

aggressive disease, and shows that despite the poor outcome, allogeneic SCT remains a potentially curative option for these patients.

Collectively, these data suggest that in patients with low or intermediate-1 IPSS risk and multilineage dysplasia, transfusion-dependency or moderate-to-severe bone marrow fibrosis, an early transplant should be considered when possible.

Much work does, however, remain to be done. The poor results observed in patients with severe bone marrow fibrosis in the study by Kroger *et al.* (18% 3-year disease-free survival) strongly support the need for further investigations aimed at improving this outcome. Such investigations must encompass an evaluation of the role of pre-transplant treatments, including hypomethylating agents and intensive chemotherapy, the most appropriate intensity of the preparative regimen and possible post-transplantation interventions aimed at preventing relapse.

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*Financial and other disclosures provided by the author using the ICMJE (www.icmje.org) Uniform Format for Disclosure of Competing Interests are available with the full text of this paper at www.haematologica.org.*

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## Pathophysiology of thrombosis in myeloproliferative neoplasms

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### **Thrombosis in myeloproliferative neoplasms: few answers and many new questions**

Life expectancy of patients with myeloproliferative neoplasms (MPNs) and particularly that of subjects with polycythemia vera (PV) and essential thrombocythemia (ET) has significantly increased over the last three decades, largely due to the use of cytoreductive treatments. Currently, polycythemia vera and essential thrombocythemia are consid-

ered relatively benign diseases in which the main objective of treatment strategy is the prevention of thrombotic events. Widespread use of routine hematologic screening and novel diagnostic tools greatly facilitate disease recognition and treatment. This helps to prevent a significant number of early vascular events which still constitute the first disease manifestation in approximately one-third of patients.<sup>1</sup> We can also expect that new therapeutic options

and appropriate use of aspirin will result in a further reduction of morbidity and mortality. The unmet needs of polycythemia vera and essential thrombocythemia subjects, however, remain significant. These include the availability of more safe and effective therapeutic agents and of validated tools for vascular risk stratification. The evaluation of the thrombotic risk in the individual patients, as underlined by Barbui *et al.* in their paper in this issue of the Journal,<sup>2</sup> is still approximate and generally relies on a limited number of variables. The role of all disease-related abnormalities and of their interaction with vascular risk factors remains largely obscure and difficult to explore in limited size epidemiological studies. In this complex scenario a significant scientific effort has recently been concentrated on studies testing new diagnostic and prognostic tools or aimed at clarifying the pathogenetic mechanisms of myeloproliferative neoplasm associated thrombophilia. This has several particular characteristics which include microcirculatory disturbances and thromboses at atypical sites often manifesting several years before diagnosis.<sup>3</sup>

**Pathophysiology of thrombosis in myeloproliferative neoplasms: what do we know?**

The pathogenesis of thrombosis in myeloproliferative neoplasms has been extensively investigated by focusing in particular on the possible contribution of disease related hemostatic abnormalities. However, the pathogenesis of thrombosis is multifactorial and even the relative role of these abnormalities, as compared with that of other individual and environmental factors, is controversial. Quantitative and qualitative red blood cell, platelet, and leukocyte abnormalities are likely to play a key-role in myeloproliferative neoplasm thrombophilia. High shear stress of the vessel wall, due to blood hyperviscosity, accounts for chronic endothelial dysfunction and platelet and leukocyte activation.

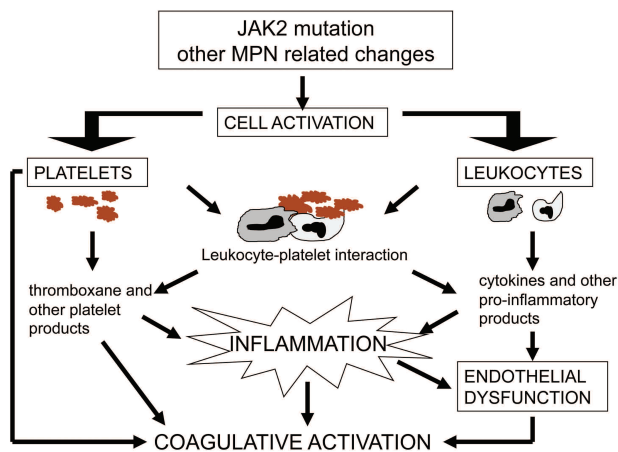
The increase of the thrombotic risk observed at progressively higher hematocrit values parallels blood viscosity, although also biochemical changes in the cell membrane and content could contribute to rheological abnormalities.<sup>4</sup> In this view, aggregates of red cells, platelets, leukocytes and endothelial cells have already been demonstrated to disturb blood flow and cause ischemia in the cerebral vessels.<sup>5</sup> Thrombocytosis can contribute to the vascular events in myeloproliferative neoplasms but qualitative platelet changes are likely to be even more important. Both polycythemia vera and essential thrombocythemia subjects were found to have an increased urinary excretion of the two major thromboxane A2 metabolites, 11-dehydro- and 2,3-dinor-Thromboxane B2, which can be suppressed by low-dose aspirin.<sup>6</sup> This finding constituted the biological background for assessing the efficacy of low-dose aspirin in the ECLAP trial.<sup>7</sup>

