Clone-specific secondary aberrations are not detected in neonatal blood spots of children with ETV6-RUNX1-positive leukemia

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MM designed and performed experiments, interpreted data and compiled results. CM and RM determined genomic breakpoints of *ETV6-RUNX1*. GM was responsible for the clinical study, provided patient samples and clinical data. OAH interpreted data, contributed to study design and wrote the paper. RG-P conceived and supervised the study, interpreted data and wrote the manuscript. All authors read and agreed to the final version of the manuscript.