

Disruption of the ASXL1 gene is frequent in primary, post-essential thrombocytosis and post-polycythemia vera myelofibrosis, but not essential thrombocytosis or polycythemia vera: analysis of molecular genetics and clinical phenotypes

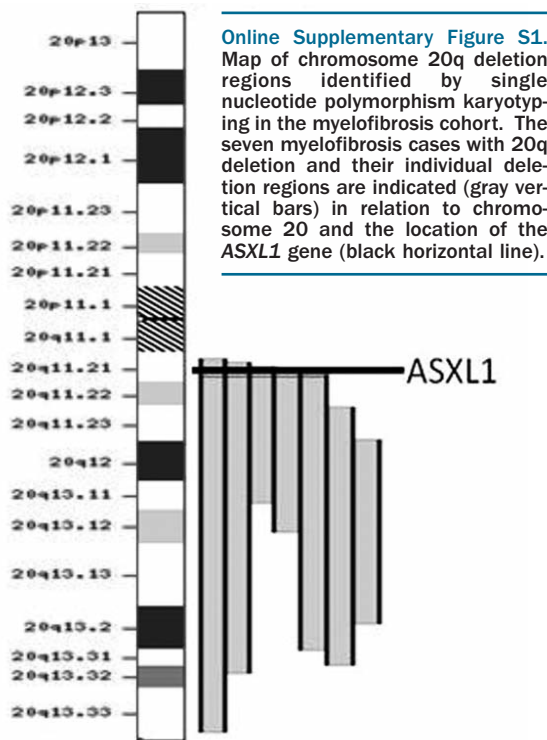
Brady L. Stein,¹ Donna M. Williams,² Christine O’Keefe,³ Ophelia Rogers,² Roxann G. Ingersoll,⁴ Jerry L. Spivak,² Amit Verma,⁵ Jarek P. Maciejewski,³ Michael A. McDevitt,² and Alison R. Moliterno²

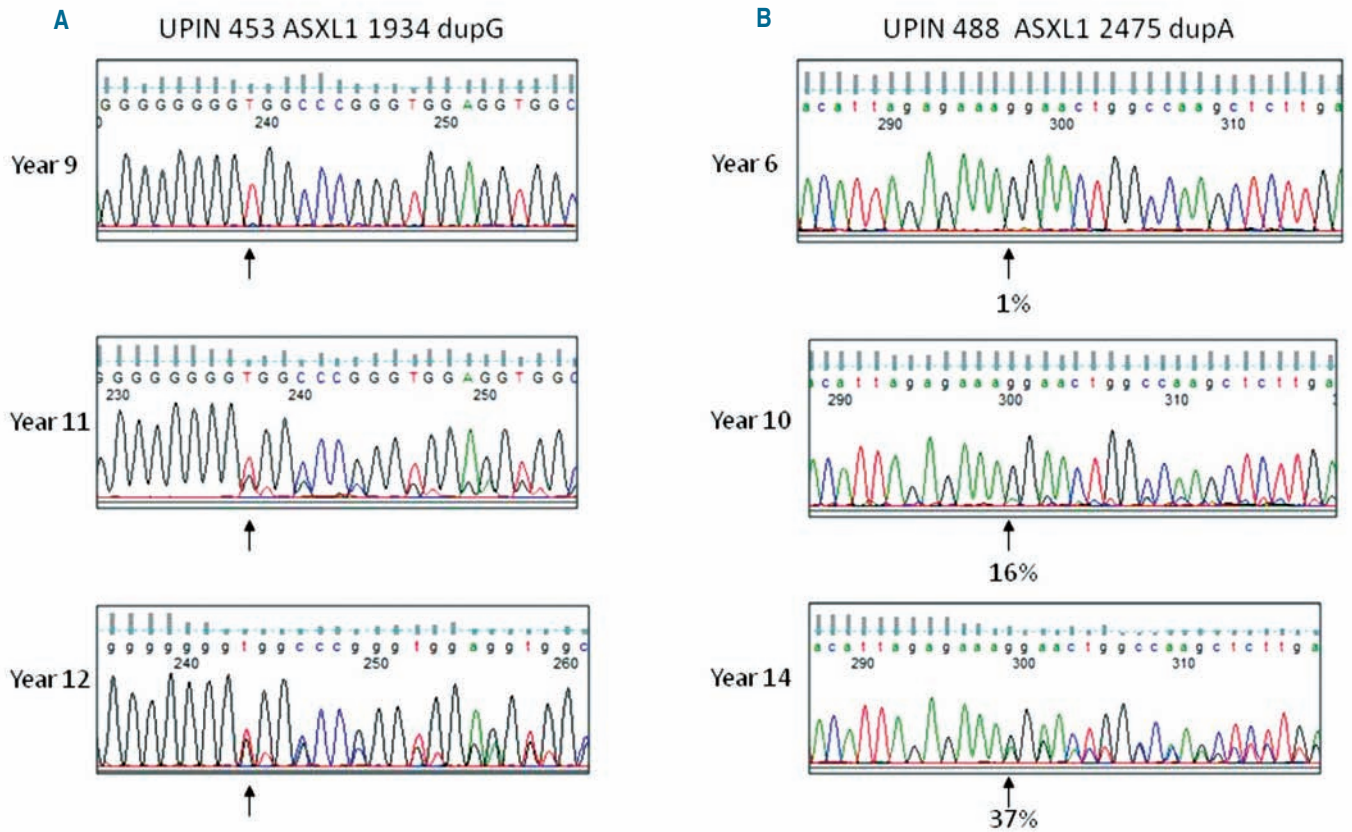
¹Medicine, Northwestern University Feinberg School of Medicine, USA; ²Medicine, Johns Hopkins University School of Medicine, USA; ³Translational Hematology and Oncology Research, Cleveland Clinic/Taussig Cancer Institute, USA; ⁴Genetics Research Core Facility, Johns Hopkins University School of Medicine, USA, and ⁵Albert Einstein College of Medicine, NY, USA

