

Article

Circulating Neutrophil Profiles Undergo a Dynamic Shift during Metabolic Dysfunction-Associated Steatohepatitis (MASH) Progression

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Supplementary Figures

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Supplemental Figure 1

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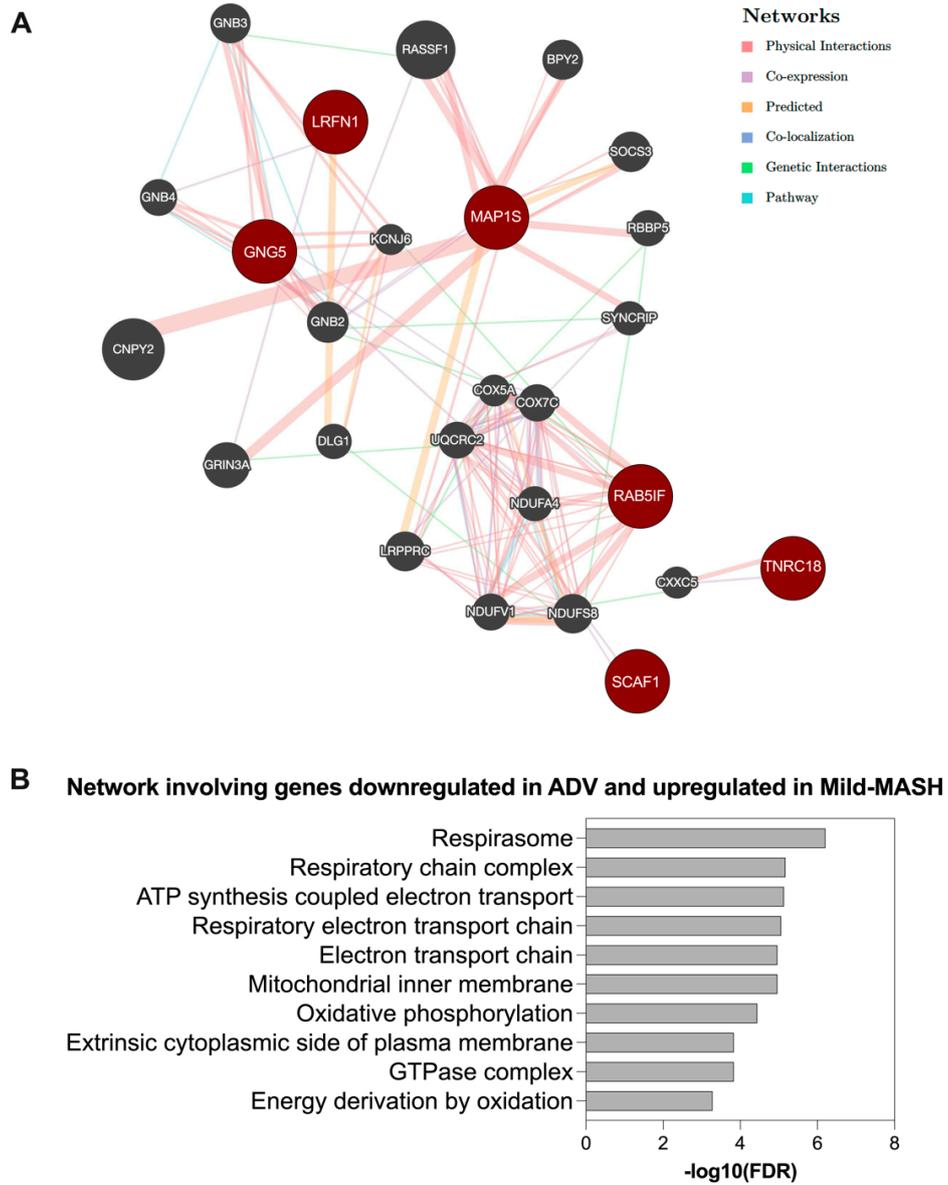


Figure S1. Identification of genes that shift expression in neutrophils from mild to advanced MASH patients. **(A)** Network analysis using GeneMANIA showed that from the 9 genes identified in our dataset, 6 genes are part of a larger network. Legend indicates the type of connection among the genes. **(B)** Pathway enrichment of genes involved in the network from Figure A.

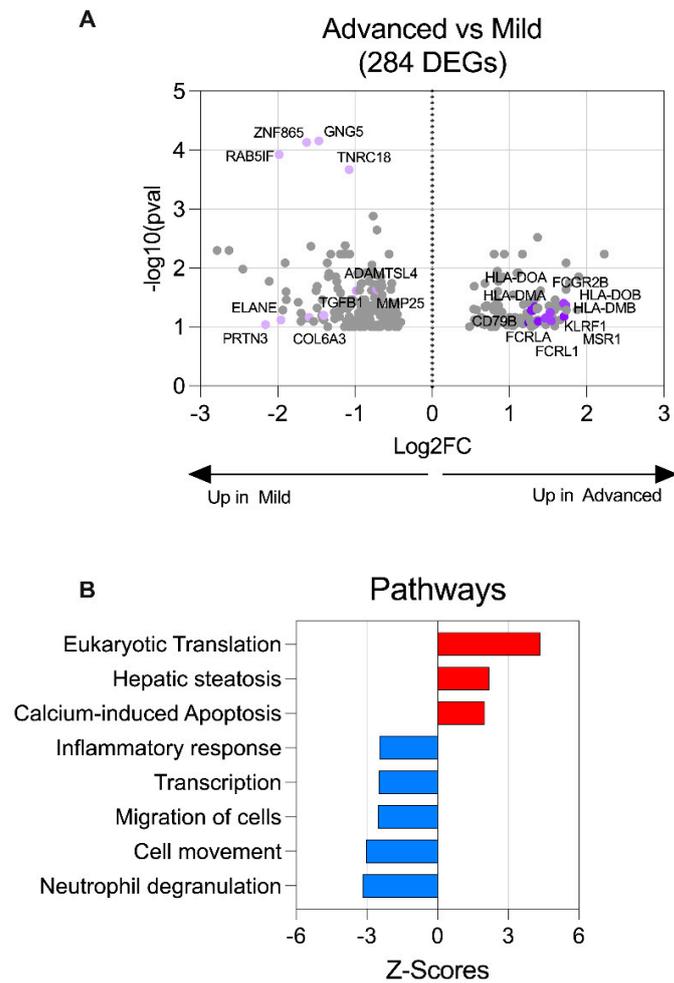


Figure S2. Direct comparison between advanced and mild MASH. **(A)** Volcano plot showing the differentially expressed genes obtained from statistical comparison of advanced and mild transcriptome groups. Arrow to the left: genes upregulated in Mild x Adv. Arrow to the right: genes upregulated in Adv x Mild. **(B)** Pathways enrichment analysis showing pathways upregulated in Mild x Adv (blue) and Adv x Mild (red).