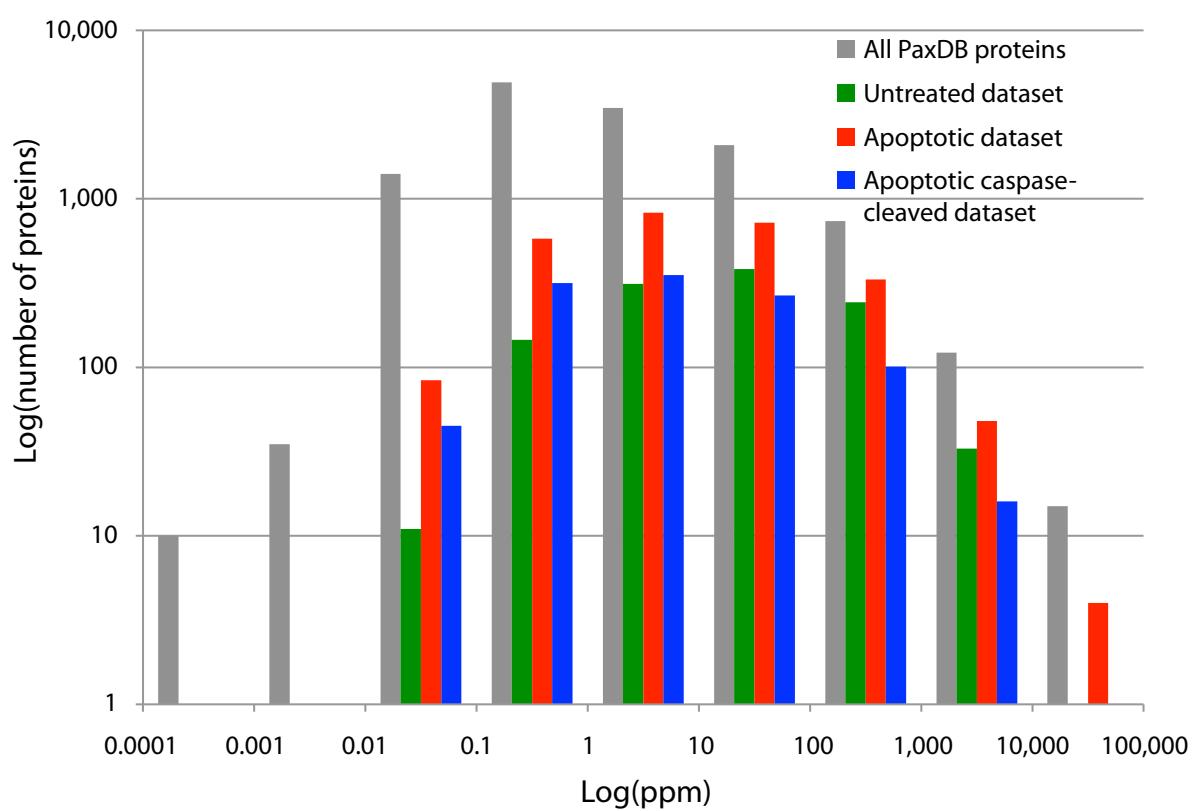


c. Abundance range of proteins in the apoptotic caspase-cleaved dataset

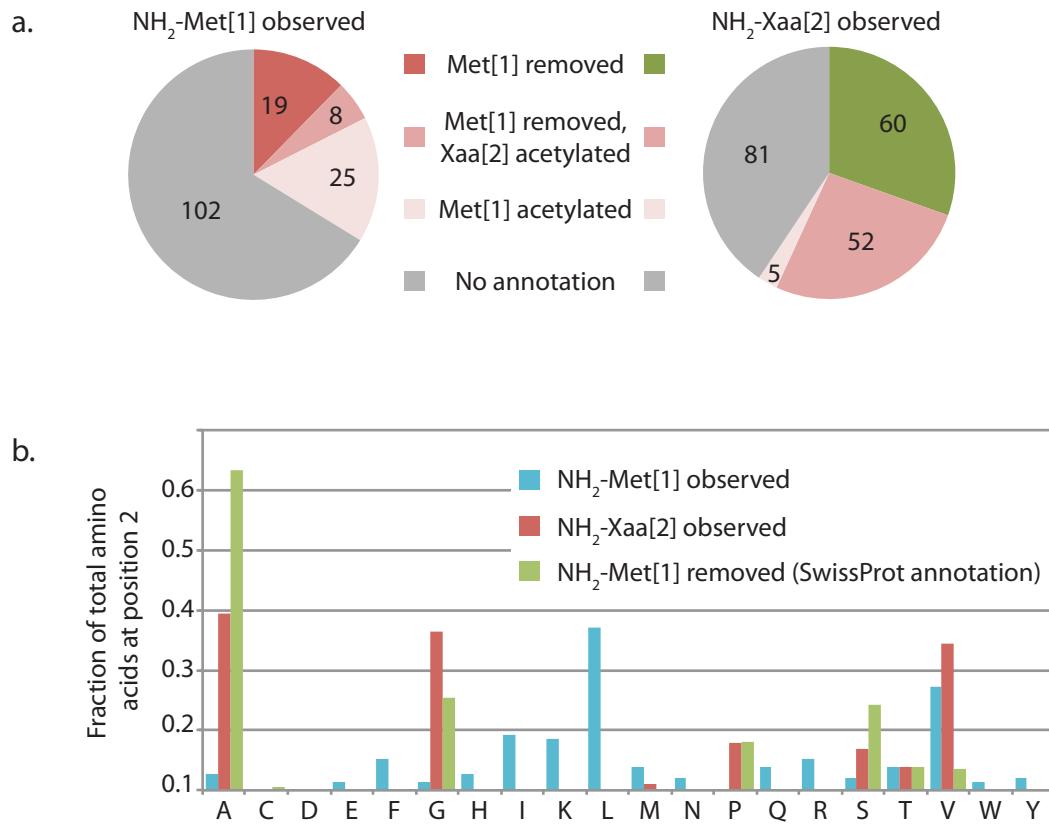


d.

Abundance histogram for all datasets



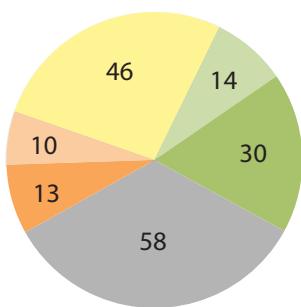
Supplemental Figure 2



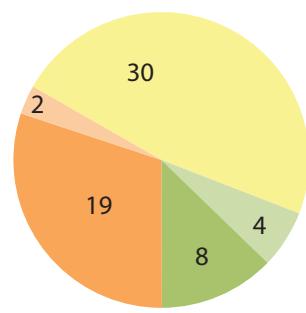
### Supplemental Figure 3

- [Yellow square] Observed same as SwissProt
- [Orange square] Observed C-term of SwissProt by 1 residue
- [Dark Orange square] Observed C-term of SwissProt by >1 residue
- [Light Green square] Observed N-term of SwissProt by 1 residue
