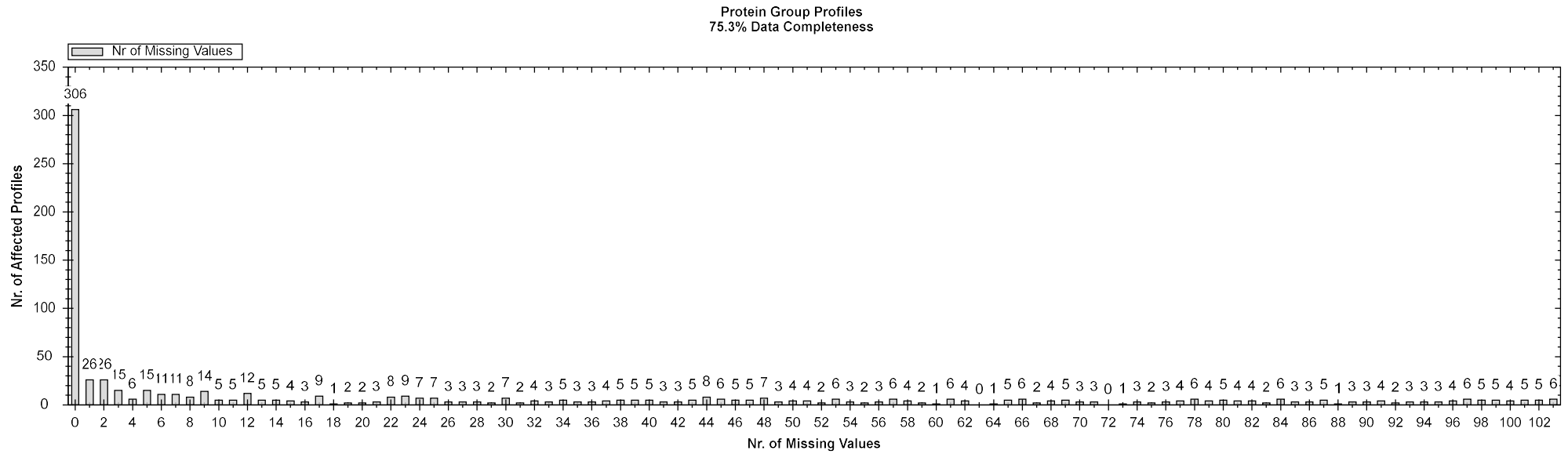
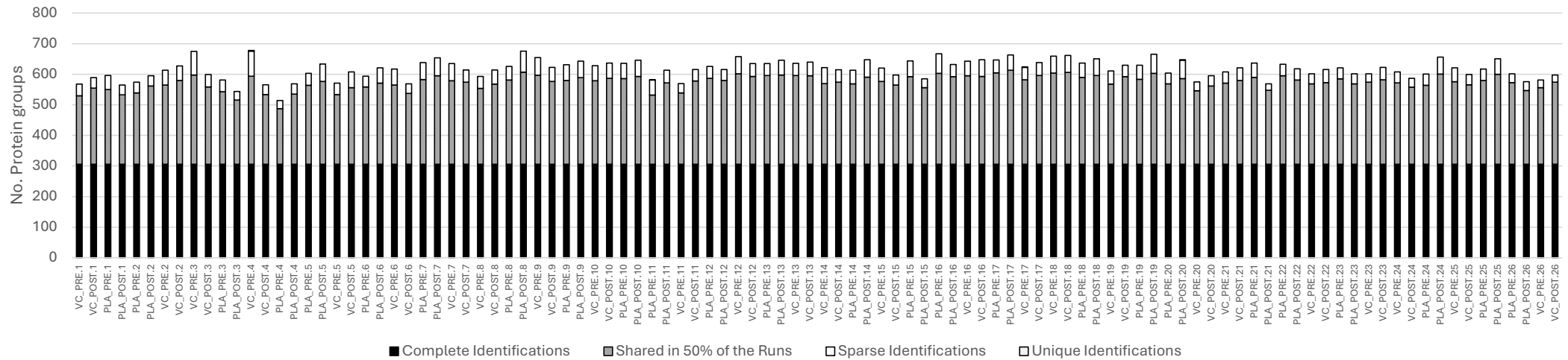


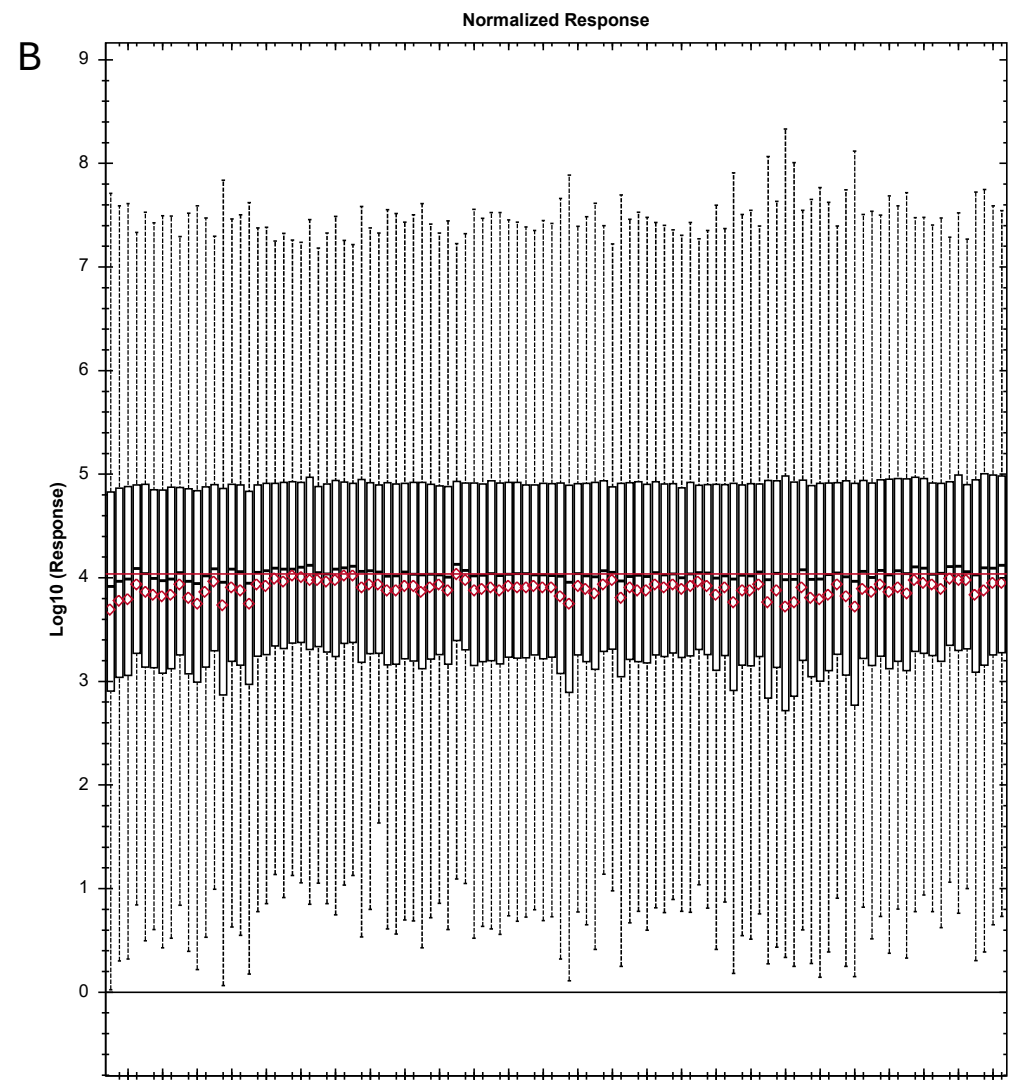
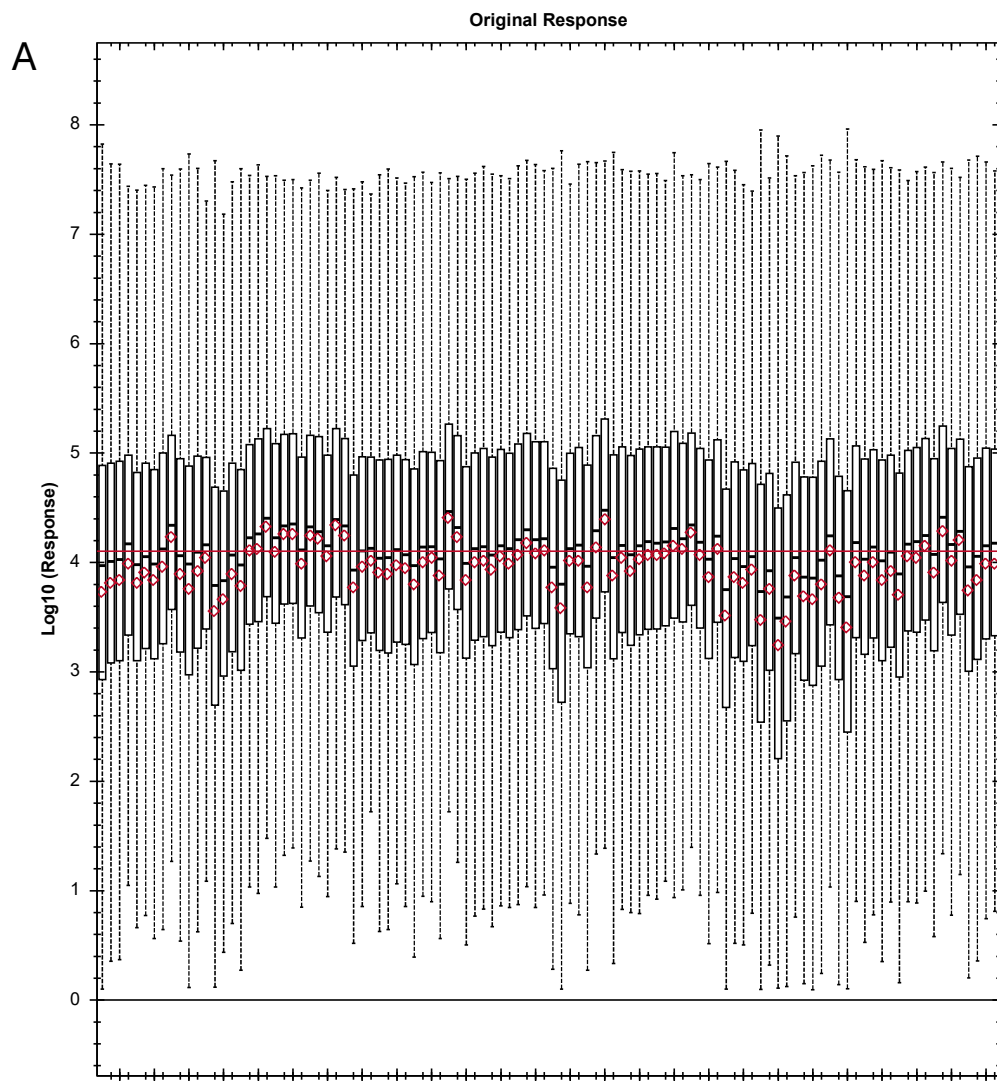
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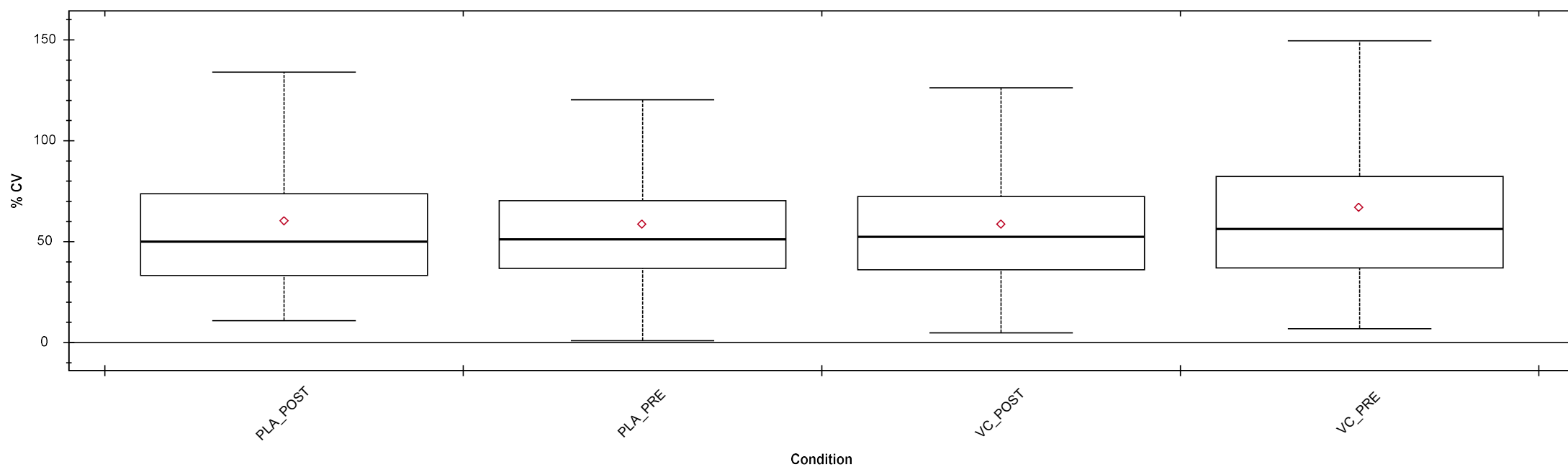
B



