**Additional Figures**

**Potential early clinical stage colorectal cancer diagnosis using a proteomics blood test**

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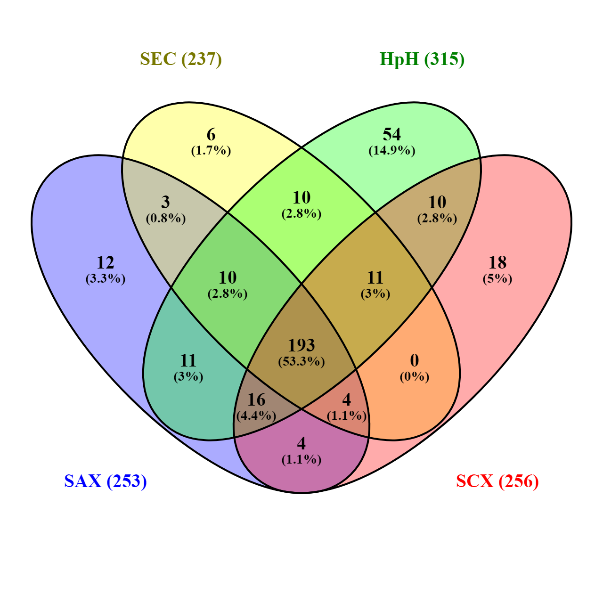
