

Extensive heterogeneity of glycopeptides in plasma revealed by deep glycoproteomic analysis using size-exclusion chromatography

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Supplementary figure 1

A. All glycopeptides were manually divided into broad glycan categories, complex, hybrid, high-mannose and truncated. Distribution of all identified glycopeptides is shown. **B.** Unique glycoforms identified for 15 most glycosylated proteins is shown.

Supplementary figure 2

A. Unique glycopeptides identified by lectin-affinity chromatography (LAC-bRPLC) and size-exclusion chromatography (SEC) are compared in a Venn diagram. **B.** unique glycan compositions combined from all glycopeptides identified are compared between LAC-bRPLC and SEC.

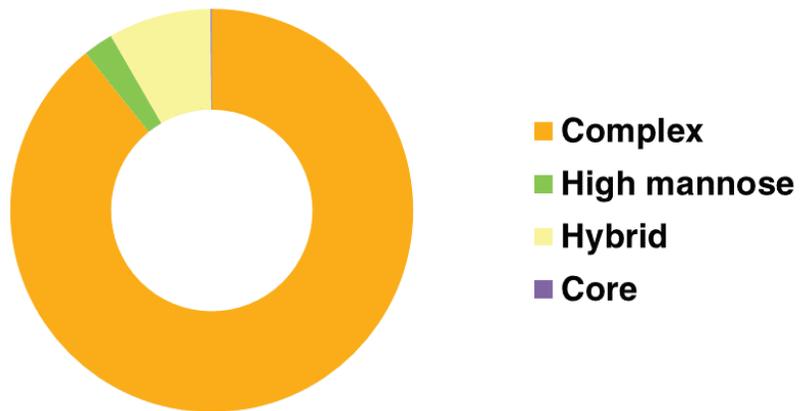
Supplementary figure 3

Venn diagram comparing this study's unique glycopeptide identification with previously published studies (Zhang et al.¹ and Sun et al.²). Sun et al.'s study was conducted on serum while this study and Zhang et. al. was conducted on plasma.

References

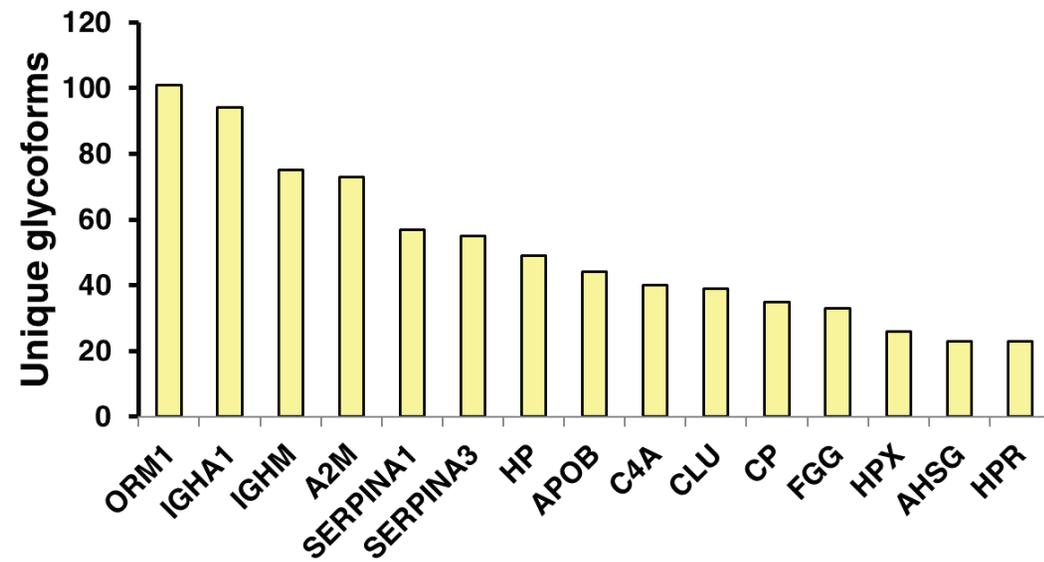
1. Y. Zhang, Y. Mao, W. Zhao, T. Su, Y. Zhong, L. Fu, J. Zhu, J. Cheng and H. Yang, *J Proteome Res*, 2020, **19**, 655-666.
2. S. Sun, Y. Hu, L. Jia, S. T. Eshghi, Y. Liu, P. Shah and H. Zhang, *Anal Chem*, 2018, **90**, 6292-6299.

A.



All glycopeptides by glycan composition

B.



A

LAC-bRPLC

SEC

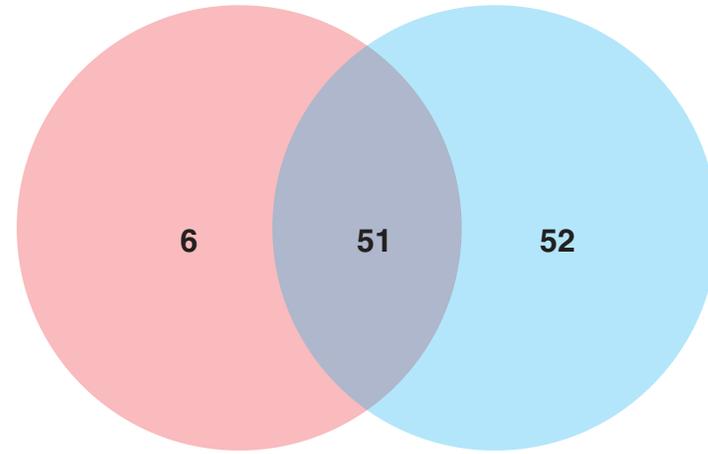


All glycopeptides by glycan composition

B

LAC-bRPLC

SEC



All glycan compositions

