

Supplementary Figure S1. KEGG pathways and Gene ontology analysis of differentially expressed plasma exosome miRNAs. Heatmaps show the top most statistically significant KEGG pathways (A) and Gene Ontology terms (B) associated with the four miRs (miR-425-5p, miR-425-3p, and miR-339-3p, miR-93-5p).