

Targeting Hodgkin and Reed-Sternberg cells with an inhibitor of heat-shock protein 90: Molecular pathways of response and potential mechanisms of resistance

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Supplementary Figure S1

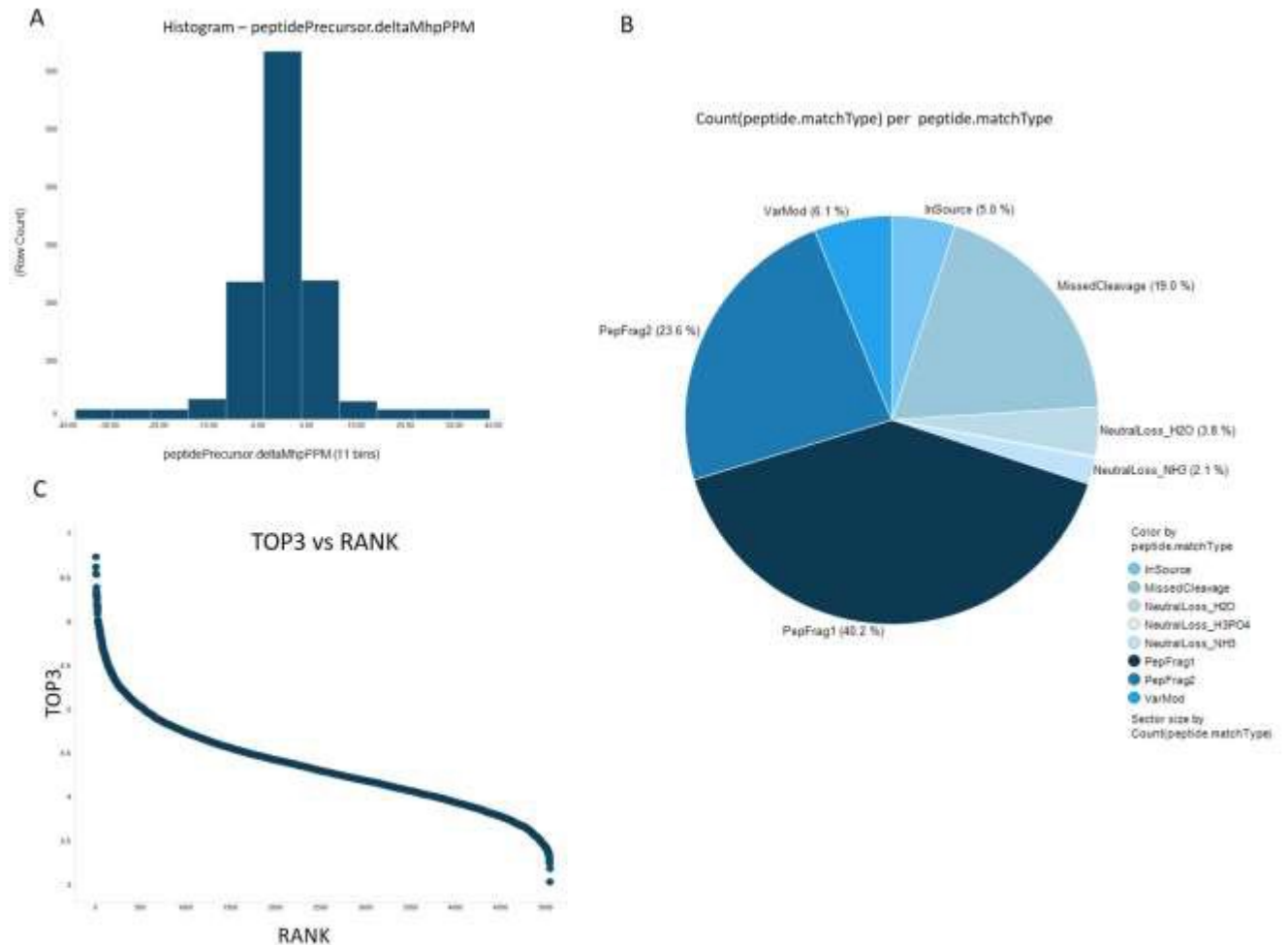


Figure S1: Label-free proteomic analysis. (A) Histogram of peptides error distribution. (B) Peptide match distribution of the identified peptides. PepFrag1 and PepFrag2: peptide matches when compared with database by PLGS; VarMod: variable modifications; InSource: fragmentation that occurred on ionization source; neutral loss H₂O: water precursor loss. (C) Dynamic range of plasma proteomics from the data obtained.