- [Dark Green square] Observed N-term of SwissProt by >1 residue
- [Grey square] No SwissProt annotation

a. Transit peptide removal sites  
(all proteins in MitoCarta):

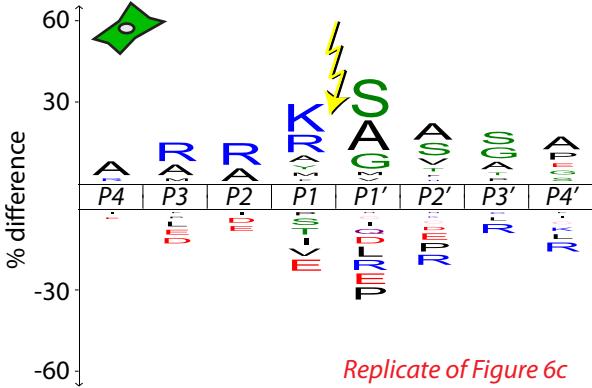


b. Signal peptide removal sites:

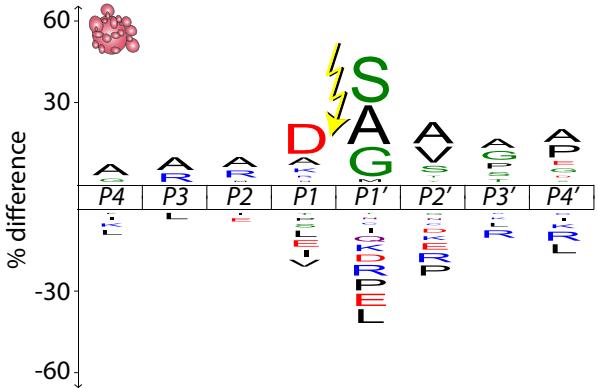


Supplemental Figure 4: IceLogos for untreated and apoptotic datasets, P1' site 3-65 and 66+

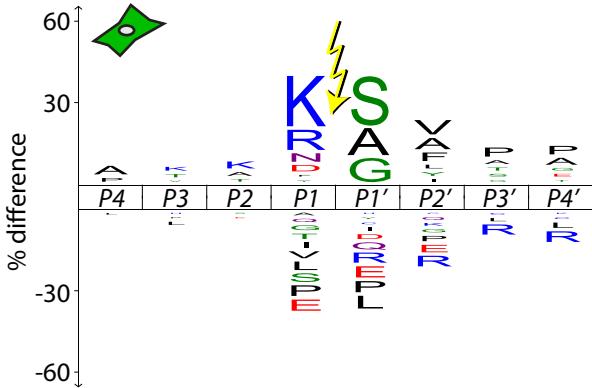
a. Untreated dataset, cleavage site P1' is 3-65



b. Apoptotic dataset, cleavage site P1' is 3-65



c. Untreated dataset, cleavage site P1' is 66+



d. Apoptotic dataset, cleavage site P1' is 66+

