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JAK2^{V617F}-mutant Vascular Niche Contributes to JAK2^{V617F} Clonal Expansion in Myeloproliferative Neoplasms

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Abstract

The myeloproliferative neoplasms (MPNs) are characterized by hematopoietic stem/progenitor cell (HSPC) expansion and overproduction of blood cells. The acquired mutation *JAK2*^{V617F} plays a central role in these disorders. Mechanisms responsible for MPN HSPC expansion is not fully understood, limiting the effectiveness of current treatments. Endothelial cells (ECs) carrying the *JAK2*^{V617F} mutation can be detected in patients with MPNs, suggesting that ECs are involved in the pathogenesis of MPNs. Here we report that *JAK2*^{V617F}-bearing primary murine ECs have increased cell proliferation and angiogenesis *in vitro* compared to *JAK2*^{WT} ECs. While there was no difference between *JAK2*^{V617F} and *JAK2*^{WT} HSPC proliferation when co-cultured with *JAK2*^{WT} EC, the *JAK2*^{V617F} HSPC displayed a relative growth advantage over the *JAK2*^{WT} HSPC when co-cultured on *JAK2*^{V617F} EC. In addition, the thrombopoietin (TPO) receptor MPL is up regulated in *JAK2*^{V617F} ECs and contributes to the maintenance/expansion of the *JAK2*^{V617F} clone over *JAK2*^{WT} clone *in vitro*. Considering that ECs are an essential component of the hematopoietic niche and most HSPCs reside in the perivascular niche, our studies suggest that the *JAK2*^{V617F}-bearing ECs form an important component of the MPN vascular niche and contribute to mutant stem/progenitor cell expansion, likely through a critical role of the TPO/MPL signaling axis.

Keywords

myeloproliferative neoplasms; hematopoietic stem/progenitor cells; microenvironment; endothelial cells; *JAK2*^{V617F}; MPL

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Author Contribution

C.L. performed experiments, analyzed data, and reviewed the manuscript; K.K. analyzed data and wrote the manuscript; and H.Z. designed and performed experiments, analyzed data, and wrote the manuscript. All authors approved the submitted version of the manuscript.

Introduction

The marrow consists of the hematopoietic cells and non-hematopoietic stromal cells, including fibroblasts, reticular cells, endothelial cells (ECs), macrophages, adipocytes and osteoblasts. In addition to its role in normal HSPC biology, an altered microenvironment is an important contributor to the development of hematologic malignancies.¹⁻³ In a reciprocal fashion, myeloid malignancies also affect the function of the marrow microenvironment to impair normal hematopoiesis while favoring malignant stem cell expansion.^{4,5}

The chronic Philadelphia chromosome (Ph¹) negative myeloproliferative neoplasms (MPNs), including polycythemia vera (PV), essential thrombocythemia (ET) and primary myelofibrosis (PMF), are clonal stem cell disorders characterized by HSPC expansion, overproduction of mature blood cells, a tendency to extramedullary hematopoiesis, and transformation to acute leukemia or myelofibrosis at variable rates. The acquired signaling kinase mutation *JAK2*^{V617F} plays a central role in the pathogenesis of MPN, but mechanism(s) responsible for MPN HSPC expansion is not fully understood, limiting the effectiveness of current treatments. Although the etiology of dysregulated hematopoiesis has been mainly attributed to the molecular alterations within the hematopoietic stem/progenitor cells, abnormalities of the marrow microenvironment are beginning to be recognized as an important factor in MPN development.^{1,5-7} The diseased niche could impair normal hematopoiesis and favor the competing malignant stem cells, which could contribute to the poor engraftment and treatment-related mortality following allogeneic stem cell transplantation, the only curative treatment for patients with MPNs.^{1-3,5,8,9}

ECs are an essential component of the hematopoietic niche and most hematopoietic stem/progenitor cells (HSPCs) reside close to a marrow sinusoid (the “perivascular niche”).¹⁰⁻¹³ In addition, ECs are an important niche component of the extramedullary (splenic) hematopoiesis, which is almost always present in patients with MPNs and is associated with MPN disease progression.^{13,14} Although the existence and cell of origin of endothelial progenitors is still a matter of debate, *JAK2*^{V617F} mutation can be detected in endothelial progenitors derived from the hematopoietic lineage (the so-called endothelial cell colony-forming units; CFU-ECs) and, in some reports, in the true endothelial colony-forming cells (ECFC) based on *in vitro* assays.¹⁵⁻¹⁹ *JAK2*^{V617F} mutation is also present in ECs isolated by microdissection from liver and spleen samples of patients with MPNs.^{17,20} In addition, we and others have shown that *JAK2*^{V617F} ECs are critical in the development of the bleeding abnormalities in a murine model of *JAK2*^{V617F}-positive MPNs in which *JAK2*^{V617F} is expressed in all hematopoietic cells and endothelial cells.²¹ All of these observations suggest that ECs are involved in the pathogenesis of MPNs.

