

Supporting Information for Publication

Deep Proteomic Deconvolution of Interferon and HBV Transfection Effects on a Hepatoblastoma Cell Line

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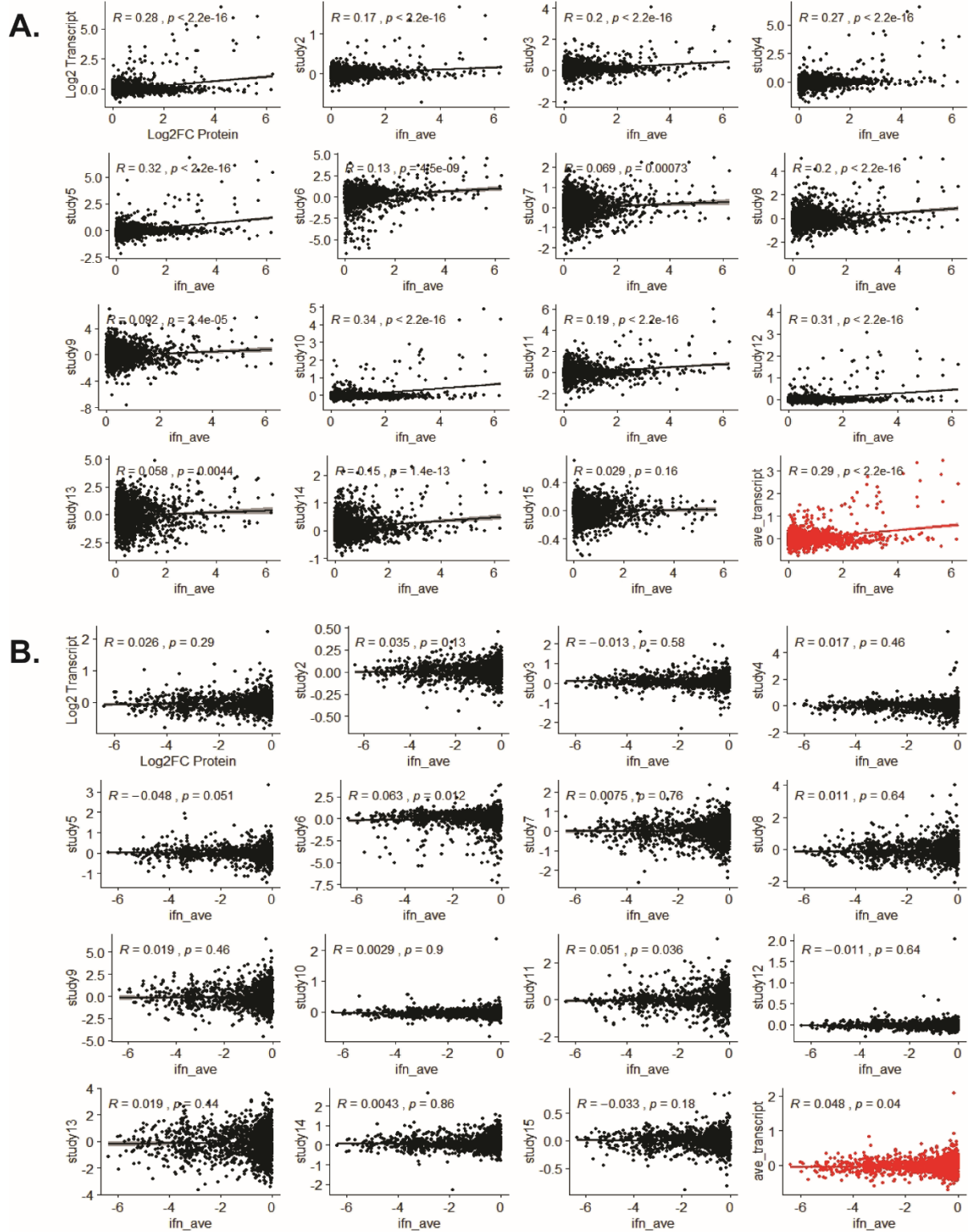


Figure S1. Pearson correlation analysis in which either upregulated (A) or downregulated (B) proteins were excluded in protein/transcript comparisons.

