## Haematologica HAEMATOL/2015/130799 Version 3

Whole-exome sequencing reveals a C-terminal germline variant in CEBPA-associated acute myeloid leukemia: 45-year follow-up of a large family

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Disclosures: None of the authors of this manuscript have any conflicts of interest to disclose.

Contributions: A.Pathak, KS, TP, A.Pemov, RD, CB, RAG, SR, BTL, MM, SS, MY and the NCI DCEG Cancer Genomics Research Laboratory performed the research. Individuals in the NCI DCEG Cancer Sequencing Working Group contributed control data to the study and revised the manuscript. NC, LG, MLM, and JM revised the manuscript. DRS designed the study. A.Pathak and DRS wrote the manuscript.