Previously, we and others have shown that the thrombopoietin (TPO) receptor MPL is essential for the development of HSPC expansion in MPNs.^{22,23} MPL is expressed in long-term HSPCs and is associated with both HSPC repopulating activity and HSPC quiescence.^{24,25} MPL is also expressed on several types of endothelium.²⁶⁻²⁸ Whether EC MPL receptor affects the vascular niche function and contributes to MPN development is not known. In this study, we examined the roles of *JAK2*^{V617F}-bearing ECs in MPN

hematopoiesis. We found that the MPN vascular niche contributes to the growth advantage of JAK2^{V617F} HSPC over the JAK2^{WT} HSPC. We also found that the EC MPL receptor is important for the JAK2^{V617F} clonal expansion in MPNs.

Materials and Methods

Experimental mice

JAK2^{V617F} Flip-Flop (FF1) mice²⁹ were kindly provided by Radek Skoda (University Hospital, Basel, Switzerland), Tie2-Cre mice³⁰ by Mark Ginsberg (University of California, San Diego), and MPL knockout mice (MPL^{-/-})³¹ by Warren Alexander (Melbourne, Australia). FF1 mice were crossed with Tie2-Cre to express JAK2^{V617F} specifically in hematopoietic cells and ECs (Tie2/FF1 mice) as we previously did.²¹ All mice used were on a C57BL/6 background and were bred in a pathogen-free mouse facility at Stony Brook University. CD45.1+ congenic mice (SJL) were purchased from Taconic Inc. (Albany, NY). Animal experiments were performed in accordance with the guidelines provided by the Institutional Animal Care and Use Committee at Stony Brook University.

Isolation of murine hematopoietic stem/progenitor cells (HSPCs)

14–18-week old mice were euthanized and the femurs and tibias removed. A 25-gauge needle was used to flush the marrow with PBS + 2% FBS. Cells were triturated and filtered through 70µm nylon mesh (BD Biosciences, San Jose, CA) to obtain a single cell suspension. For depletion of mature hematopoietic cells, the Lineage Cell Depletion Kit (Miltenyi Biotec, Cat. 130-090-858, San Diego, CA) was used. The lineage (CD5, CD45R, CD11b, Ter119, and GR-1) negative cells were collected and then positively selected for CD117⁺ (cKit⁺) cells using CD117 microbead (Miltenyi Biotec, Cat. 130-091-224) to yield Lineage^{neg}cKit⁺ (Lin⁻cKit⁺) HSPCs.

Isolation of murine lung endothelial cells

Primary murine lung EC isolation was performed as we previously did.³² Briefly, 14–18wk old mice were euthanized using 100% CO₂ inhalation followed by cervical dislocation. The chest was immediately opened through a midline sternotomy. The left ventricle was identified and the ventricular cavity was entered through the apex with a 27-gauge needle. The right ventricle was identified and an incision was made in the free wall to exsanguinate the animal and to allow the excess perfusate to exit the vascular space. The animal was perfused with 30 ml of cold PBS. The lung tissue was collected and minced finely with scissors. The tissue fragments were digested in DMEM medium containing 1 mg/mL Collagenase D (Roche, Switzerland), 1 mg/mL Collagenase/Dispase (Roche) and 25 U/mL DNase (Sigma, St. Louis, MO) at 37°C for 2hr with shaking, after which the suspension was homogenized by triturating. The homogenate was filtered through a 70µm nylon mesh (BD Biosciences, San Jose, CA) and pelleted by centrifugation (400g for 5 min). Cells were first depleted for CD45⁺ cells (Miltenyi Biotec) and then positively selected for CD31⁺ cells (Miltenyi Biotec) using magnetically labeled microbeads according to the manufacturer's protocol. Isolated ECs (CD45⁻CD31⁺) were cultured in EC culture medium with no medium change for the first 72hrs to allow EC attachment followed by medium change every 2–3

days. Cells were re-selected for CD31⁺ cells when they reach >70–80% confluence (usually after 3–4 days of culture).

Flow cytometry

CD45.1 (A20) antibody or isotype control was used for chimerism studies. All staining steps were performed in ice-cold PBS containing 2% fetal bovine serum. All samples were analyzed by flow cytometry using a FACSCalibur flow cytometer (Becton Dickinson, Franklin Lakes, NJ). Postacquisition data analysis was performed with FlowJo software V9.2.3 (Treestar, CA).

Polymerase Chain Reaction

Human JAK2 cDNA-specific primers (5'-GAAGAACTTCAGCAGTCTTAAAGATC-3' and 5'-CCATGCCAACTGTTTAGCAACTTC-3') were used to detect the expression of human JAK2V617F in ECs from Tie2/FF1 mice using reverse transcription polymerase chain reaction (RT-PCR). The primers amplify a 573bp fragment that would be detected on 2% agarose gel.

The TaqMan® Gene Expression Assay (Applied Biosystems) was used for real-time quantitative polymerase chain reaction (qPCR) to verify differential expression of MPL (Mm00440306_g1) on an ABI ViiA™ 7 Real-Time PCR machine (Applied Biosystems). The gene expression levels were normalized to Actin beta (Actb) expression and relative fold changes was calculated by the 2^{-CT} method. All assays were performed in triplicate.

