Figure S4
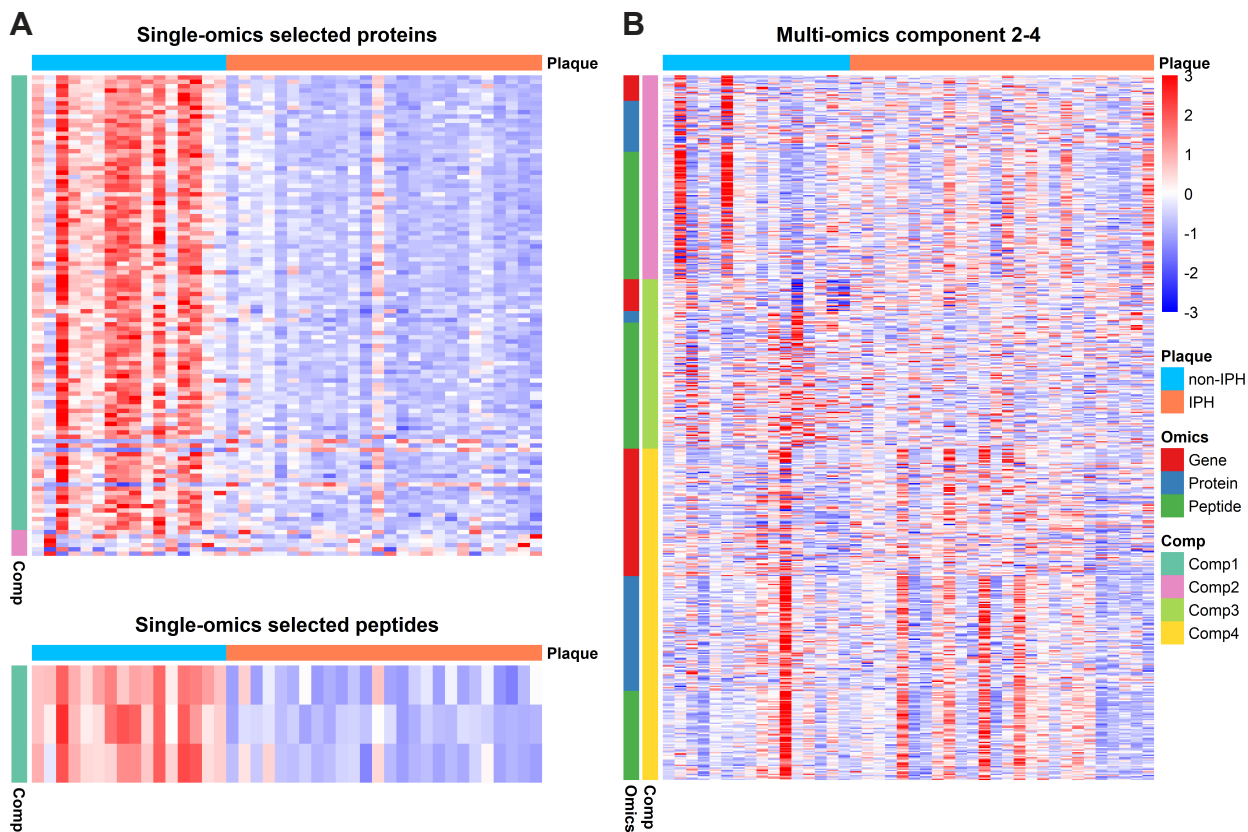

Figure S4 Heatmap of the selected features

(A) Heatmaps showing the expression level of the selected proteins from proteomics and peptides from peptidomics by single-omics analysis. (B) Heatmap showing the expression level of the selected features for component 2–4 of multi-omics analysis. Expression values were z-normalized per row.