Brief report
The JAK2-V617F mutation is frequently present at diagnosis in patients with essential thrombocythemia and polycythemia vera

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We determined the allelic frequency of the JAK2-V617F mutation in DNA and assessed the expression levels of the mutant and wild-type JAK2 mRNA in granulocytes from 60 patients with essential thrombocythemia (ET) and 62 patients with polycythemia vera (PV) at the time of diagnosis. Using allele-specific quantitative polymerase chain reaction (qPCR), we detected JAK2-V617F in 75% of ET and 97% of PV at diagnosis. The total JAK2 mRNA levels were elevated in ET, PV, and secondary and idiopathic erythrocytosis, suggesting that hyperactive hemopoiesis alters JAK2 expression. The expression levels of JAK2-V617F mRNA were variable but strongly correlated with the allelic ratio of JAK2-V617F determined in DNA. Thus, differences in JAK2-V617F expression, markedly lower in ET than in PV, reflected different percentages of granulocytes carrying the mutation. Moreover, allelic ratios higher than 50% JAK2-V617F, indicating the presence of granulocytes homozygous for JAK2-V617F, were found in 70% of PV at diagnosis but never in ET.

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Study design
With informed consent and before treatment, blood and bone marrow (BM) samples from 62 PV and 60 ET, as well as 76 controls (24 secondary erythrocytosis [SE], 28 idiopathic erythrocytosis [IE], 24 reactive thrombofusosis), were collected in 5 centers. Blood was also obtained from 40 healthy donors (HDs) and 18 patients hospitalized for minor surgery. The study was approved by local Comité Consultatif de Protection des Personnes dans le Recherche Biomedicale de Bourgogne ethics committees.

Results and discussion
Patient characteristics and frequency of JAK2-V617F in ET and PV at diagnosis
Patients with erythrocytosis or thrombocytosis were diagnosed with PV, SE, ET, or RT according to Polycythemia Vera Study...
Table 1

<table>
<thead>
<tr>
<th>JAK2-V617F status</th>
<th>Negative (P)</th>
<th>Less than 25% (P)</th>
<th>25% to 40% (P)</th>
<th>41% to 74% (P)</th>
<th>75% or more (P)</th>
<th>All positive</th>
</tr>
</thead>
<tbody>
<tr>
<td>No. of patients</td>
<td>17</td>
<td>42</td>
<td>31</td>
<td>26</td>
<td>20</td>
<td>58</td>
</tr>
<tr>
<td>PV JAK2-V617F</td>
<td>408 / 421</td>
<td>377 / 373</td>
<td>306 / 326</td>
<td>60 / 33</td>
<td>616 / 619</td>
<td>629 / 634</td>
</tr>
<tr>
<td>ET JAK2-V617F</td>
<td>408 / 421</td>
<td>377 / 373</td>
<td>306 / 326</td>
<td>60 / 33</td>
<td>616 / 619</td>
<td>629 / 634</td>
</tr>
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**Expression levels of JAK2-WT and JAK2-V617F mRNA in granulocytes**

Relative expression of JAK2-WT and JAK2-V617F was quantitated against plasmidic standard dilutions and normalized for ABL expression. Granulocytes of IE and SE patients expressed significantly higher levels of JAK2-WT (medians: 348 and 452 JAK2-WT/100 ABL, respectively) than healthy donors and presurgery patients (197 and 174 copies/100 ABL, respectively). When both JAK2-WT and JAK2-V617F were considered, PV and ET granulocytes also expressed significantly more total JAK2 than healthy donors and presurgery patients (median total JAK2/100 ABL: 680 in PV, P < .001; 303 in ET, P = .008), suggesting that chronic stimulation of hematopoiesis may up-regulate JAK2 expression.

The percentage of total JAK2 represented by JAK2-V617F (%V617F) in ET and in PV was then analyzed (Figure 1; Table 1). This percentage can be assimilated to the allelic ratio since %V617F in genomic DNA was similar when assessed in cDNA and in genomic DNA from granulocytes of 40 patients (r = 0.90, P < .01; [%V617F in cDNA] = 1.03 × [%V617F in genomic DNA]). The percentages of mutant were not close to 50 or 100 as would be the case if all cells were exclusively heterozygous or homozygous for the mutant allele. Rather, they were distributed continuously, as if fractions of granulocytes had different allelic status. Thus, a percentage of mutants more than 50 is necessary and sufficient to affirm homozygosity. In positive PV, JAK2-V617F represented on average 62% of total JAK2 (median: 61%; range: 8%-98%). In all but one positive PV, JAK2-V617F represented more than 25% of total JAK2. Seventy percent of PV expressed more than 50% JAK2-V617F, implying that at least part of the granulocytes was homozygous for the V617F allele. In contrast, JAK2-V617F represented an average of 20% of total JAK2 in ET (P < .001...
compared with PV), and all but one positive ET expressed less than 40% JAK2-V617F (median: 19%; range: 3%-50%). Thus, the 50% JAK2-V617F threshold revealed more “homozygous” PV at diagnosis than previously reported using sequencing and nonquantitative PCR.1,3 In ET, either the mechanisms leading to JAK2-V617F homozygosity do not exist, or homozygous clones are repressed.

**Effects of JAK2-V617F levels on hematopoietic lineages and disease phenotype**

Percentages of JAK2-V617F correlated with granulocyte expression of PRV-1 (n = 92, r = 0.543, P < .001), confirming PRV-1 expression as a target of JAK2 signaling.1,9 In patients at diagnosis, JAK2-V617F levels correlated only with leukocyte counts (PV: n = 56, r = 0.496, P = .001; ET: n = 42, r = 0.314, P = .043), not with hemoglobin level, hematocrit level, platelet counts, numbers of EECs, or numbers of EMCs. Consistently, the level of mutant expression was not sufficient per se to determine PV or ET phenotype: comparison of PV and ET with similar levels of mutant (25%-40%) showed similar leukocyte counts, but PV patients had significantly higher EPO, EEC formation, red cell mass (152% vs 78%, P = .001), hematocrit, and hemoglobin level and lower platelet counts (Table 1). However, as recently described for ET,3 in our series of PV and ET at diagnosis, JAK2-V617F was associated with stimulation of erythropoiesis and repression of thrombopoiesis: PV with 75% or more mutant and positive ET differed from other PV and negative ET by higher hematocrit and hemoglobin levels, but lower platelet counts (Table 1).

In summary, sensitive qPCRs detected JAK2-V617F in 97% of PV and 75% of ET at diagnosis, with higher levels of expression in PV than in ET; cells homozygous for the mutation were present in 70% of PV. This demonstrates the interest of precise and sensitive assessment of JAK2-V617F for the diagnosis of MPD.

**Acknowledgments**

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**References**