Assays to examine endothelial cell in vitro angiogenesis

EC tube formation assay was performed as a measure of angiogenesis *in vitro*.³² Matrigel® matrix (10mg/ml, Corning Inc., Corning, NY) were thaw overnight at 4°C and kept on ice until use. 150ul Matrigel per well was added to pre-chilled 48-well culture plate. After gelation at 37°C for 30minutes, gels were overlaid with 6 × 10⁴ JAK2^{V617F} ECs (from Tie2/FF1 mice) or JAK2^{WT} ECs (from control mice) (passage 3–4) in 300ul of complete EC medium. Tube formation was inspected after a period of 2, 4, 6, and 8hrs and images were captured with a phase-contrast microscope (AMEX-1200, AMG, Bothell, WA). The quantification of the capillary tube formation was performed using the ImageJ® software (National Institute of Health, Bethesda, MD) by counting the number of nodes (or branch points), loops, and tubes in 4 non-overlapping areas at ×40 magnification in two duplicate wells.

In vitro cell culture

Lin⁻cKit⁺ HSPCs were cultured in StemSpan® serum-free expansion medium (SFEM) containing 100 ng/mL recombinant mouse SCF, 6 ng/mL recombinant mouse IL3 and 10 ng/mL recombinant human IL-6 (all from Stem Cell Technologies, Vancouver, BC)

ECs were cultured on 1% gelatin coated plates in complete EC medium which is consisted of advanced DMEM/F12 (ThermoFisher, Waltham, MA) medium containing 20% fetal bovine serum, 50µg/ml endothelial cell growth supplement (Alfa Aesar, Ward Hill, MA), 1% Antibiotic-antimycotic solution (Cat. 15240-062, ThermoFisher), 10mM HEPES buffer

(ThermoFisher), 5 μ M SB431542 small molecule (R&D, Minneapolis, MN), 50 μ g/ml Heparin (Sigma), 1% Glutamax 100 \times solution (ThermoFisher), 1% non-essential amino acid (ThermoFisher), recombinant mouse VEGF 10ng/ml (PeproTech, Rocky Hill, NJ; add fresh when changing medium) and recombinant human FGF2 20ng/ml (PeproTech; add fresh when changing medium). Cells were cultured at 37°C in a humidified 5% CO₂ atmosphere with medium change every 2–3 days until they reach 70–80% confluence. 0.05 % trypsin-EDTA solution was used for cell passaging. EC-conditioned media (ECCM) was collected when cell density reach 50–60% confluency.

Co-culture of Lin-cKit⁺ cells with EC

Three days prior to co-culture with HSPCs, 2 \times 10⁴ primary murine lung ECs (passage 1–3) were seeded into 1% gelatin coated 24-well plate in complete EC medium. On Day 0, 10⁴ Lin-cKit⁺ cells were seeded onto the EC monolayer in 0.3ml SFEM containing 100 ng/mL recombinant mouse SCF, 6 ng/mL recombinant mouse IL3 and 10 ng/mL recombinant human IL-6. 0.3ml fresh SFEM with cytokines was added on Day 3 and cells were counted on Day 5.

In vitro competitive growth assay

On Day 0, 10³ WT Lin-cKit⁺ HSPCs (CD45.1) and 10³ Tie2/FF1 Lin-cKit⁺ HSPCs (CD45.2) were mixed and culture together in 24-well plate with 0.2ml SFEM, or SFEM + WT ECCM (1:1 vol:vol), or SFEM + MPL^{-/-} ECCM (1:1 vol:vol) containing 100 ng/mL recombinant mouse SCF, 6 ng/mL recombinant mouse IL3 and 10 ng/mL recombinant human IL-6. 0.2ml fresh medium with cytokines was added on Day 3. After 6 days of culture, cells were stained with CD45.1 antibody for flow cytometry analysis.

Statistical Analysis

Statistical analyses were performed using Student's unpaired, 2-tailed *t* tests using Excel software (Microsoft). A *p* value of less than 0.05 was considered significant. For all bar graphs, data are presented as mean \pm standard error of the mean (SEM).

Results

JAK2^{V617F} mutant ECs show increased cell proliferation and angiogenesis *in vitro* compared to JAK2^{WT} ECs

FF1 mice were crossed with Tie2-Cre mice to generate mice that express JAK2^{V617F} specifically in hematopoietic cells and ECs (Tie2/FF1 mice) as was previously described.²¹ Primary lung ECs were isolated from Tie2/FF1 mice (JAK2^{V617F} ECs) and age-matched littermate control mice (JAK2^{WT} EC). The expression of human JAK2^{V617F} mutation in Tie2/FF1 ECs was confirmed by RT-PCR. (Figure 1A) Consistent with a previous report on lentiviral-transduced JAK2^{V617F}-mutant human umbilical vein endothelial cells (HUVECs),³³ primary murine lung ECs carrying the JAK2^{V617F} mutation proliferated to a greater extent than JAK2^{WT} ECs (2.0-fold, *p* = 0.031). (Figure 1B) To study the effects of JAK2^{V617F} mutation on EC function, tube formation assay was performed on JAK2^{WT} and JAK2^{V617F} ECs as a measure of their capacity for *in vitro* angiogenesis. JAK2^{V617F} ECs had significantly increased angiogenesis in MatrigelTM compared to JAK2^{WT} EC. (Figure 1C) In

