# Exploration and validation of a combined immune and metabolism gene signature for prognosis prediction of colorectal cancer

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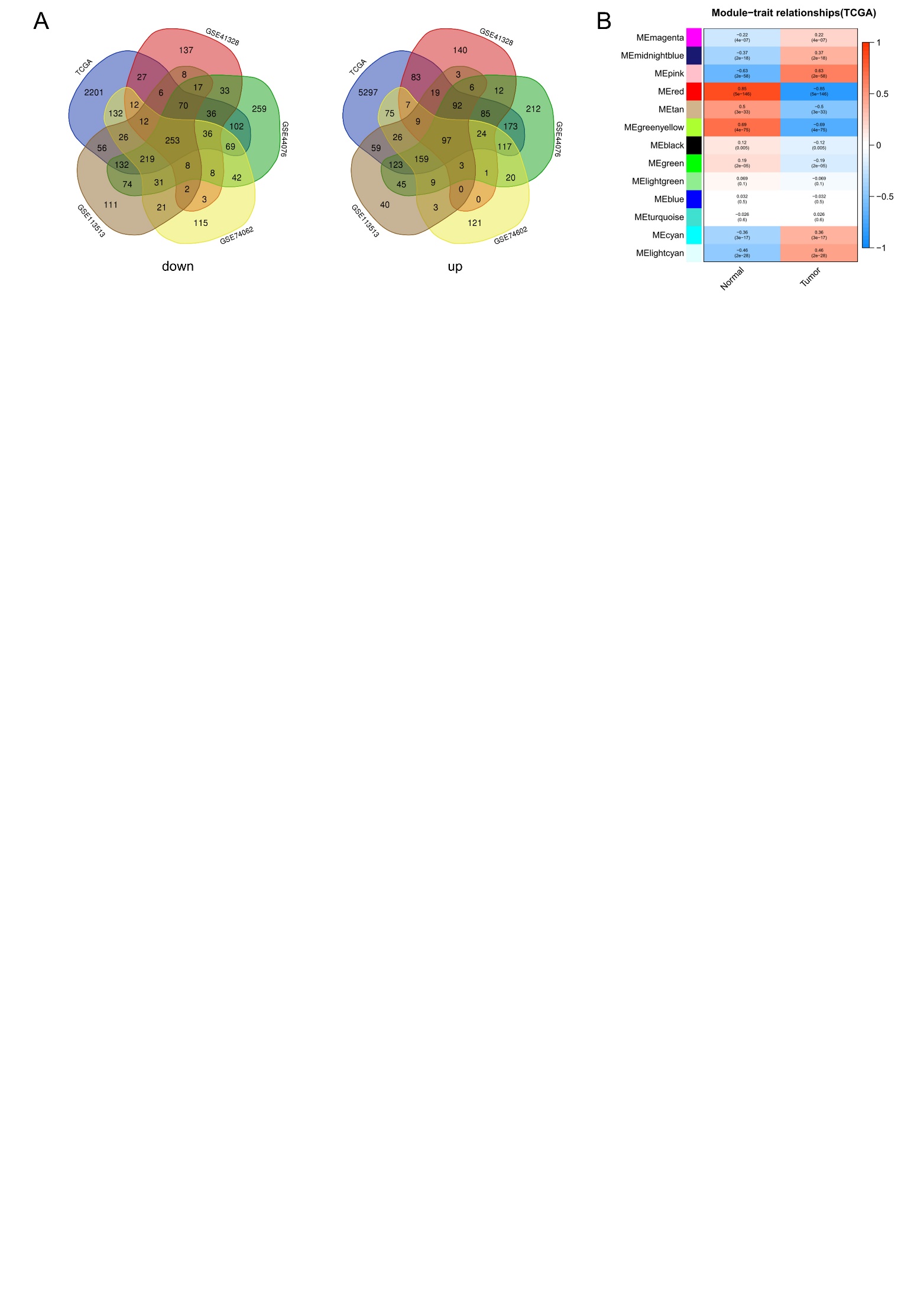
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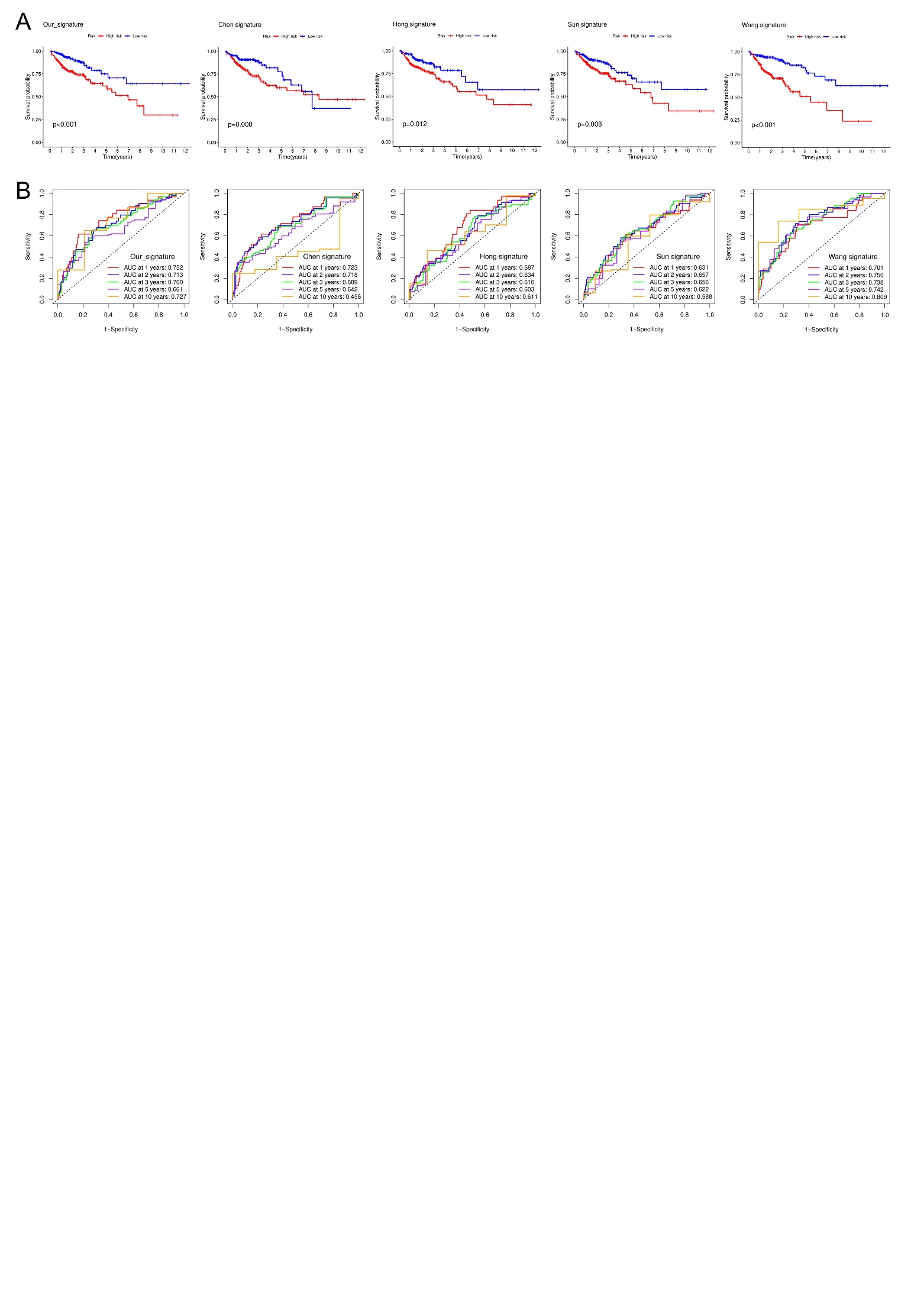
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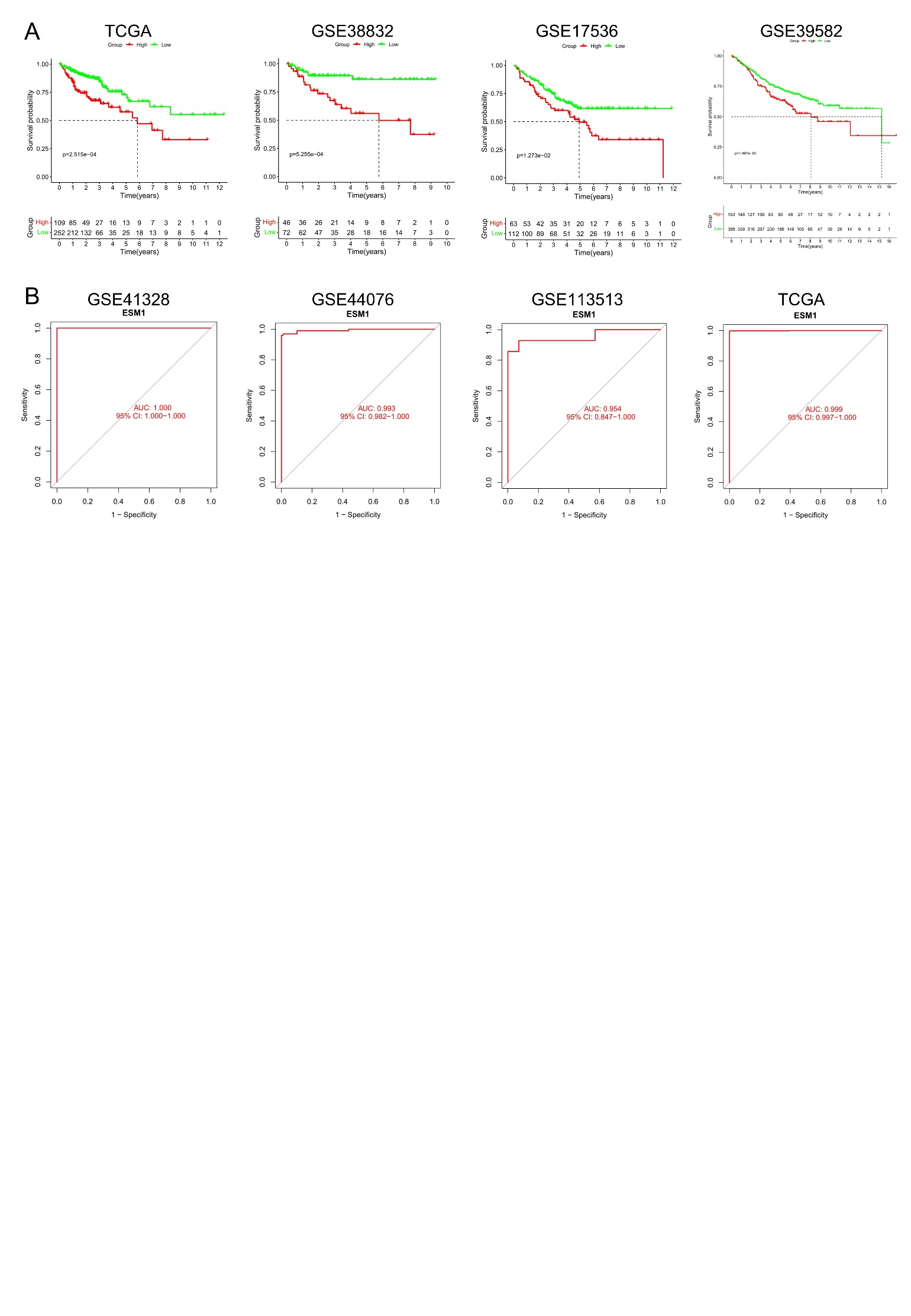
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**Figure S1** (A) Venn diagram of the downregulated and upregulated differentially expressed genes (DEGs) among CRC patients. (B) Relationship between the modules and the tumor or nomal. Different color blocks represent different clusters of genes. Color shades represent the correlation.



**Figure S2** (A) Kaplan-Meier OS curves of our signatures and other existing signatures. (B) ROC curves of our signatures and other existing signatures.



**Figure S3** (A) Kaplan-Meier curves of ESM1 in different datasets. (B) ROC curves of ESM1 to diagnose CRC in different datasets

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