

Lipopolysaccharide and Statin Mediated Immune-responsive Protein Networks in Macrophage Revealed Through Affinity Purification Spacer-Arm Controlled Cross-linking (AP-SPACC) Proteomics

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Supplementary Information

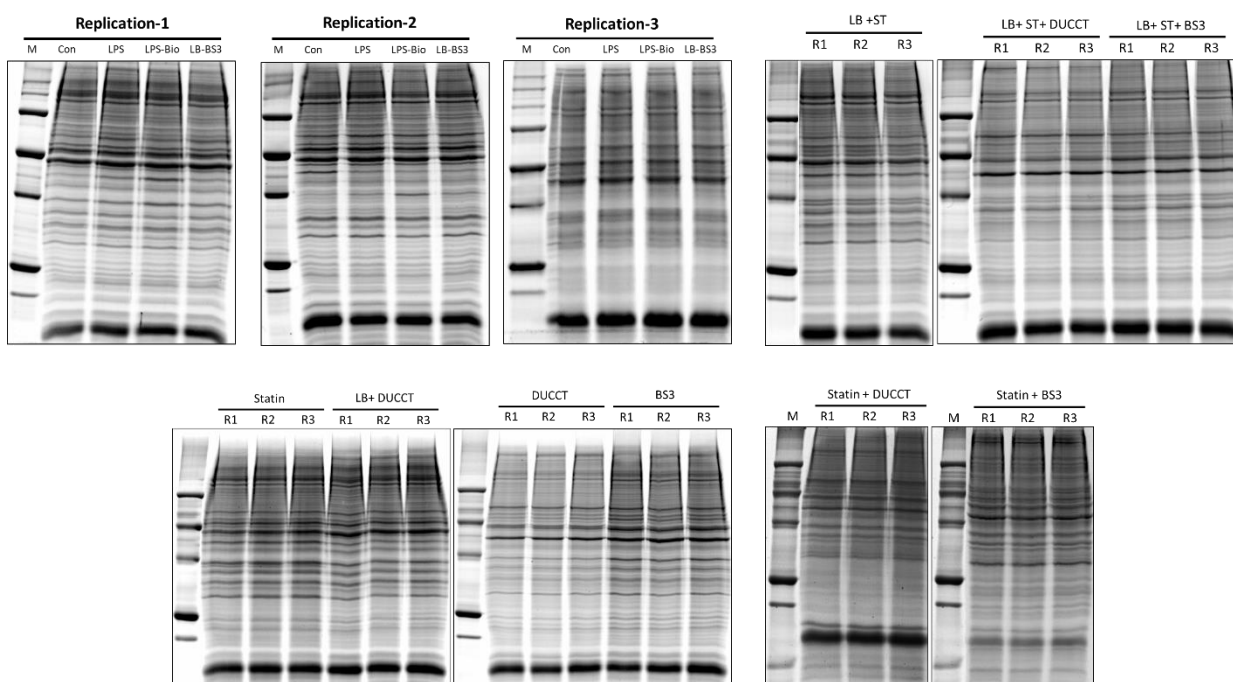


Figure S1 Representative SDS-PAGE gels with the treatment of bio-LPS, statin-bio-LPS, and statin along with cross-linkers in Raw 264.7 macrophage cells.

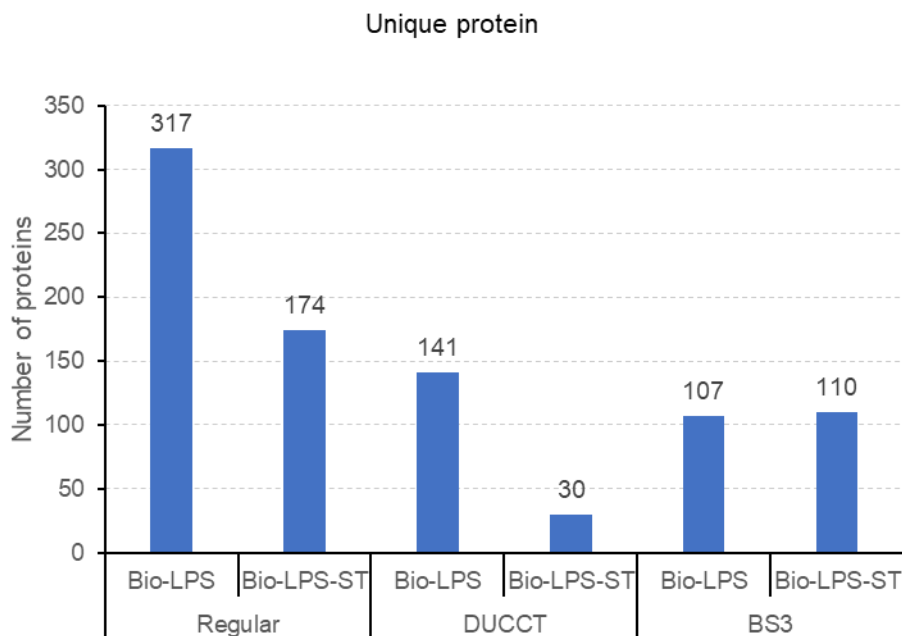
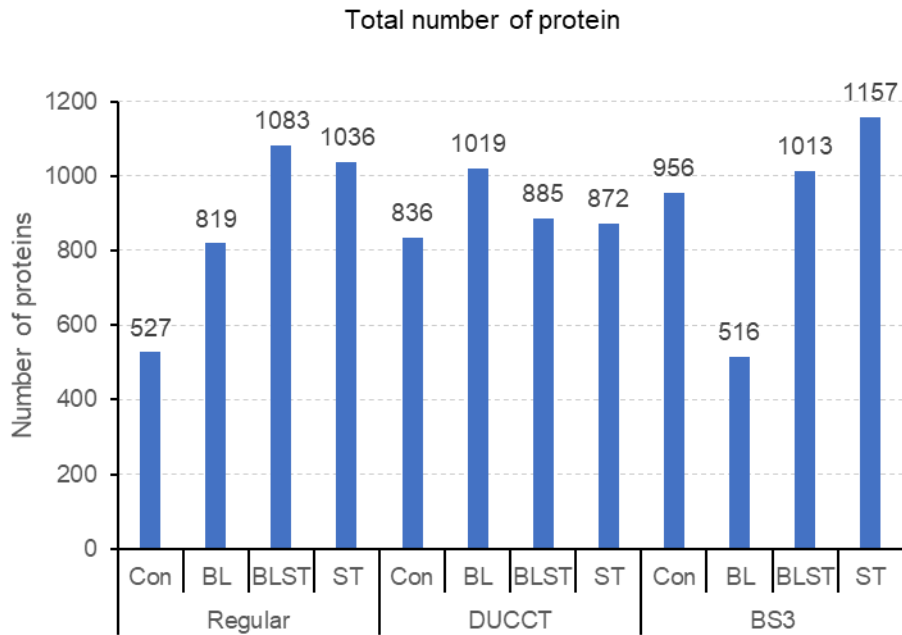


Figure S2 Distribution of identified proteins in RAW 264.7 macrophages during the treatment of bio-LPS, statin-bio-LPS, and statin along with cross-linkers using gel-based affinity proteomics coupled to mass spectrometry.