addition, the tubular structures formed by the JAK2^{V617F} ECs *in vitro* were more stable than those from JAK2^{WT} ECs. (Figure 1D) This result suggests that the JAK2^{V617F} mutation in EC might be responsible for the increased microvascular density in the marrow of patients with MPNs.^{34–36} Since ECs are an essential component of the hematopoietic niche, and most HSPCs reside in the perivascular niche,^{10–13} these results prompted us to study how JAK2^{V617F}-bearing ECs affect hematopoiesis in MPNs.

The JAK2^{V617F} vascular niche contributes to the maintenance/expansion of JAK2^{V617F} HSPC in preference to JAK2^{WT} HSPCs

Lin-cKit+ HSPCs were isolated from Tie2/FF1 and age-matched littermate control mice. To examine the role of JAK2^{WT} and JAK2^{V617F} ECs in MPN hematopoiesis, we used an *ex vivo* co-culture system in which Lin-cKit+ HSPCs were cultured on a feeder layer of JAK2^{WT} or JAK2^{V617F} ECs under serum-free conditions. We compared JAK2^{WT} and JAK2^{V617F} HSPC proliferation on JAK2^{WT} or JAK2^{V617F} ECs respectively. Consistent with previous reports that JAK2^{V617F} does not confer a significant growth advantage to HSPCs,^{37–42} we did not observe any significant difference between JAK2^{WT} and JAK2^{V617F} HSPC proliferation *in vitro* in SFEM. (Figure 2A) While there was no difference between JAK2^{V617F} and JAK2^{WT} HSPC proliferation when co-cultured with JAK2^{WT} EC, the JAK2^{V617F} HSPC displayed a relative growth advantage over the JAK2^{WT} HSPC (1.24-fold, $p = 0.041$) when co-cultured on JAK2^{V617F} EC. (Figure 2B–C) These results suggest that the MPN vascular niche (i.e. JAK2^{V617F} EC) contributes to the maintenance/expansion of the JAK2^{V617F} HSPCs in preference to JAK2^{WT} HSPCs. This is consistent with other reports that the marrow microenvironment of myeloid malignancies are altered to impair normal hematopoiesis while favoring malignant stem cell expansion.^{4,5}

The EC MPL receptor contributes to the maintenance/expansion of the JAK2^{V617F} HSPC over JAK2^{WT} HSPCs

Thrombopoietin (TPO) and its receptor, the proto-oncogene MPL, are key regulators of HSPC activity.^{24,25,43–46} MPL is expressed in long-term HSPCs and is associated with both HSPC repopulating activity and HSPC quiescence.^{24,25} Previously, we and others have shown that MPL is essential for the development of an increased neoplastic stem cell pool in MPNs.^{22,23} Specifically, reducing MPL expression attenuated MPN severity and stem cell numbers, suggesting a gene dosage effect of receptor expression levels on the disease process. MPL is also expressed on endothelial cells (ECs) and TPO can stimulate EC growth and angiogenesis.^{26–28} Both TPO knockout (TPO^{-/-}) and MPL knockout (MPL^{-/-}) mice have significantly decreased HSPC populations and impaired angiogenesis.^{46–48} We checked MPL expression levels in JAK2^{WT} and JAK2^{V617F} ECs by qPCR. We found that MPL expression was increased by 46% in the JAK2^{V617F} ECs compared to JAK2^{WT} ECs ($p = 0.003$). (Figure 3A) Therefore, we hypothesized that MPN JAK2^{V617F}-mutant clone expansion might depend on the endothelial microenvironment, as mediated by the endothelial MPL receptor.

To test this hypothesis, we obtained primary murine lung ECs isolated from the MPL knockout mice (MPL^{-/-} EC)³¹ and performed *ex vivo* co-culture with JAK2^{V617F} Lin-cKit+ HSPCs. We found that JAK2^{V617F} HSPC (Lin-cKit+) cell proliferation was significantly

decreased when co-cultured with MPL^{-/-} EC, compared to when co-cultured with JAK2^{WT} ECs or JAK2^{V617F} ECs. (Figure 3B) This suggests that the EC MPL receptor is important for the JAK2^{V617F} HSPC expansion in the MPN vascular niche.