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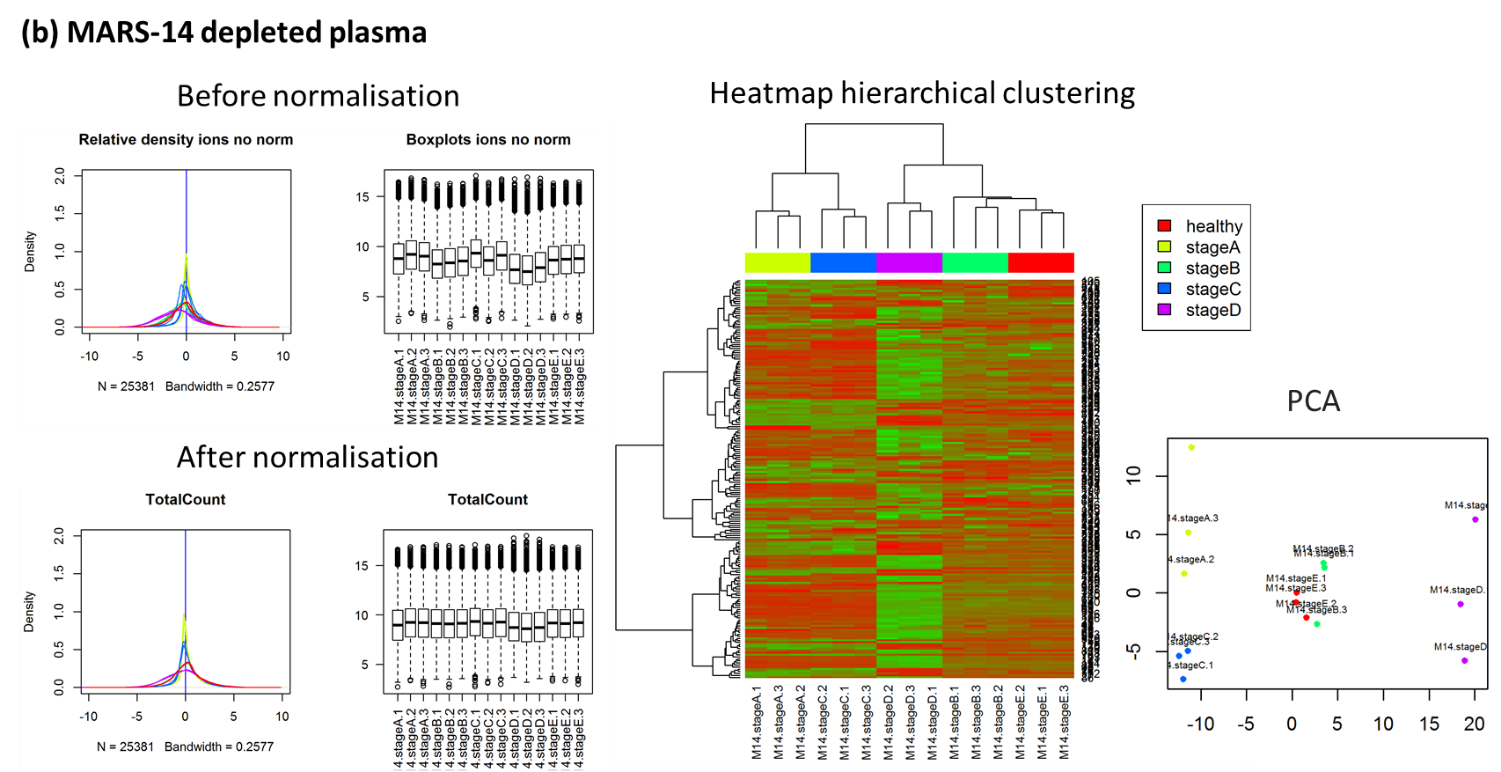
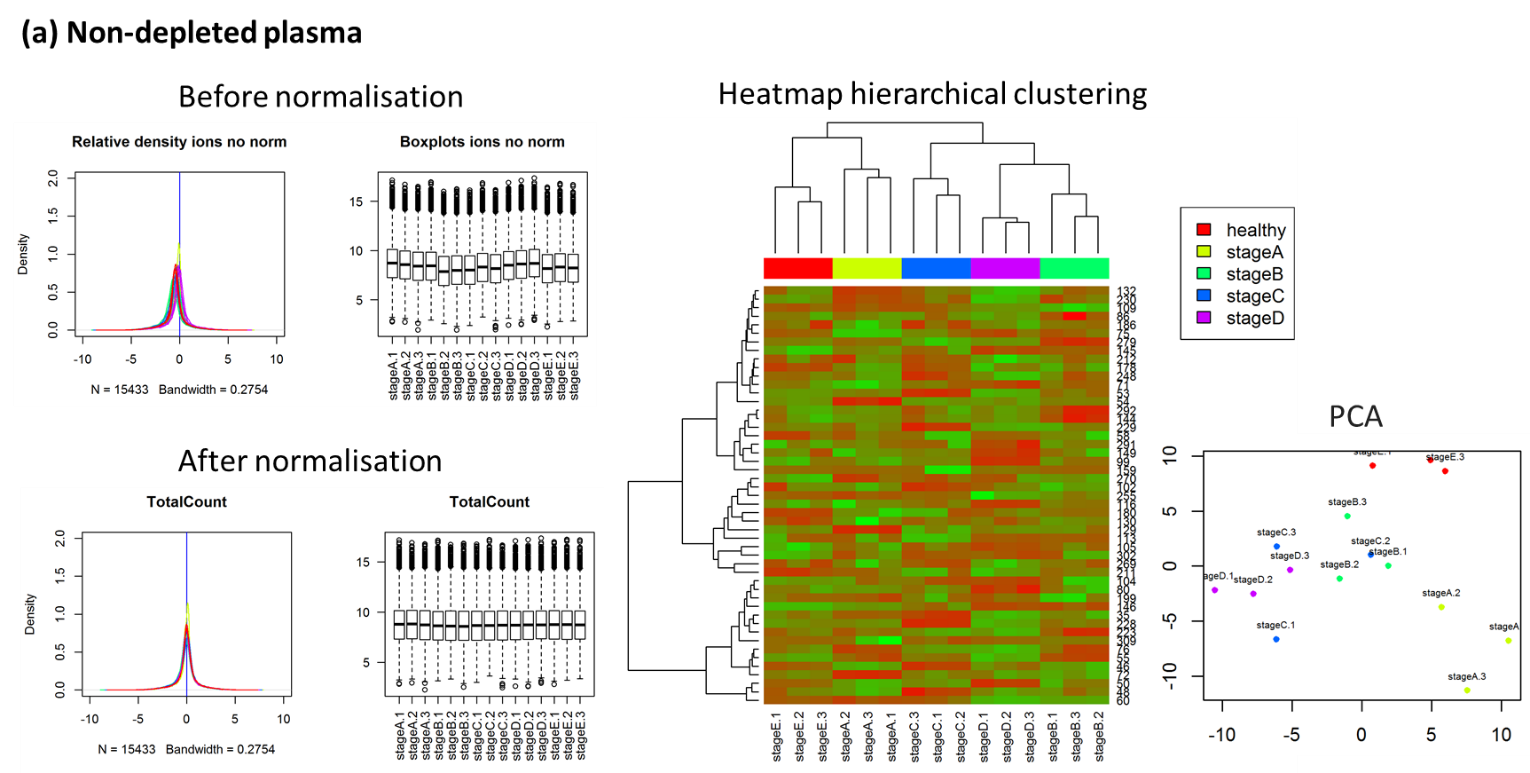
+ These authors contributed equally

