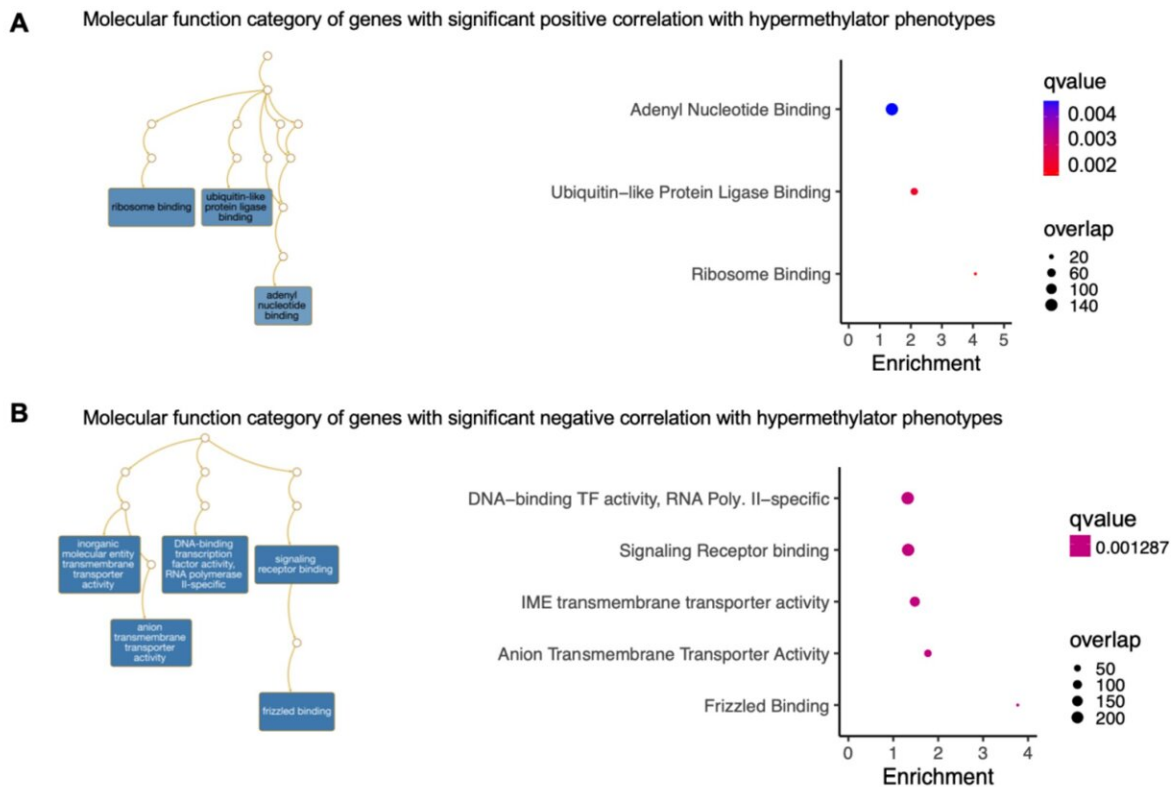


Transcription factors found to contribute to subtypes of colorectal cancers

July 27 2023



Molecular function categories of genes correlated with hypermethylator phenotypes. Gene ontology molecular function categories of the top protein coding genes (Benjamini-Hochberg adjusted p-value < 0.05) whose expression is positively (1115 genes) (A) or negatively (1813 genes) (B) correlated (Spearman rank-order correlation test) with the hypermethylator subtypes of CRCs. The left panels show the Directed Acyclic Graph (DAG) outputs summarizing the key nodes representing the enriched pathways that are significantly enriched (FDR

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