Title | Comprehensive whole-genome sequencing of an early-stage primary myelofibrosis patient defines low mutational burden and nonrecurrent candidate genes
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Specific contributions of each author are listed on page 2 of this document.

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All authors either drafted the manuscript or critically reviewed it for important intellectual content, and all authors approved the final version of the manuscript to be published. The following individuals also made these contributions:

Jason D. Merker – recruited, consented, and obtained specimens from the patients; performed experiments; data analysis and curation; coordinated the research; generation of figures; interpretation of data; drafted initial version of manuscript
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