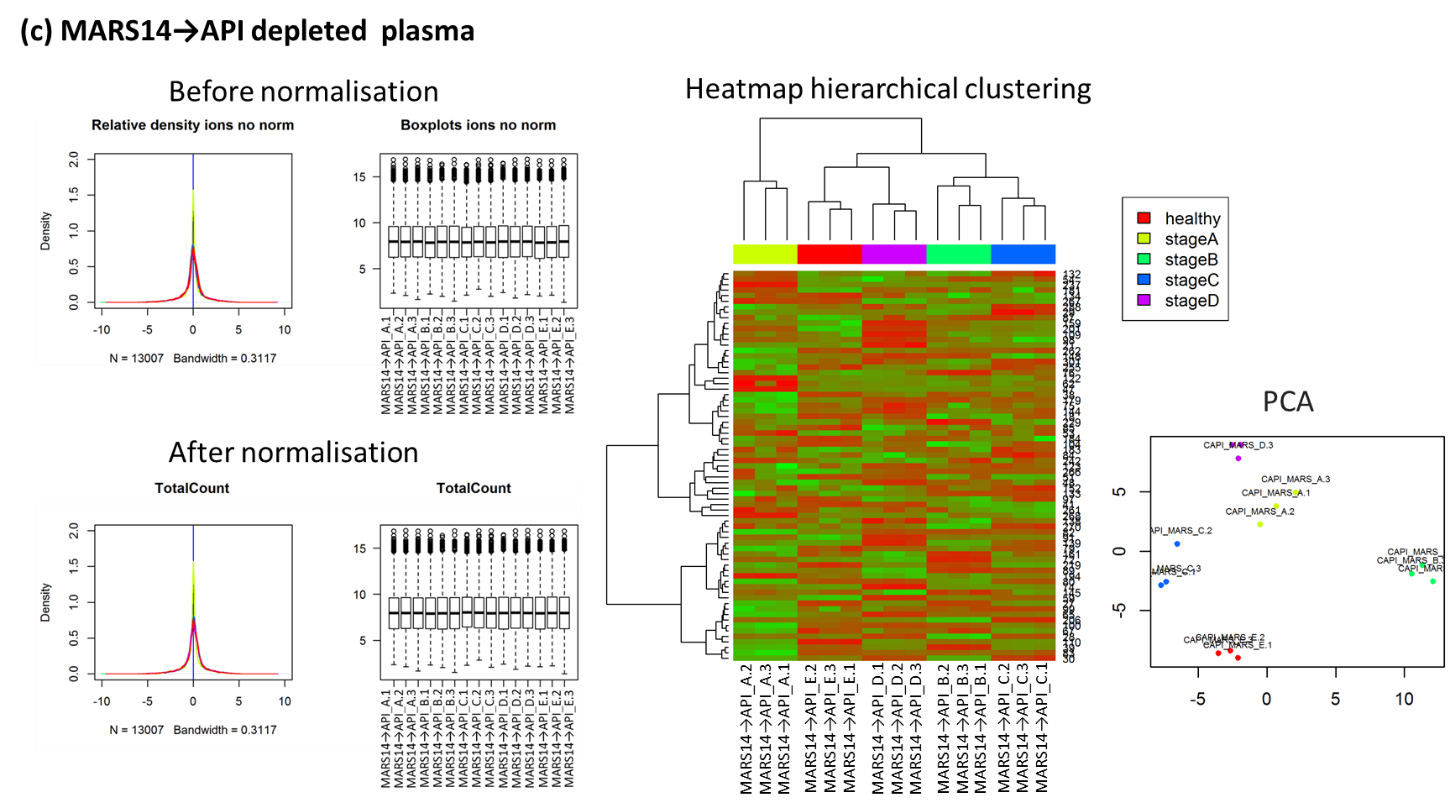
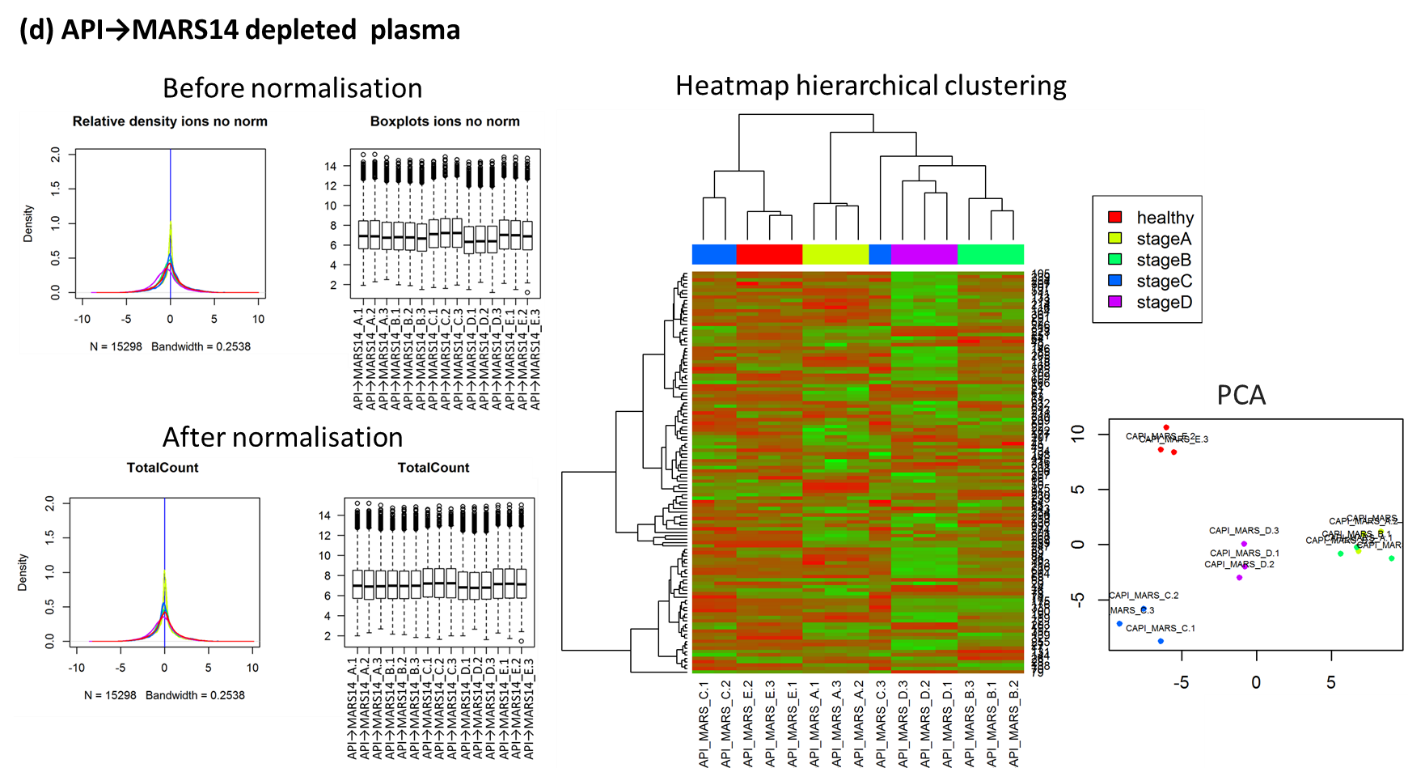
Additional Information contains:

Additional Figures S1-S4

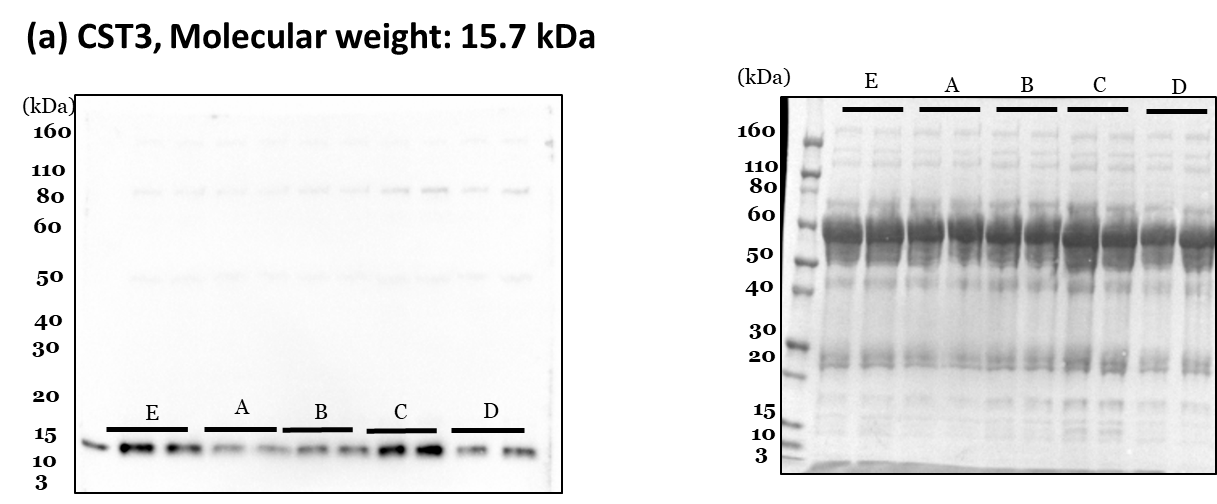