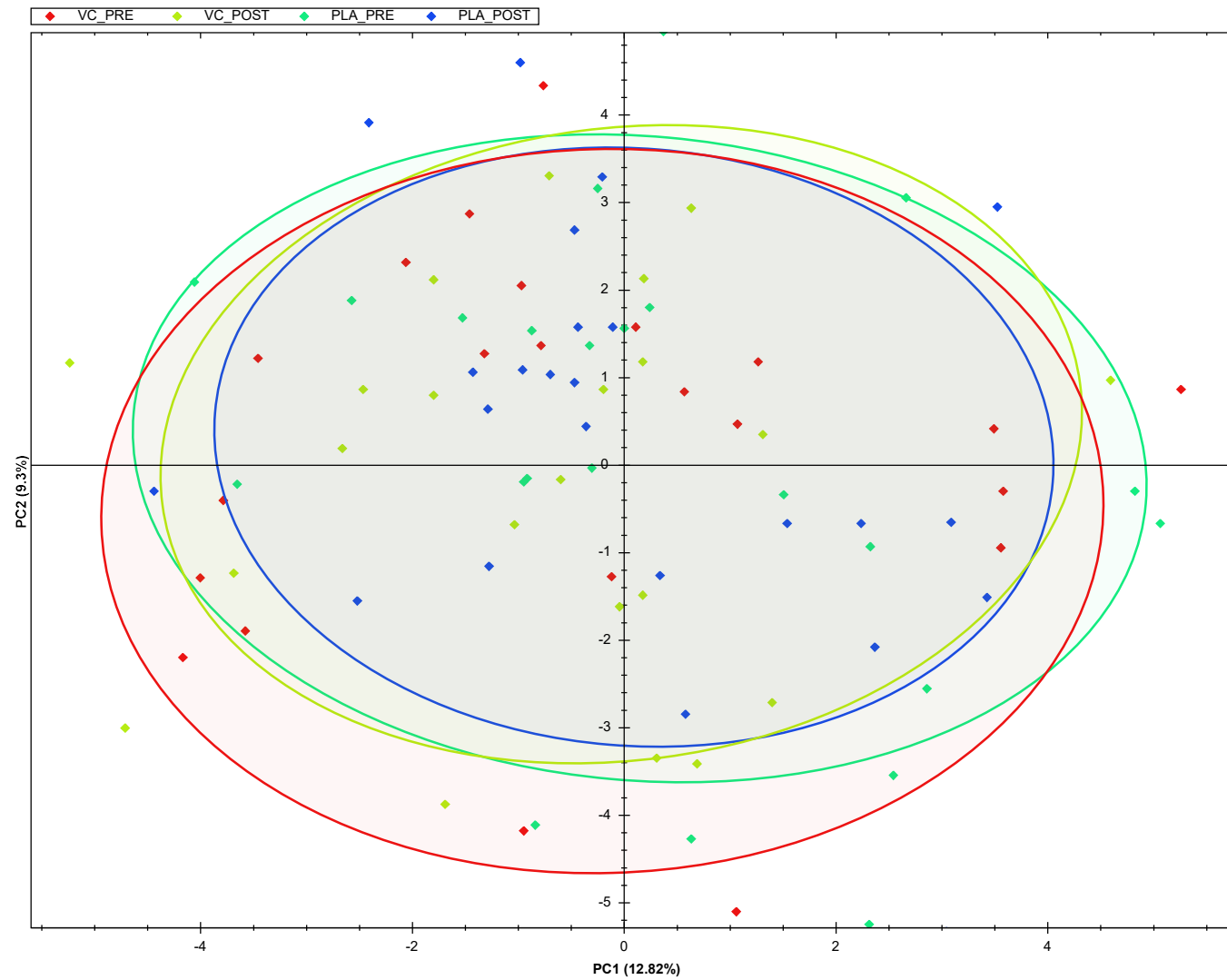
Supplementary Figure S1. Protein group data completeness across samples showing missing data values (A) and number of protein groups found across all samples noting complete, 50% shared, sparse, and unique identifications (B) (Figure generated using Spectronaut v19.0).



Supplementary Figure S2. Original (A) and normalised (B) peptide signal responses across n=104 samples (Figures generated using Spectronaut v19.0 and adapted).



Supplementary Figure S3. % coefficient of variation for each condition. Data shows median, mean, interquartile range, and range of data within each condition (Figure generated using Spectronaut v19.0).



Supplementary Figure S4. Principal component analysis including comparison of all conditions (Figure generated using Spectronaut v19.0).