Platelets and endothelial cells play a pivotal role in regulating blood flow, both cells might contribute to determine a prothrombotic microenvironment in myeloproliferative neoplasm patients by producing more soluble selectins and less nitric oxide, likely as a consequence of inflammation (see below).<sup>8</sup>

Leukocytosis has been shown to represent an independent risk factor for thrombosis.<sup>9-10</sup> In a retrospective analysis of the ECLAP study, a leukocyte count greater than  $10 \times 10^9/L$ , adjusted for potential confounders such as cytoreductive and antithrombotic treatment, was associated with an increased rate of thrombosis.<sup>9</sup> A similar association was demonstrated in essential thrombocythemia patients.<sup>11</sup>

However, as for platelets, qualitative abnormalities might play a more critical role in the activation of the hemostatic system. Leukocyte activation in myeloproliferative neoplasm patients is demonstrated by overexpression of CD11b antigen and by the raised plasma levels of serine proteases cathepsin G, elastase and myeloperoxidase.<sup>12</sup> Expression of integrins and selectins increases the leukocyte adhesion to endothelium and platelets, stimulates the formation of mixed aggregates and the production of reactive oxygen species and inflammatory cytokines.<sup>13</sup>

These events link blood cell activation to inflammation and the latter is likely to play a significant role in the thrombotic risk of myeloproliferative neoplasms. Some mechanisms linking cell activation, inflammation and thrombosis are summarized in Figure 1.



**Figure 1.** Mechanisms which, in myeloproliferative neoplasms (MPN), can increase the thrombotic risk through cell activation and inflammation.

**Table 1.** Old and emerging biomarkers related to inflammation pathways with a potential role in major cardiovascular events.<sup>14-19</sup>

Parameter
Total and HDL-cholesterol levels
Oxidized LDL cholesterol
hs-C reactive protein
Pentraxin 3
Fibrinogen
TNF $\alpha$ , Interleukine 6 and 18
ICAM-1, VCAM, selectins, serine proteases
Soluble CD 40 ligand
Nitric oxide
Endothelin

CD: cluster denomination; hs: high-sensitive; ICAM: intercellular adhesion molecule; HDL: high-density-lipoprotein; LDL: low-density-lipoprotein; TNF: Tumor Necrosis Factor.

### Seeking novel vascular biomarkers in myeloproliferative neoplasms, the role of inflammatory parameters

The role of inflammation among the pathogenetic mechanisms of myeloproliferative neoplasm thrombophilia is also intriguing for the possible use of inflammatory parameters as vascular biomarkers. Barbui *et al.*<sup>2</sup> tested the hypothesis that blood levels of the inflammatory biomarkers C-reactive protein (CRP) and pentraxin-3 (PTX3) could be correlated with thrombosis and JAK2V617F allele burden in essential thrombocythemia and polycythemia vera subjects.

Within the limitations of the retrospective nature of the study and of the limited number of events, the results do suggest that C-reactive protein and pentraxin-3 measurements deserve scientific attention and might be helpful both for a more accurate risk stratification and for exploring the possible link between inflammation and thrombosis in myeloproliferative neoplasms.

In the general population this link is well established.<sup>14-15</sup> Over the last decade, a number of inflammatory biomarkers have been investigated as tools for identifying subjects at high risk for arterial vascular events (Table 1).<sup>14-16</sup>

More recently, high C-reactive protein levels have been shown to also influence the risk of venous thromboembolism and this has led to the hypothesis of a direct causative role of C-reactive protein on venous thrombophilia.<sup>17</sup> A recent study, however, ruled out this role.<sup>18</sup> Thus, one has to conclude that an increased inflammatory state can play a causative role in both arterial and venous thromboses.

In the study of Barbui *et al.*,<sup>2</sup> C-reactive protein plasma levels were interestingly related with JAK2 mutation allelic burden thus suggesting a role of this mutation on inflammation and cytokine production, as reflected by C-reactive protein plasma levels. Raised levels of this protein were associated with an increased thrombotic risk but, unfortunately, the size of the study did not allow a comparison to be made on the predictive value of C-reactive protein on venous and arterial thromboses. Also pentraxin-3, like C-reactive protein, was significantly correlated with JAK2 allele burden. But in this case, pentraxin-3 levels higher than 4.5 ng/mL were associated with a lower thrombotic risk.