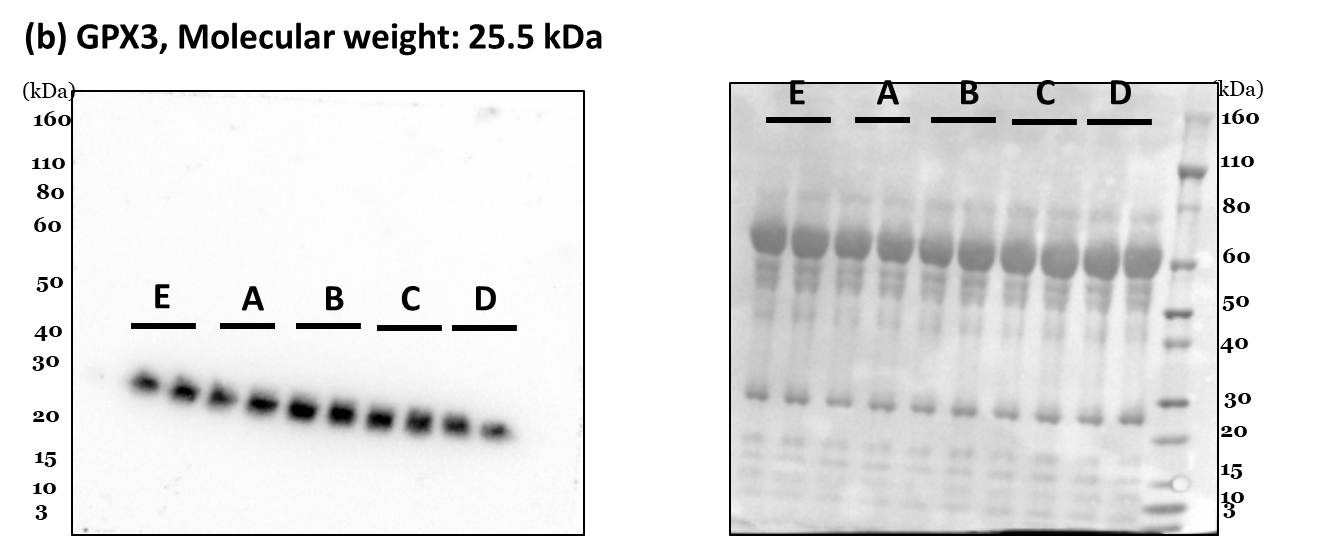
**Figure S1**: Venn diagram comparison of number of common, unshared and shared identified proteins (containing >2 uniquely mapping non-nested peptides of amino acid length >9) between four peptide fractionation methods. **HpH**: High pH C18 reversed phase, **SEC**: Size exclusion chromatography, **SAX**: Strong anion exchange, **SCX**: Strong cation exchange.

