

Supplementary Figures:

Identification of macrophage-related genes in sepsis-induced ARDS using bioinformatics and machine learning

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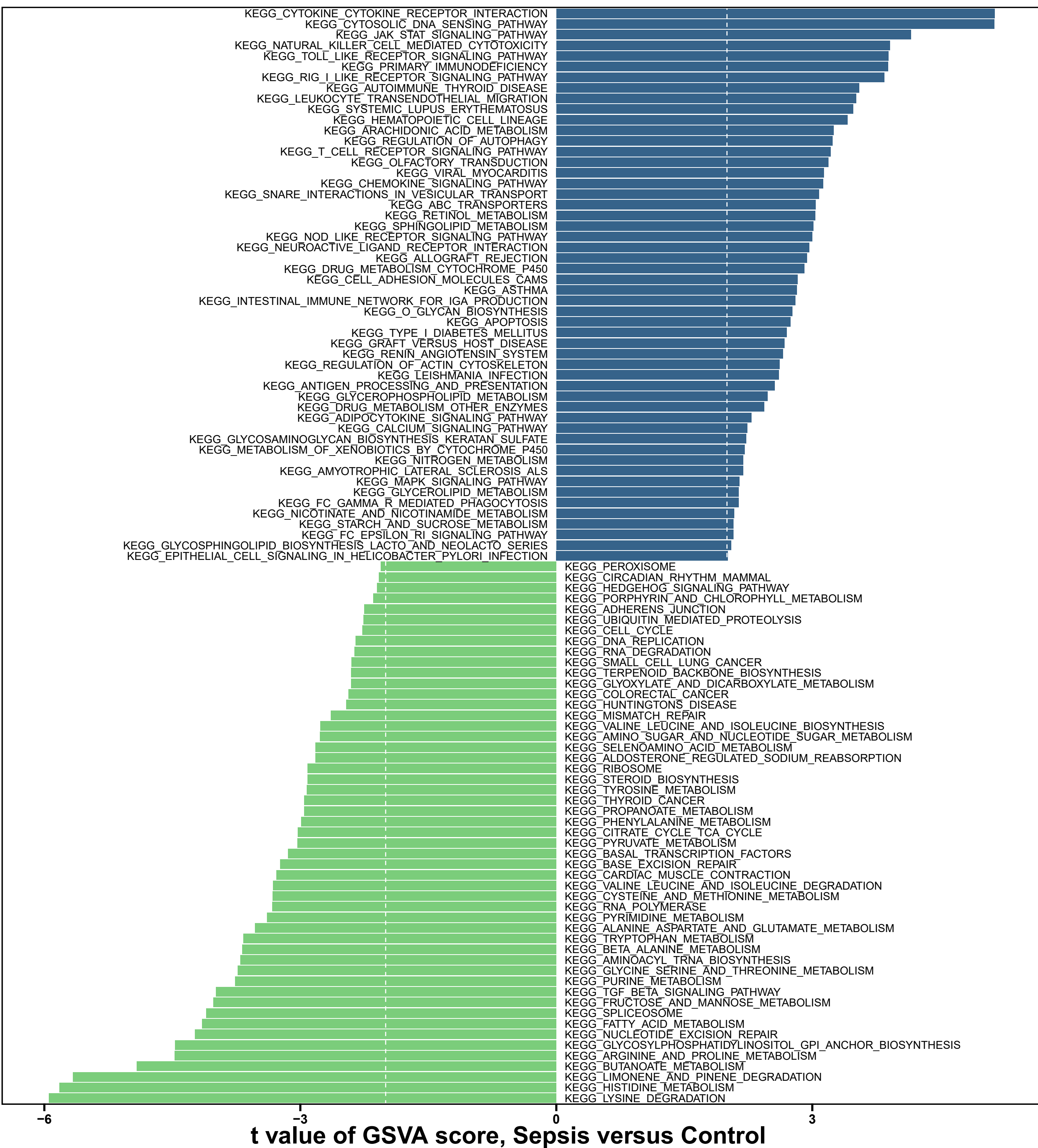
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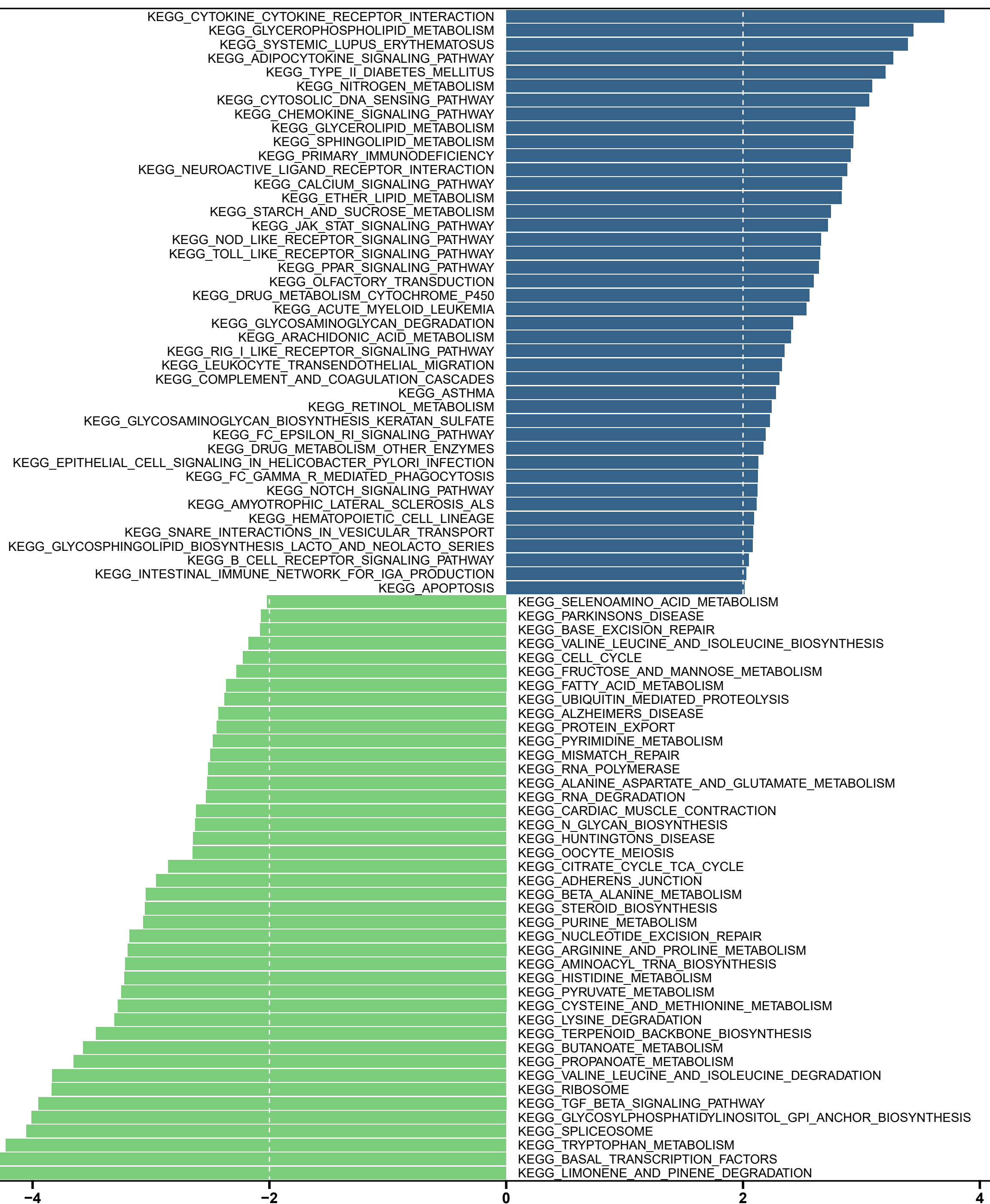
Figure legend:

Supplementary Figure 1. GSVA enrichment pathway differences between the sepsis group and the control group in GSE32707.

Supplementary Figure 2. Differences in enriched pathways according to GSVA between the sepsis-induced ARDS group and the control group in GSE32707.

Supplementary Figure 3. miRNA prediction. A. Database prediction of SGK1 target miRNAs in the intersection of the Venn diagram. B. Database prediction of MSRB1 target miRNAs in the intersection of the Venn diagram. C. Database prediction of DYSF target miRNAs in the intersection of the Venn diagram. D. Correlation diagram between SGK1, MSRB1, DYSF and miRNAs.





t value of GSVA score, sepsis-induced ARDS versus Control

