

Global Proteomics Analysis Of Circulating Extracellular Vesicles Isolated From Lung Transplant Recipients

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Supporting Information

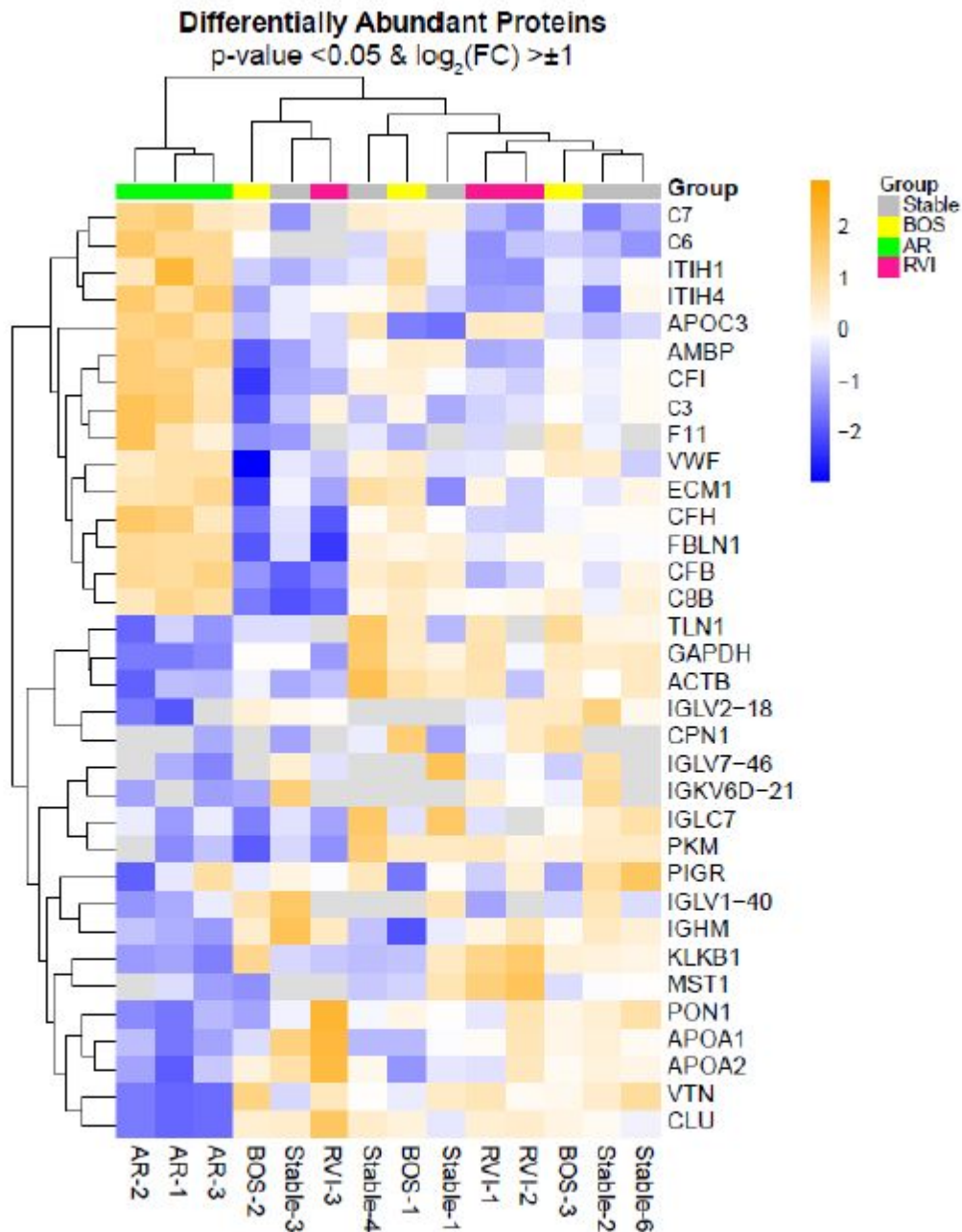


Figure S1: Unsupervised clustering of lung transplant conditions from 35 significant proteins in lung transplant recipients with bronchiolitis obliterans syndrome (BOS), acute rejection (AR), and respiratory viral infection (RVI) compared to stable lung transplant recipients. The AR patients show clearly distinct protein profiles for the 31 differentially abundant proteins compared to the rest of the LTxR groups.