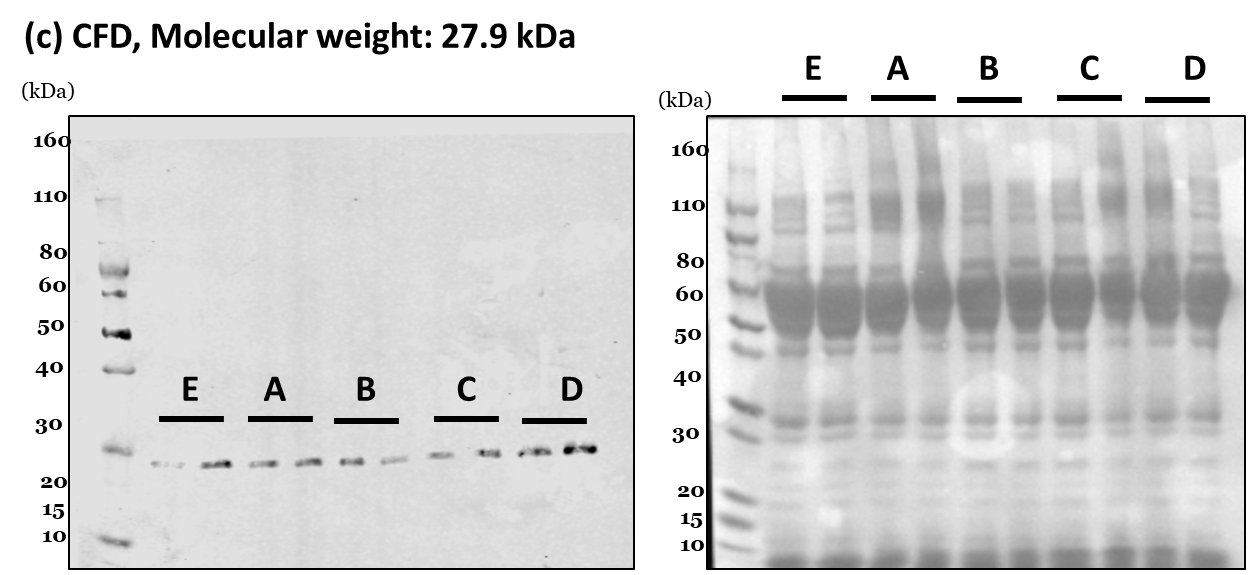


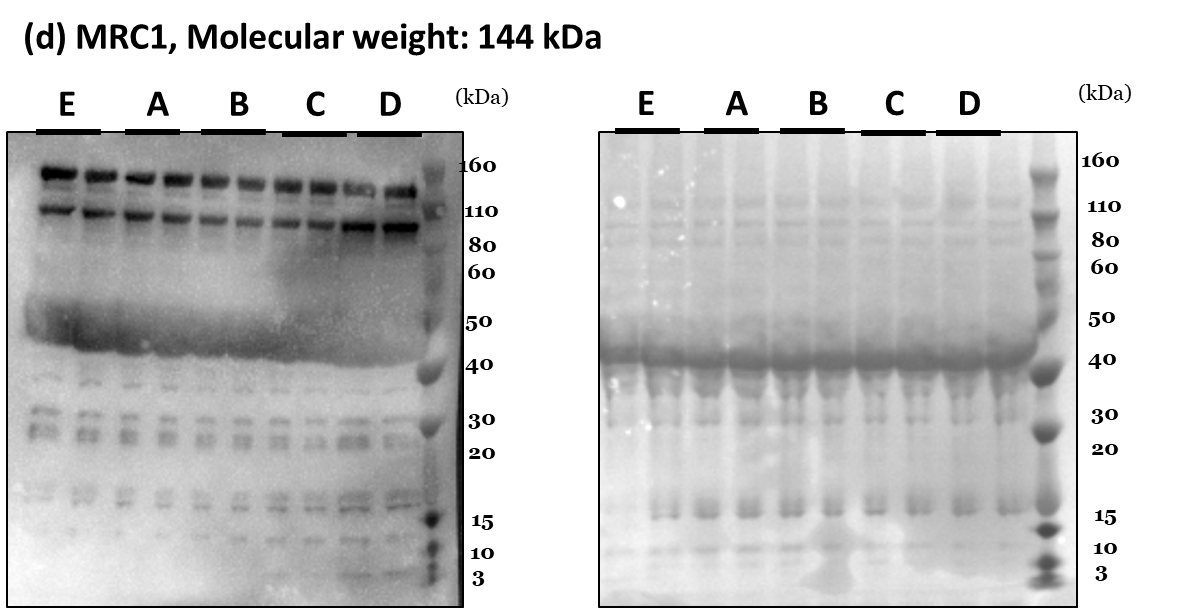


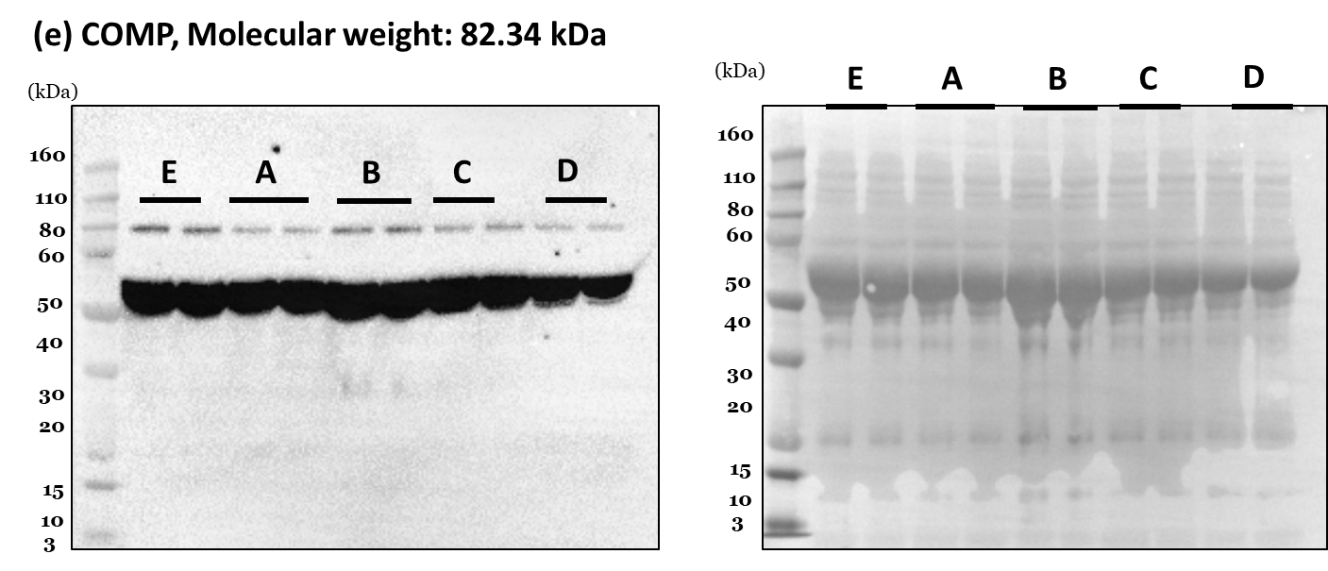
**Figure S2**: Extracted SWATH dataset from **(a)** non-depleted, **(b)** MARS-14 depleted, **(c)** MARS-14→API depleted **(d)** API→MARS-14 depleted experiments were independently normalized using total area normalization. The data distribution was examined using density plots and boxplots. The consistency of the sample replication was examined visually using heatmap hierarchical clustering and PCA plots. Healthy controls represented as stage E in the images. PCA: Principal component analysis.

