

*Article*

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

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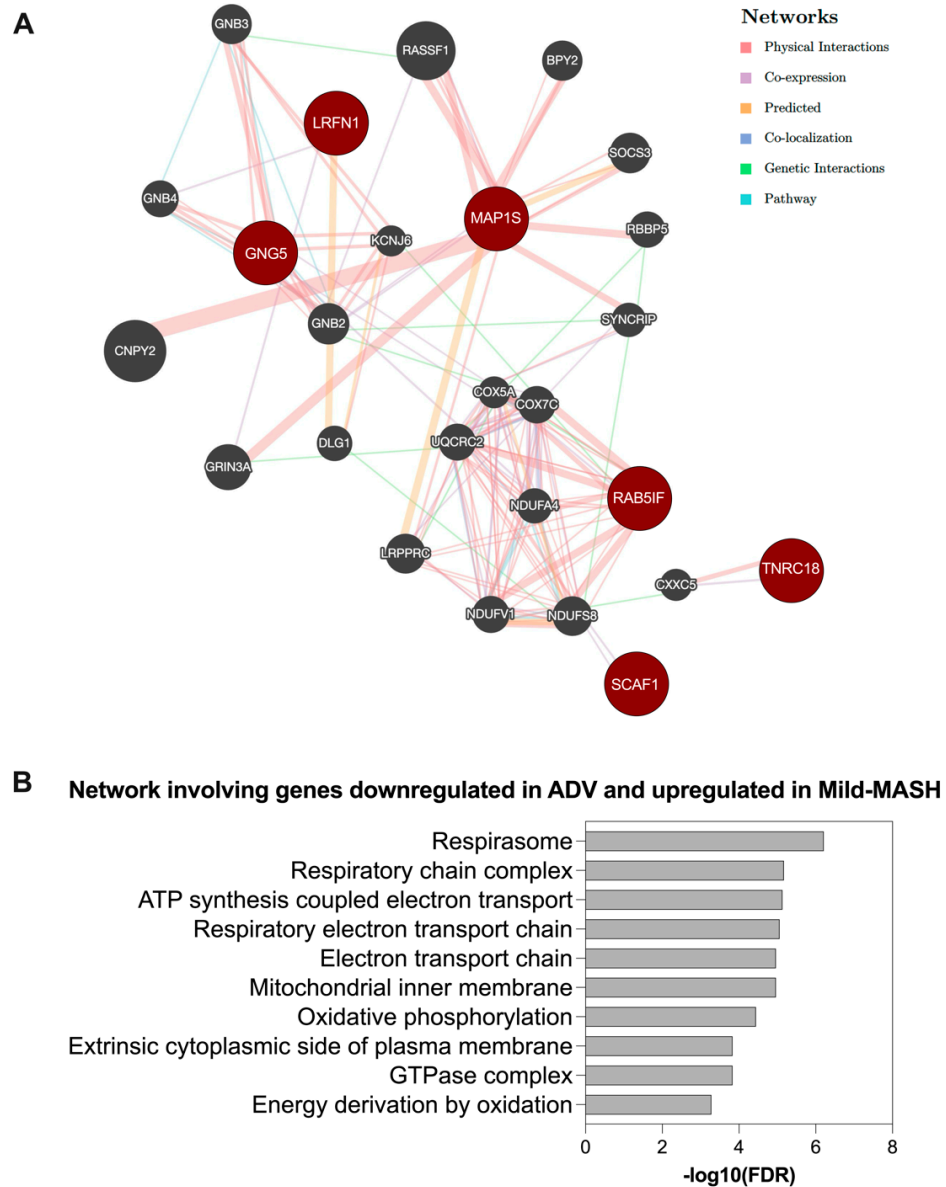
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## **Supplementary Figures**

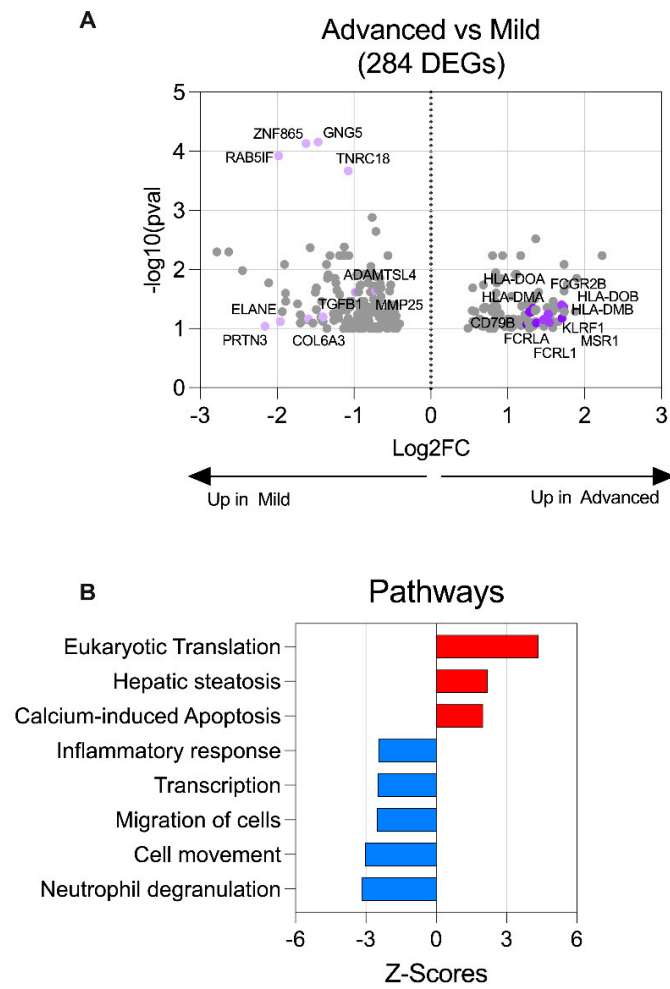
### **Content List:**

Supplemental Figure 1

Supplemental Figure 2



**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).