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author contribution form

Title	Archival bone marrow trephines are suitable for high-throughput mutation analysis using next generation sequencing technology
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Specific contributions of each author are listed on page 2 of this document.

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Please describe the contributions of each author, indicating who was responsible for each part of the study and the preparation of the manuscript (collection of data, experiments, data analysis, generation of figures, collection of images, interpretation of data, preparation of the text, etc.)

RG and UL designed the study, including selection of target genes and design of enrichment probes

BH and RG performed the sequencing, including DNA isolation and library preparation.

BH, RG and UL processed the raw data and evaluated the sequence variances with crucial support from SB.

BS and HK provided crucial support in the design of the study, selection of patient samples and interpretation of the results.

UL wrote the manuscript with support from BH, RG, SB, and HK.

All authors approved the final version of the manuscript.