Next, since both JAK2^{WT} clones and JAK2^{V617F} mutant clones coexist in most patients with MPNs, we asked whether the EC MPL receptor contributes to the JAK2^{V617F} mutant clonal expansion over JAK2^{WT} clones. We designed a *in vitro* competitive growth experiment where both JAK2^{WT} HSPCs (CD45.1) and JAK2^{V617F} HSPCs (CD45.2) were cultured together (1:1 mix) in the presence of EC conditioned medium (ECCM) collected from either WT EC or MPL^{-/-} EC. At the end of the 6-day culture, there were equal numbers of JAK2^{WT} HSPCs (CD45.1 48.9±2.6%) and JAK2^{V617F} HSPCs in the presence of WT ECCM, which was what we expected since there was no significant difference between JAK2^{V617F} and JAK2^{WT} HSPC proliferation when co-cultured with JAK2^{WT} EC (Figure 2B). However, there were significantly more JAK2^{WT} HSPCs (CD45.1 77.3±2.6%) than JAK2^{V617F} HSPCs in the presence of MPL^{-/-} ECCM compared to in WT ECCM ($p = 0.002$), suggesting that EC MPL receptor contributes to the maintenance/expansion of the JAK2^{V617F} clone over the JAK2^{WT} clone in MPNs. (Figure 3C)

Discussion

ECs carrying the JAK2^{V617F} mutation can be detected in patients with MPNs.^{16,17,20} In addition, ECs are an important niche component of the extramedullary (splenic) hematopoiesis, which is almost always present in patients with MPNs and is associated with MPN disease progression.^{13,14} All of these observations suggest that ECs are involved in the malignant process leading to MPNs. Previously, we and others have shown that JAK2^{V617F} ECs are critical in the development of the bleeding abnormalities in a murine model of JAK2^{V617F}-positive MPNs in which JAK2^{V617F} was expressed in all hematopoietic cells and endothelial cells (Tie2-Cre/FF1).^{21,29,30} In this study, using primary ECs isolated from wild-type control (JAK2^{WT} EC), Tie2/FF1 (JAK2^{V617F} EC), and MPL^{-/-} mice (MPL^{-/-} EC), we have found that JAK2^{V617F}-bearing ECs contribute to the competitive advantage of JAK2^{V617F} HSPC over the JAK2^{WT} HSPC, and the MPL receptor on vascular ECs is important for JAK2^{V617F} HSPC maintenance/expansion *in vitro*.

Consistent with the enhanced angiogenesis and increased microvascular density in the marrow of patients with MPNs compared to normal marrow,³⁴⁻³⁶ JAK2^{V617F}-bearing ECs proliferated to a greater extent than JAK2^{WT} EC (2.0-fold, $p = 0.031$) (Figure 1B) and displayed increased angiogenesis *in vitro*, as measured by the tube formation assay. (Figure 1C) In addition, we found that JAK2^{V617F} HSPC gained a competitive advantage compared to JAK2^{WT} HSPC when co-cultured on JAK2^{V617F} EC, while there was no difference between JAK2^{V617F} and JAK2^{WT} HSPC proliferation when co-cultured on JAK2^{WT} EC. Both JAK2^{WT} and JAK2^{V617F}-mutant HSPC cell proliferation was decreased when co-cultured on ECs compared to cultured alone in SFEM, suggesting that our EC co-culture system inhibits HSPC proliferation or promotes HSPC quiescence by the elaboration of secreted factors and/or by cell-cell contacts. We hypothesize that the JAK2^{V617F} mutation alters both HSPC and EC function to provide the relative growth advantage of the JAK2^{V617F}-bearing HSPCs over JAK2^{WT} HSPCs when co-cultured on JAK2^{V617F} EC. As

no significant difference was detected between JAK2^{V617F} and JAK2^{WT} HSPC proliferation when co-cultured with JAK2^{WT} EC, our study suggests that both mutant HSPCs and mutant vascular niche are required for the development of MPNs. The stem cell compartment in MPN is heterogeneous with the presence of both JAK2^{WT} clones and JAK2^{V617F} mutant clones in most patients with MPNs. The mechanism(s) that contribute to the JAK2^{V617F}-positive HSPC proliferation and expansion is poorly understood, limiting the effectiveness of current treatments. Since ECs are an essential component of the hematopoietic niche, and most HSPCs reside in the perivascular niche,^{10–13} our results suggest that the JAK2^{V617F}-bearing vascular niche contributes to the JAK2^{V617F}-mutant clonal expansion in MPNs. Therefore, targeting the MPN vascular niche could provide more effective adjuvant therapies to treatments directly focusing on the MPN neoplastic cells.

The TPO receptor, MPL, is a key regulator of HSPC activity.^{24,25,44–46,49} MPL is expressed in long-term HSPCs and is associated with both HSPC repopulating activity and HSPC quiescence.^{24,25} MPL is also expressed on several types of endothelium.^{26–28} Our previous study showed that the EC MPL receptor does not contribute significantly to the regulation of TPO levels or to steady-state platelet counts.⁵⁰ Whether the EC MPL receptor could affect vascular niche function, and contribute to the critical role of TPO/MPL signaling in HSPC maintenance, is not known. In this study, we found that MPL expression was increased in the JAK2^{V617F} ECs compared to JAK2^{WT} ECs. (Figure 3A) Ablating the MPL receptor in ECs (i.e. MPL^{-/-} ECs) significantly decreased cell proliferation of the co-cultured JAK2^{V617F} HSPCs compared to JAK2^{WT} ECs or JAK2^{V617F} ECs did. (Figure 3B) In addition, an *in vitro* competitive growth experiment showed that the EC MPL receptor contributes to the maintenance/expansion of the JAK2^{V617F} clone over the JAK2^{WT} clone when cultured together. (Figure 3C) Together with our previous report,²² we have shown that the MPL receptor on HSPCs and ECs is important for JAK2^{V617F} HSPC expansion in MPNs. Since MPL relies on JAK2 protein for its downstream signaling pathways, and JAK2 can affect MPL receptor stability and cell surface expression, we hypothesize that the JAK2^{V617F} mutation could alter the vascular niche function and HSPC-EC interactions through altered TPO/MPL signaling, which in turn could contribute to altered hematopoietic factor and adhesion molecule production/expression in the MPN vascular niche. Considering that most HSPCs reside adjacent to a marrow sinusoid (the “perivascular niche”),^{10–13} our results suggest that JAK2^{V617F}-bearing endothelial niche could maintain/expand the JAK2^{V617F} HSPC clone via altered TPO/MPL signaling.