#### How can we interpret these data?

For a convincing interpretation of Barbui's data further investigations both in larger myeloproliferative neoplasm cohorts and in the general population are needed. In fact, compared to C-reactive protein, pentraxin 3 has undergone less extensive investigation and its role is still uncertain.<sup>19</sup> Moreover, while C-reactive protein is produced by the liver, pentraxin 3 is produced at the site of inflammation by various cell types, including endothelial cells, neutrophils, monocytes and macrophages, and its level increases in several pathological conditions (sepsis, autoimmune disorders, small vessel vasculitis).<sup>19-20</sup>

Pentraxin 3 is induced by IL-1, TNF- $\alpha$ , oxidized low-density lipoprotein and lipopolysaccharide and its plasma levels are not correlated with C-reactive protein levels.

Recent studies on humans and mice reported that high levels of pentraxin 3 in cardiovascular diseases seem to reflect a protective response to severe ischemic injury.<sup>19-21</sup> Moreover, pentraxin 3 deficiency in mice is associated with

increased atherosclerosis and increased macrophage accumulation in atherosclerotic lesions. In addition, myocardial infarcts are significantly larger in PTX3-deficient than in wild-type mice and the use of recombinant pentraxin 3 can reduce heart damage and inflammation and block neointimal thickening after balloon injury of rat carotid arteries.<sup>19</sup>

It has been hypothesized that an increased pentraxin 3 production is a feedback mechanism which inhibits leukocyte recruitment, modulates complement activation and balances the overactivation of a pro-inflammatory and pro-atherogenic cascade.<sup>22</sup> This hypothesis suggests that the increased levels of pentraxin 3 in subjects with cardiovascular diseases could reflect both inflammation and a protective response. On the basis of the model proposed above one can hypothesize that when the pentraxin 3 response to inflammation is blunted by genetic or acquired factors the thrombophilic state linked to inflammation is further enhanced. The hypothesis is too simplistic to apply to all clinical settings but Barbui's data suggest that it could fit myeloproliferative neoplasm patients.

#### Perspectives

The study by Barbui *et al.* raises several questions which need to be addressed by future studies. These questions include the type of thromboses which are more prevalent in subjects having high C-reactive protein, low pentraxin 3 plasma levels or both conditions. This issue seems relevant since inflammation in the general population seems particularly linked with the risk of acute coronary syndromes and also the increased risk associated with high leukocyte count in polycythemia vera was found to be particularly linked with coronary events.<sup>9</sup> Another interesting issue is the association of inflammatory biomarkers with JAK2 allelic burden. This association is biologically plausible<sup>12-13</sup> but we do need more extensive clinical data to clarify this field also in view of the rather controversial association between the risk of thrombosis and allele burden. According to the data of Barbui *et al.*,<sup>2</sup> it is intriguing to consider the possibility that pentraxin 3 response to inflammation in subjects with high JAK2 burden might contribute to lower or enhance the thrombotic risk. More generally the association between JAK2 mutation, inflammation and thrombotic risk deserves scientific attention also for other speculative and practical purposes. For example, there is still no convincing explanation for the well-known association between venous thromboses at atypical sites and JAK2 mutation in patients who do not meet criteria for myeloproliferative neoplasm diagnosis and this also raises the question of interventional strategies. To date, this relationship suggests a direct influence of the JAK2 mutation on the hemostatic system and raises further expectations from molecules that selectively target JAK2 kinase.

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*Financial and other disclosures provided by the author using the ICMJE (www.icmje.org) Uniform Format for Disclosure of Competing Interests are available with the full text of this paper at www.haematologica.org.*

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## Macrophages predict treatment outcome in Hodgkin's lymphoma

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Although Hodgkin's lymphoma (HL) can be considered a successful paradigm of modern treatment strategies, about 15-20% of patients with advanced-stage HL still die following relapse or progressive disease and a similar proportion of patients are over-treated,<sup>1,2</sup> leading to treatment-related late sequelae including solid tumors and end-organ dysfunction.<sup>3,4</sup> To help guide treatment decisions, the distinction between classical HL and nodular lymphocyte-predominant HL and the separation of limited and advanced-stage disease are widely used in clinical practice. Most patients present with advanced-stage classical HL and for this group of patients the International Prognostic Score was introduced more than a decade ago to improve risk stratification.<sup>5</sup> However, neither the International Prognostic Score itself nor any of the individual clinical variables can predict the majority of patients in whom standard therapy will fail to eradicate the disease. Novel biological markers that

improve on primary treatment outcome prediction across all clinical stages will, therefore, be critical to advancing the field.

The histological hallmark of HL is the presence of the malignant mononuclear Hodgkin and polynucleated Reed Sternberg (HRS) cells in classical HL and so-called lymphocyte predominant cells in nodular lymphocyte-predominant HL.<sup>6</sup> These malignant cells are, however, greatly outnumbered by the reactive cells in the tumor microenvironment.<sup>7</sup> Because of the prominent and abnormal immune reaction that creates this variable microenvironment, the biology of HL can be considered unique among lymphomas. Many studies in HL have, therefore, focused on the cellular composition of the microenvironment, not only to gain more insight into the pathobiology of the disease, but also to explore whether these immune-related cells in some way contribute to outcome prediction.

In this issue of the journal, Kamper *et al.* used immuno-