



S9 Fig. Gene Ontology (GO) enrichment analysis results for the identified gene sets in pediatric acute myeloid leukemia (AML). This figure illustrates the results of the enrichment analysis for high-risk and low-risk gene signatures in pediatric acute myeloid leukemia. The enrichment encompasses three primary GO categories: Molecular Function, Biological Process, and Cellular Component. These enriched GO terms highlight the intricate interplay of genetic regulation, immune response, and cellular interactions in pediatric AML, providing insights into potential therapeutic targets and the biological foundations of risk stratification.