In this study, primary lung ECs were used to model the hematopoietic vascular niche *in vitro*. Since lung ECs might have different structural, phenotypic, or functional attributes from the marrow or spleen ECs,⁵¹ we have also used primary murine splenic ECs from wild-type control (JAK2^{WT} EC), Tie2/FF1 (JAK2^{V617F} EC), and MPL^{-/-} mice (MPL^{-/-} EC) and verified selected findings in splenic ECs (data not shown). In contrary to previous reports that ECs (e.g. HUVECs, human brain ECs, transfected/immortalized HUVECs, or primary murine ECs) support hematopoietic stem/progenitor cell (HSPC) expansion *in vitro*, our study found decreased Lin-cKit⁺ HSPC cell expansion when cultured on lung ECs.^{52–57} The different results could be due to differing HSPC populations, differences in the type of ECs employed, as well as culture medium and growth factors used in these studies. Further investigation will be required to explore whether the MPN vascular niche (with the

JAK2^{V617F}-bearing ECs) contributes to the JAK2^{V617F} HSPC clonal expansion *in vivo* and whether this process also depends on the EC MPL receptor. Based on results reported here (i.e. JAK2^{WT} HSPC did not display any growth advantage on JAK2^{V617F} EC compared to on JAK2^{WT} EC), competitive marrow transplantation experiments are likely required to model the human diseases and quantify JAK2^{WT} and JAK2^{V617F} HSPC numbers and functions in the JAK2^{V617F}-bearing vascular niche.

In summary, our studies have shown that JAK2^{V617F}-bearing ECs form an important part of the MPN HSPC niche and contribute to JAK2^{V617F} clonal expansion *in vitro*. Further, we have demonstrated that the EC MPL receptor is important for the maintenance/expansion of the JAK2^{V617F} HSPC over JAK2^{WT} HSPCs, providing a potential mechanism through which the absence of MPL prevents MPN development.^{22,23} Considering that both JAK2^{WT} clones and JAK2^{V617F} mutant clones coexist in most patients with MPNs, our findings provide a possible new mechanism for the mutant clone expansion seen over time in patients with MPNs, and suggests that the MPN vascular niche and the EC MPL receptor may be excellent therapeutic targets to eradicate the malignant clone in patients with MPNs.

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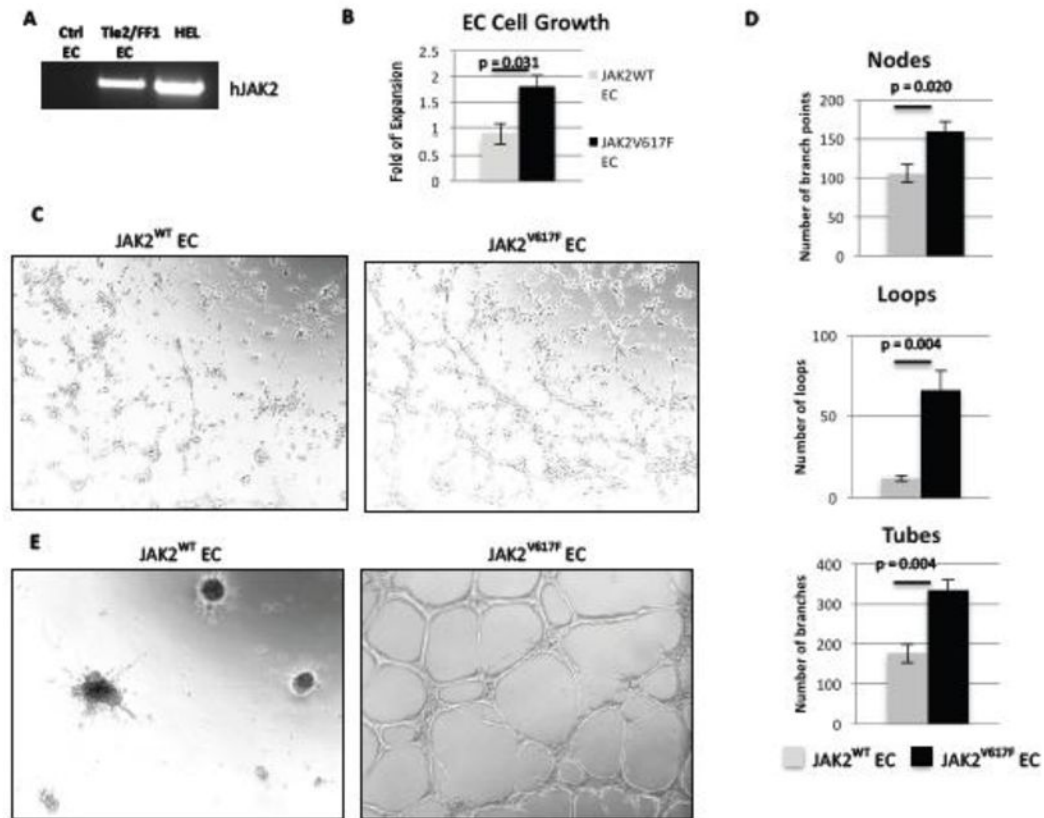