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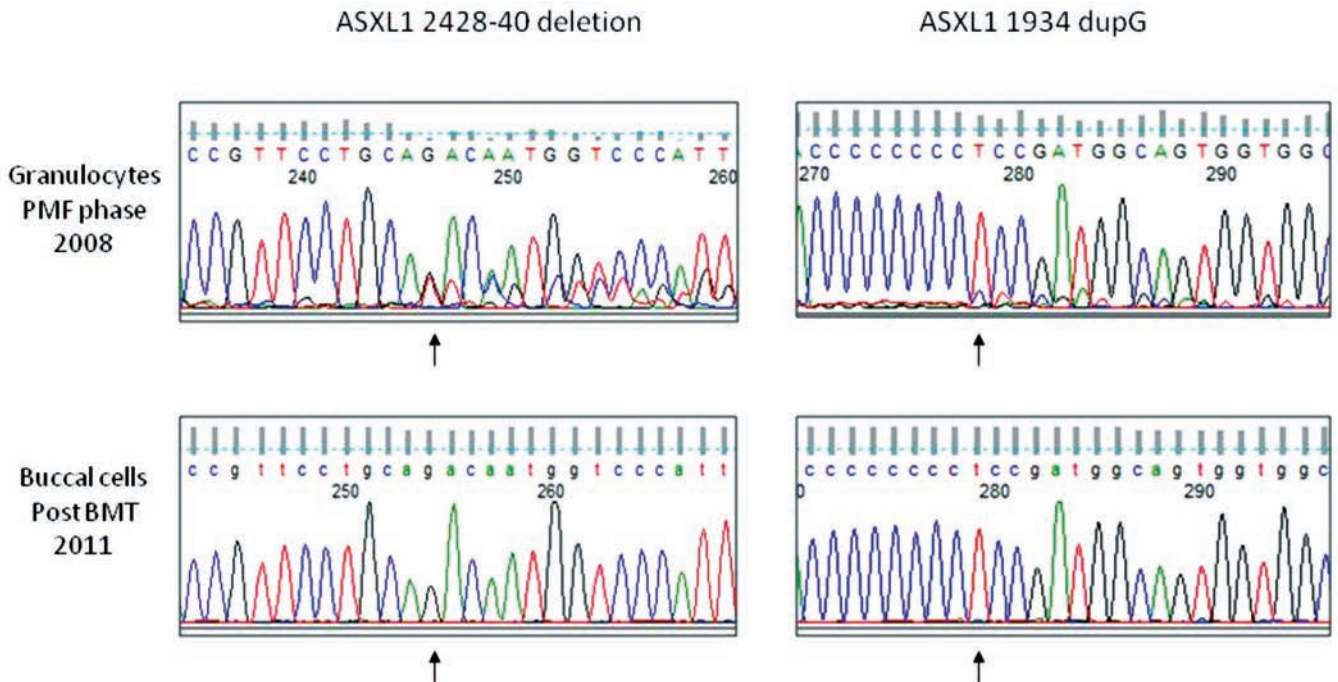
Online Supplementary Table S1. Pyrosequencing primers.

Lesion	Forward Primer	Reverse Primer	Sequencing Primer
JAK2 V617F	GCAGAGAGAATTTTCTGAACTAT	CTCTGAGAAAAGGCATTAGAAAAG	GGTTTTAAATTATGGAGTATGT
ASXL1 1954G->A	AGGTCCGAGGGGCGAGAG	CCACAGGCCTCACCACCAT	GGTGCCCGGGTGGGA
ASXL1 1963-1976del	AGGTCCGAGGGGCGAGAG	CCACAGGCCTCACCACCAT	GGTGCCCGGGTGGGA
ASXL1 2475dupA	ATTCCGTCTCTAGTGGGAGATGA	CCTTCATAGTGGGATGACTGTCAA	GAGATGATACATTAGAGAAA
ASXL1 2846-47del	GCATTGCCTGGGGATTTG	GGCACAGTCCAGAGTGAAGTAAGG	TGACAGCTGAGGAGG





Online Supplementary Figure S2. Electropherograms identifying ASXL1 lesions over time. (A) The ASXL1 lesion 1934 dupG identified in increasing intensity after year 9 in UPIN 453. (B) The ASXL1 nonsense lesion 2475 dupA identified in increasing intensity after year 6 in UPIN 488 (Manuscript Figure 1A). Arrows indicate the start of the sequence variation; percentages indicate the pyrosequencing result of the mutant allele from the same sample which generated the electropherogram in the case of ASXL1 2475 dupA.



Online Supplementary Figure S3. Electropherograms identifying two separate ASXL1 nonsense lesions. Electropherograms from granulocyte DNA from UPIN 1327 obtained in 2008 during the PMF phase of the MPN are shown in the upper panels, with arrows indicating the start of the respective frameshift lesions. Electropherograms from buccal cell DNA obtained from the same patient in 2011, after bone marrow transplantation with donor engraftment and PMF remission (lower panels) do not identify frameshift lesions apparent in the upper panels. The ASXL1 1934 dupG sequence is in reverse orientation.