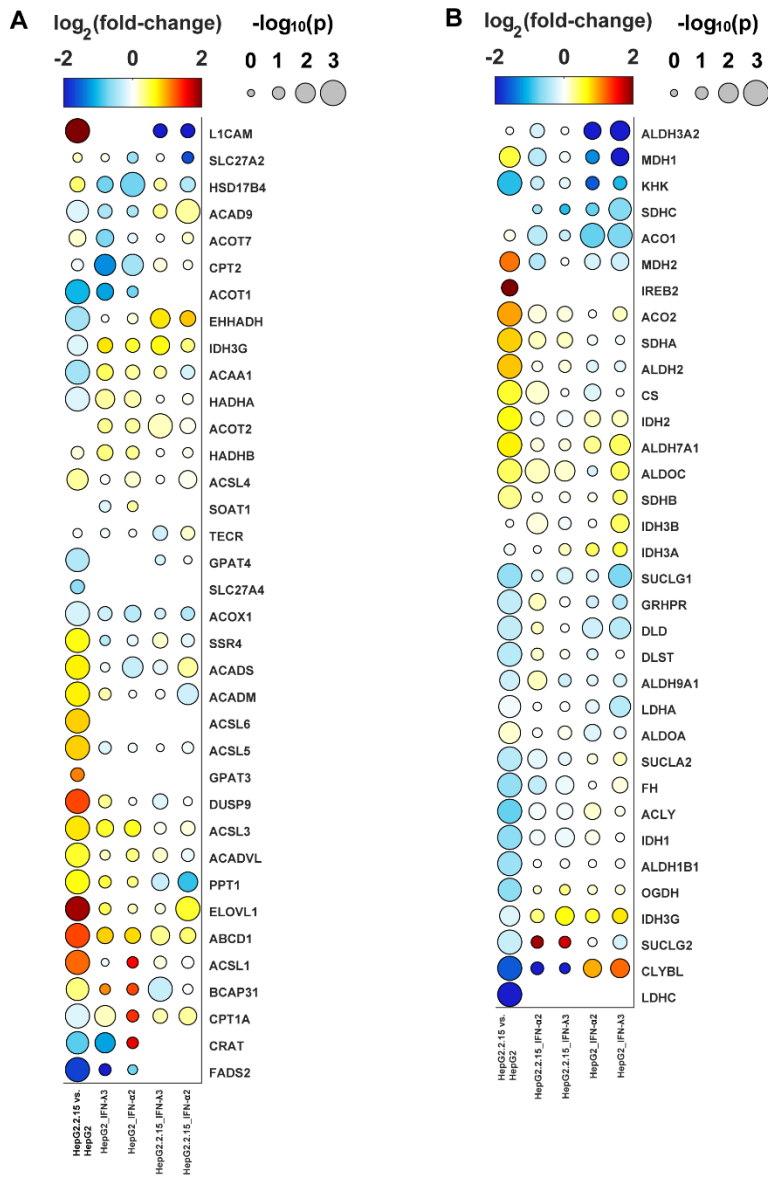


Figure S2. (A) Comparison of expression changes in proteins associated with acyl-CoA metabolites (13Z)-eicosenoyl-CoA, arachidonoyl-CoA, or linoleoyl-CoA. Circle colors indicate \log_2 fold-change value, and the size represents the fold-change significance (p-value). Missing circles correspond to proteins that were not sufficiently detected. (B) Comparison of expression changes in proteins associated with the TCA cycle. Circle colors indicate \log_2 fold-change value, and the size represents the fold-change significance (p-value). Missing circles correspond to proteins that were not sufficiently detected.