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A 29-gene and cytogenetic score for the prediction of resistance to
induction treatment in acute myeloid leukemia

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Contributions: T.H., U.M. and K.S. conceived and designed the experiments. T.H., M.R.-T., B.K., L.H. and K.H.M. performed experiments. T.H., V.J., A.M.N.B., S.A.B., M.R.-T. and K.H.M. analyzed data. V.J., A.M.N.B., S.A.B. and U.M. provided bioinformatics support. J.P.-M., S.K. and H.B. managed the Genome Analyzer Iix platform and the RNA sequencing. M.R.-T., B.K., L.H., P.A.G., S.S., N.K., J.T., M.S., J.B., S.K.B. and K.S. characterized patient samples; M.C.S., D.G., J.B., S.A., W.E.B., B.J.W. and W.H. coordinated the AMLCG clinical trials. U.M. and K.S. supervised the project. T.H., V.J., S.A.B., K.H.M. and S.K.B. wrote the manuscript.