Figure 1. JAK2^{V617F}-mutant ECs have increased cell proliferation and angiogenesis *in vitro* compared to JAK2^{WT} ECs

(A) As determined by RT-PCR, human JAK2^{V617F} was expressed in ECs from Tie2/FF1 mice, but not in ECs from control mice. Human erythroleukemia (HEL) cells were used as the positive control. (B) JAK2^{V617F} ECs proliferated more than JAK2^{WT} EC (2.0-fold, $p = 0.031$). Results of 2 independent experiments (with triplicate in each experiments) are shown here. (C) JAK2^{WT} and JAK2^{V617F} ECs (6×10^4) were seeded in Matrigel matrix. EC tube formation was observed after an 8-hour incubation. A representative picture is shown. Magnification: 100 \times . (D) Quantification of tube formation was performed on images taken at 40 \times magnification by counting the number of nodes (or branch points), loops, and tubes in 4 non-overlapping fields. Results are expressed as the mean \pm SEM ($n=4$). Data are from one of two independent experiments that gave similar results. (E) The tubular structures formed by the JAK2^{V617F} ECs *in vitro* were more stable than the JAK2^{WT} ECs after 24-hour incubation.

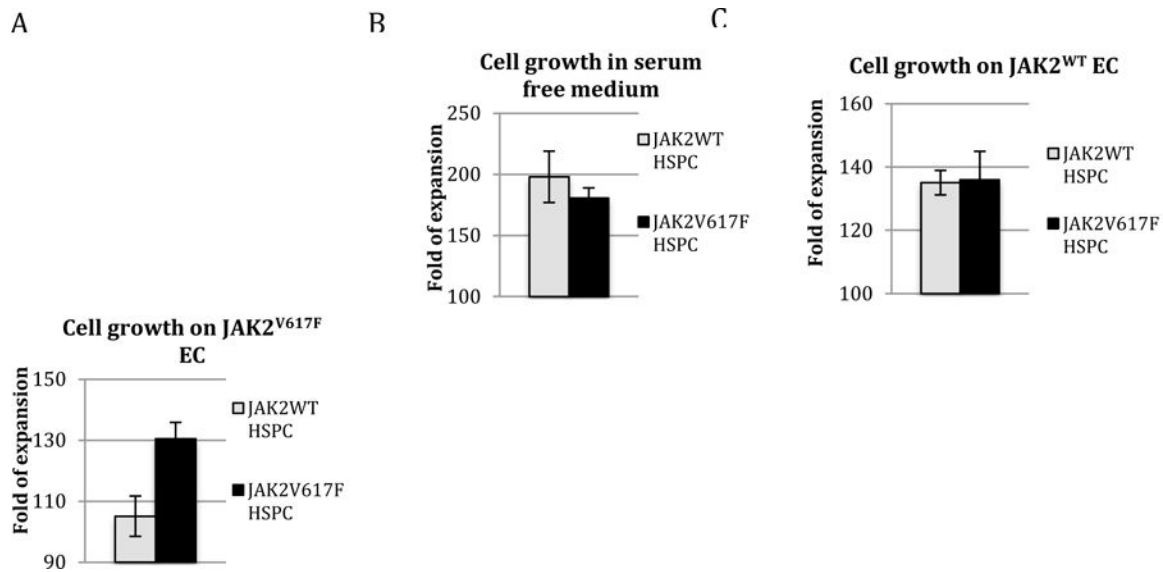


Figure 2. JAK2^{V617F} vascular niche contributes to the growth advantage of JAK2^{V617F} HSPC over JAK2^{WT} HSPC

JAK2^{WT} (from control mice) and JAK2^{V617F} (from Tie2/FF1 mice) Lin⁻cKit⁺ HSPCs were cultured on a feeder layer of JAK2^{WT} or JAK2^{V617F} ECs under serum-free conditions. (A) There was no significant difference between JAK2^{WT} and JAK2^{V617F} HSPC proliferation *in vitro* in SFEM. (B) There was no significant difference between JAK2^{WT} and JAK2^{V617F} HSPC proliferation when co-cultured with JAK2^{WT} EC. (C) JAK2^{V617F} HSPC displayed a relative growth advantage over the JAK2^{WT} HSPC when co-cultured with JAK2^{V617F} EC. Cell proliferation was shown as fold of expansion which is the ratio of the final cell count to starting cell count. The results were expressed as mean \pm s.e.m. (n=3) Data are from one of two independent experiments (with triplicates in each experiment) performed by two investigators (C.L. and H.Z.) that gave similar results. * p < 0.05

