Rare coincident NPM1 and RUNX1 mutations in intermediate risk AML display similar patterns to single mutated cases

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<td>First author</td>
<td>Fasan, Annette</td>
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Specific contributions of each author are listed on page 2 of this document.

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AF analyzed the data and wrote the manuscript; CH was responsible for chromosome analysis, AK, FD and CE performed molecular analyses; WK was responsible for immunophenotyping and was involved in statistical analyses; TH was responsible for cytomorphologic analysis and was involved in the collection of clinical data. SS was the principle investigator of the study and wrote the manuscript. All authors read and contributed to the final version of the manuscript.