

show that among PRV-1-positive ET patients, those with normal c-Mpl levels appear to be at highest risk of thromboembolic events. The results of this retrospective analysis must now be corroborated in a large prospective trial of newly diagnosed patients such as that being implemented by the MPD Research Consortium

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Acknowledgments: the authors wish to thank Prof. Dr. K. Geiger for his continuing support. This work was supported by grants from the Else Kröner Fresenius-Stiftung and by the Alfried Krupp Förderpreis für junge Hochschullehrer, both awarded to H.L.P.

Key words: essential thrombocythemia, complications, c-Mpl, PRV-1.

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Acute Myeloid Leukemia

Mutations of *PTPN11* are rare in adult myeloid malignancies

The *PTPN11* gene encodes the phospho-tyrosine phosphatase protein SHP-2. Constitutional mutations of this gene are involved in Noonan's syndrome, a developmental disorder in which children have a predisposition to develop a myeloid disorder called juvenile myelomonocytic leukemia. Recently, studies have shown that somatic mutations of *PTPN11* can be found in children with myeloid malignancies. We evaluated the incidence of acquired mutation of *PTPN11* in 76 adults with acute or chronic myeloid malignancies and summarized our results together with others published recently.

haematologica 2005; 90:853-854

(<http://www.haematologica.org/journal/2005/6/853.html>)

The *PTPN11* gene (protein-tyrosine phosphatase, nonreceptor-type, 11), localized in 12q24, encodes a protein with tyrosine phosphatase activity called SHP-2 (Src homology 2 domain-containing phosphotyrosine phosphatase 2). The SHP-2 protein has 3 functional domains: the PTPase catalytic domain and two SH2 domains (src homology 2), one in the C-terminal part (C-SH2) and the other in the N-terminal part (N-SH2) of the protein. In its inactive form, the PTPase catalytic domain is masked by N-SH2; binding of phospho-tyrosyl peptides on N-SH2 induces a conformational change uncovering the catalytic site. SHP-2 is implicated in signal transduction pathways, particularly those induced by growth factors; activation of SHP-2 generally leads to activation of the RAS/RAF/ERK pathway.¹ The SHP-2 protein is strongly expressed in blood cells and is implicated in the response to KIT-ligand, interleukin-3, granulocyte-monocyte colony-stimulating factor and erythropoietin. The *PTPN11* alterations so far reported in human diseases were exclusively missense mutations. In Noonan's syndrome, *PTPN11* carries constitutional mutations preferentially localized in exon 3, the coding sequence for the domain of N-SH2 implicated in the inhibition of the PTPase catalytic site.² Children with this syndrome have a developmental disorder as well as a predisposition to develop juvenile myelomonocytic leukemia (JMML). Furthermore, in children, somatic mutations of *PTPN11* have been reported in 34% of non-syndromic JMML, 18.5% of refractory anemia with excess blasts (RAEB) in transformation, 4% of acute myeloid leukemia (AML), and 6.5% of acute lymphocytic leukemia (Table 1); 98% of those mutations were localized in exon 3.³⁻⁷ In this study, we performed mutational analysis of exon 3 of the *PTPN11* gene to assess the incidence of mutations in adult myeloid disorders. We tested 84 patients: 38 had chronic myelomonocytic leukemia (CMML), 18 RAEB, 4 RAEB in transformation, 4 AML with monosomy 7, 12 AML post-myelodysplastic syndrome and, as a control, 8 non-syndromic JMML. After DNA extraction from bone marrow or blood samples, exon 3 of *PTPN11* was amplified as previously described.³ Amplification products were purified and sequenced. As expected, 3 of the 8 cases of JMML tested carried mutations (G60R, D61V and G503A). Among the 76 remaining patients, only one had a *PTPN11* mutation (missense mutation: D61N). This patient had AML with monosomy

Table 1. Incidence of *PTPN11* mutations (mutated cases/studied cases).

Children						
JMML	NS-JMML	RAEB-T	Other MDS	AML	ALL	Ref.
1/4	-	-	-	-	-	Johan <i>et al.</i> ⁵
21/62	5/5	5/27	0/23	1/24	-	Tartaglia <i>et al.</i> ³
-	-	-	-	4/69	23/362	Tartaglia <i>et al.</i> ⁴
16/49	2/2	-	-	2/95	-	Loh <i>et al.</i> ⁶
-	-	-	-	11/278	-	Loh <i>et al.</i> ⁷
3/8	-	-	-	-	-	this study
33% (41/123)	100% (7/7)	18,5% (5/27)	0% (0/23)	4% (18/466)	6,5% (23/362)	Total
Adults						
RAEB-T	Other MDS	AML	CMML	CML		Ref.
0/2	0/70	1/64	0/35	-		Johan <i>et al.</i> ⁵
2/7	-	4/38	1/4	0/11		Loh <i>et al.</i> ⁶
0/15	0/26	0/49	0/1	-		Watkins <i>et al.</i> ⁸
0/4	0/18	1/16	0/38	-		this study
7% (2/28)	0% (0/114)	3,5% (6/167)	1% (1/87)	0% (0/11)		Total

JMML: juvenile myelomonocytic leukemia; NS-JMML: Noonan's syndrome - JMML; RAEB-T: refractory anemia with excess of blasts in transformation; MDS: myelodysplasia; AML: acute myeloid leukemia; ALL: acute lymphoid leukemia; CMML: chronic myelomonocytic leukemia; CML: chronic myeloid leukemia; -: no case.

7. As previously described by Loh *et al.* in pediatric AML7, monosomy 7 is frequently associated with *PTPN11* mutations. Our results confirm that alteration of *PTPN11* is a rare event in the leukemogenesis of adult myeloid malignancies. Indeed, in published studies^{5,6,8} (Table 1), only 1% of CMML, 3.5% of AML and 7% of RAEB-T had somatic mutations of *PTPN11*. Interestingly, *PTPN11* mutations occurred preferentially in leukemia with a monocytic component and/or in the presence of monosomy 7.^{4,6,7} Moreover, as in Noonan's syndrome, all somatic mutations described were missense mutations and they were almost exclusively localized in exon 3, rarely in exon 13. As demonstrated by Tartaglia *et al.*,² such mutations modify the zone of interaction between N-SH2 and PTPase domains and release enzymatic activity of SHP-2.

Contrary to pediatric myeloid malignancies in which different mechanisms can induce activation of the RAS-MEK-ERK pathway (RAS mutations, *NF1* deletions or *PTPN11* mutations), in adult myeloid malignancies this activation

only exceptionally involves *PTPN11* mutations and therefore seems related mostly to RAS mutations.

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Funding: this work was supported by the "Fondation de France" and by the "Ligue contre le cancer, Comité Nord".

Key words: *PTPN11*, SHP2, somatic mutation, adult, myeloid malignancies.

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Malignant Lymphomas

The relative levels of cyclin D1a and D1b alternative transcripts in mantle cell lymphoma may depend more on sample origin than on *CCND1* polymorphism

The relative levels of cyclin D1 (*CCND1*) (a) and (b) transcripts were determined by real-time reverse transcription polymerase chain reaction (RT-PCR) and found to vary according to the tissue origin in both control and tumor samples. A five-fold overexpression of both isoforms was observed in 28/38 cases of mantle cell lymphoma (MCL) and of only one isoform in 10/38 MCL. No correlation was observed between expression of cyclin D1 isoforms and *CCND1* genotype at position 870.

haematologica 2005; 90:854-856

(<http://www.haematologica.org/journal/2005/6/854.html>)