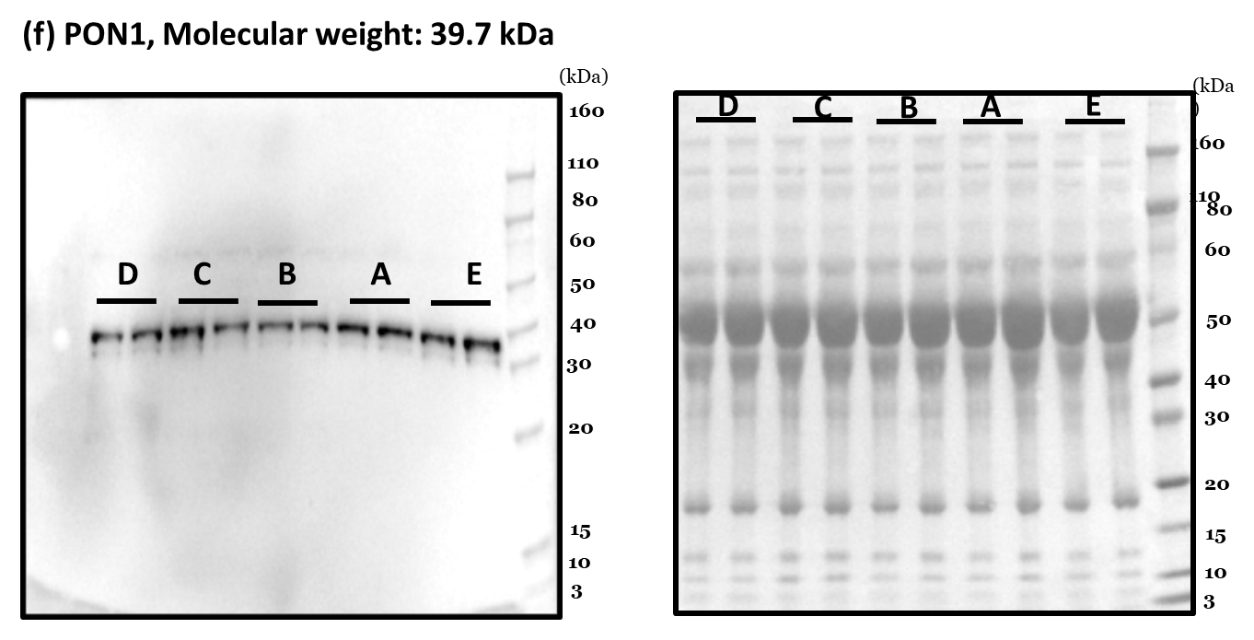




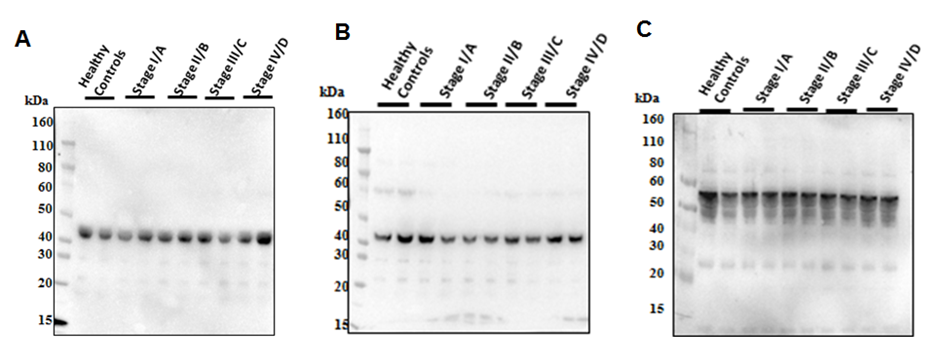
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**Figure S3**: Western blotting images (left) and Ponceau stained images (right) of **(a)** Complement factor D, CST3, **(b)** Glutathione peroxidase 3, GPX3, **(c)** Complement factor D, CFD, **(d)** Macrophage mannose receptor 1, MRC1, **(e)** Cartilage oligomeric matrix protein, COMP and **(f)** Serum paraoxonase/arylesterase 1, PON1.



**Figure S4**: Isotype controls for western blots. (a) IgG controls for CST1, MRC1 and COMP (polyclonal Rabbit, IgG). (b) IgG control for CFD and GPX3 (polyclonal goat, IgG). (c) IgG1 control for PON1 (monoclonal rabbit, IgG1).