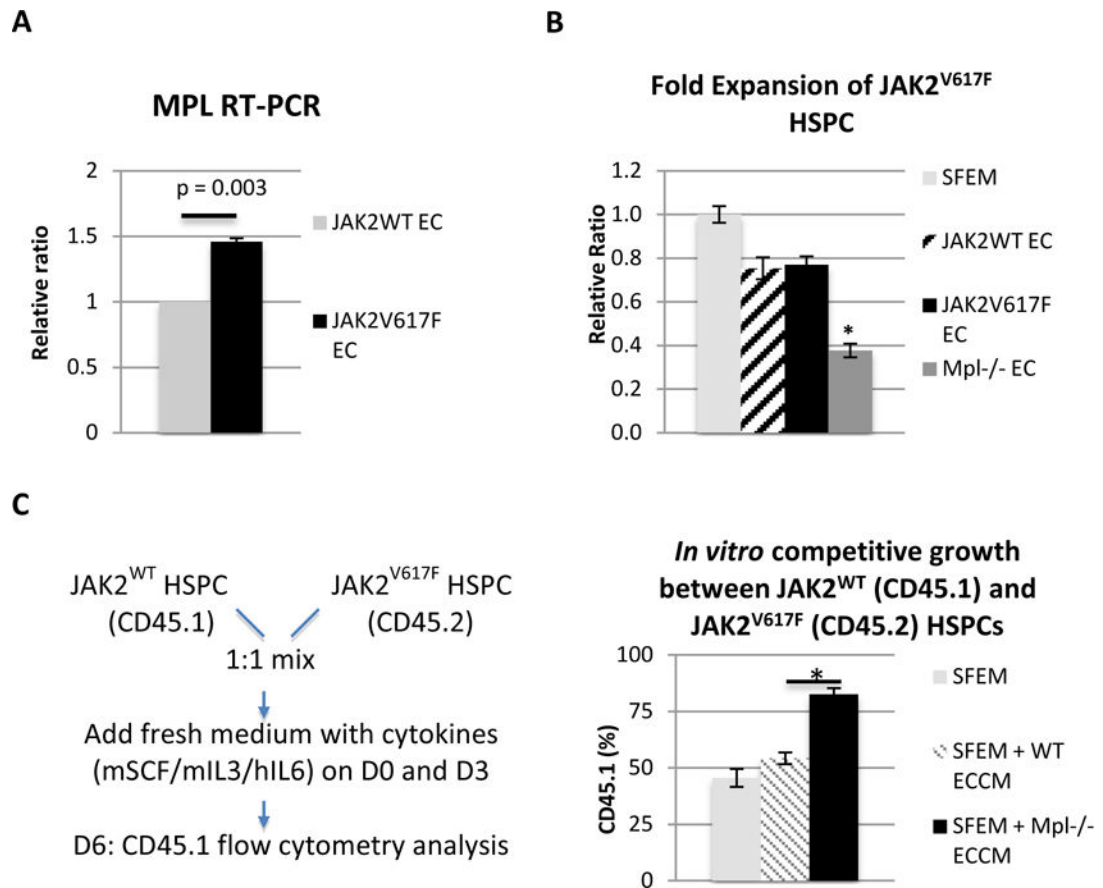


Figure 3. MPL receptor on vascular ECs contributes to JAK2^{V617F} HSPC maintenance/expansion in MPN

(A) MPL expression was increased by 46% in the JAK2^{V617F} ECs compared to JAK2^{WT} ECs ($p = 0.003$). MPL expression in JAK2^{V617F} ECs was shown as the relative ratio compared to its expression in JAK2^{WT} ECs which was set as “1”. (B) JAK2^{V617F} HSPC (Lin-cKit⁺) cell proliferation was significantly decreased when co-cultured with MPL^{-/-} EC, compared to when co-cultured with JAK2^{WT} ECs or JAK2^{V617F} ECs. Cell proliferation in EC co-culture was shown as the relative ratio compared to cell proliferation in SFEM which was set as “1”. (C) Left: experimental design of an *in vitro* competitive growth assay where JAK2^{WT} HSPCs (CD45.1) and JAK2^{V617F} HSPCs (CD45.2) were cultured together (1:1 mix) in the presence of ECCM collected from either WT EC or MPL^{-/-} EC. Right: while there were equal numbers of JAK2^{WT} HSPCs (CD45.1) and JAK2^{V617F} HSPCs (CD45.2) in the presence of WT ECCM, there were significantly more JAK2^{WT} HSPCs than JAK2^{V617F} HSPCs in the presence of MPL^{-/-} ECCM. The results were expressed as mean \pm s.e.m. ($n=3$). Data were from one of two independent experiments (with triplicates in each experiment) performed by two investigators (C.L. and H.Z.) that gave similar results.