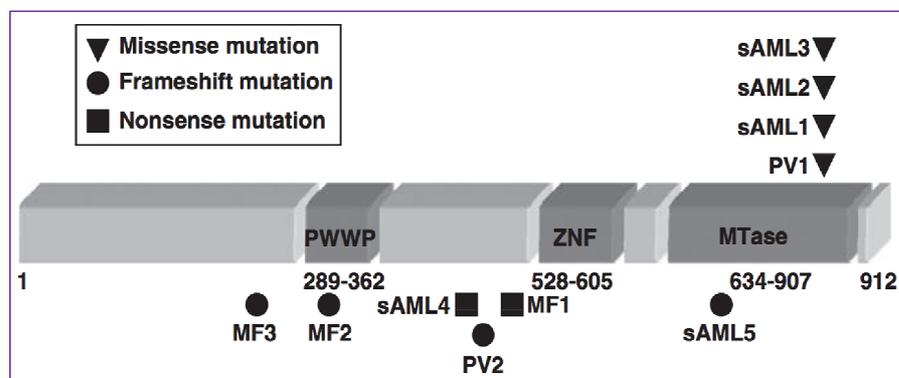


DNA Methyl Transferase 3a (DNMT3a)

DNMT are enzymes that catalyze the transfer of a methyl group on cytosine residues localized in CpG dinucleotides. Four isoforms of DNMT are characterised. DNMT3a is involved in *de novo* methylation of DNA. Methylation of DNA constitutes a repressive mark and decreases gene expression.¹ Abnormalities of DNA methylation are associated with many pathologies, particularly cancer.

In hematological malignancies, *DNMT3A* mutations have initially been described in acute myeloid leukemias^{2,3} (AML) and in myelodysplastic syndroms (MDS).⁴ In myeloproliferative neoplasms (MPNs), *DNMT3A* mutations are infrequent during the chronic phase (<5% in polycythaemia vera and essential thrombocythaemia, 4-7% in primary myelofibrosis).⁵⁻⁸ *DNMT3A* mutations seem to be associated with more advanced disease (secondary myelofibrosis, post-MPN AML).⁵ Heterozygous missense, frameshift or nonsense mutations can be observed. They generally affect regions that encode the methyltransferase domain, arginine 882 being the most affected residue.⁵⁻⁷ *DNMT3A* mutations can be associated with mutations in *JAK2*, *TET2*, *IDH* and *ASXL1*.⁵

The mechanisms linking *DNMT3A* mutations to leukemogenesis remain unclear despite decreased methyltransferase activity (at least for mutations that affect Arg882). There is no significant alteration in gene expression profile in *DNMT3A*-mutated cells compared to non-mutated cells.³ The pejorative prognostic impact of these mutations in AML and in MDS^{3,4} has not been demonstrated in MPNs.⁶



DNMT3A mutations in MPNs (Stegelmann et al. Leukemia 2011)

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