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JAK2 V617F Analysis in Indonesian Myeloproliferative Neoplasms Patients

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ABSTRACT

Background: Three subtypes of myeloproliferative neoplasms (MPNs): Polycythemia vera (PV), essential thrombocythemia (ET), and primary myelofibrosis (PMF) showed overlapping phenotype. There has been no specific cytogenetic marker identified in these subtypes. JAK2 gene has a critical role in the pathogenesis of MPNs. Similar mutation, namely JAK2 V617F mutation, was found in PV, ET and PMF.

Objective : This study was done to define the prevalence of JAK2 V617F mutation in Indonesian MPNs patients.

Methods : This is a cross-sectional study of 187 patients who were referred to Center for Biomedical Research (CEBIOR) for JAK2 V617F mutation analysis. The study period was November 2010 until November 2015. It was analysed using Amplification Refractory Mutation System Polymerase Chain Reaction (ARMS-PCR) from peripheral blood vein. Clinical data were secondary data retrieved from hospital medical records.

Results : The prevalence of JAK2 V617F mutation in Indonesian MPNs patients was 107 out of 188 patients (56.92 %). Mutation prevalence distribution for each subtypes were 43 out of 70 (61.43 %) in PV, 25 out of 53 (47.17 %) in ET, 4 out of 6 (66.67 %) in PMF, whereas in unspecified MPN/MPD/MDS 35 out of 59 (59.32 %).

Conclusion : The prevalence of JAK2 V617F mutation was found comparable with previous studies in Indian MPNs. JAK2 V617F testing should be incorporated in the management therapy of MPNs in Indonesia.

Keywords: ARMS-PCR, JAK2 V617F, ET, MPN, PMF, PV

INTRODUCTION

Myeloid malignancies are stem cell-derived and clonal disorders, consist of three wide-ranging clinicopathologic categories: acute myeloid leukemia (AML), myelodysplastic syndrome (MDS) and myeloproliferative neoplasm (MPNs). MPNs were first acknowledged by William Damashek in 1951¹ The classic MPNs were grouped into four subtypes, namely polycythemia vera (PV), essential thrombocythemia (ET), primary myelofibrosis (PMF) and chronic myelogenous leukemia (CML).

They were initially grouped based on their common phenotype of proliferation. Because of their similarities in increasing mature peripheral blood cells and overlapping phenotype, diagnosis has been difficult to be established in the past. It was believed that it came from similar unknown mechanism.¹ The North American Association of Central Cancer Registries (NAACCR) stated that the age-adjusted incidence rate was 2.1 per 100,000 in 2001-2003.² For each disease subtype, an earlier study in Sweden reported the annual prevalence per 100,000 inhabitants for PV, ET, and PMF were 2-2.8; 1.5; and 0.4 respectively.^{3,4}

The important issues in the course of MPNs are thrombosis, hemorrhage, evolution to post-polycythemic or post-thrombocytopenic myelofibrosis and AML transformation.² Thrombosis and bleeding are the leading causes of morbidity in MPNs.⁵ In one-third of MPNs patients, early vascular events constitutes first disease manifestation.⁶ Even though thrombosis is the most frequent complication in MPNs, but bleeding is more observed in ET.⁷

The understanding of the molecular pathogenesis of myeloid malignancies has fundamentally derived from the identification of t(9;22)(q34;q11) or Philadelphia chromosome in CML. However, in many patients with MPNs, no specific abnormality has been identified to date. The frequency of cytogenetic abnormalities in the Philadelphia-negative MPNs varies from approximately 40% in PMF to 3% in ET.⁸ The spectrum of aberrations is heterogeneous, ranging from gains and losses of genetic material to structural changes including unbalanced translocations.⁹ The role of cytogenetic abnormalities as a prognostic marker in PMF has been suggested, both at the time of diagnosis and later during disease course.¹⁰

In molecular level, Philadelphia chromosome is derived from two genes fusion, BCR-ABL on the 22q- and the reciprocal ABL-BCR on 9q-, resulting a chimeric gene BCR-ABL. The identification of the BCR-ABL gene and consequent protein led to the production of small-molecule drugs, proposed to hinder BCR-ABL tyrosine kinase activation by competitive binding at the ATP-binding site: Imatinib mesylate (IM).¹¹ IM turn out to be the first drug of choice in chronic phase CML, as a result of its high efficacy, low toxicity and capacity to preserve strong hematological and cytogenetic responses.¹¹

Several recent discoveries have identified a central role of protein tyrosine kinase (PTK) in the pathogenesis of MPNs. Several groups reported the discovery of *JAK2* V617F mutation in early 2005.¹²⁻¹⁴ Baxter *et al* (2005) found a single base substitution, guanine to thymine change at 1849, which resulted in the change of valine to phenylalanine in exon 14 of the pseudokinase domain of tyrosine kinase *JAK2* (Janus Kinase 2) gene in 97% PV, 57% ET and 50% PMF.¹² This mutation results in a gain of function due to the constitutive activation of tyrosine kinase-dependent cellular signaling pathways, particularly of the JAK-STAT (Signal Transducers and Activators of Transcription). The pathway is principal in regulation of cell proliferation, differentiation and apoptosis in.¹⁴

JAK2 V617F mutation as a common genetic abnormality in PV, ET, and PMF had pointed the possibility of using tyrosine kinase as a valid therapy target. The use of drug that targets the tyrosine kinases is expected to follow the efficacy of IM and other tyrosine kinase inhibitors in CML.^{2,7}

MPNs patients who carried *JAK2* V617F mutation have been associated with older age at diagnosis (ET and PMF),¹⁵ higher hemoglobin level (ET and PMF),¹⁶ leukocytosis (ET and PMF),¹⁵ lower platelet count (ET), larger spleen size (PV, ET and PMF),¹⁷ the need for splenectomy,¹⁷ and leukemic transformation¹⁷. Patients with mutation have been associated with shorter survival in PMF, but less likely to require blood transfusion.¹⁸

Dunlap *et al* studied the correlation between cytogenetic abnormalities with disease stage and *JAK2* V617F status in MPNs and MDS/MPNs patients. Cytogenetic data were available in ninety-seven out of 179 cases (54,19%). *JAK2* V617F positive group showed higher frequency of chromosomal abnormalities (51% vs. 27%). In *JAK2* V617F positive group, the commonest abnormalities were found in chromosome 9, chromosome 7, chromosome 20q, while 13q and trisomy 21 were frequent in *JAK2* V617F negative group. Chromosome 7 and complex abnormalities were associated with blastic transformation.¹⁹

This study aim was to define the prevalence of *JAK2* V617F mutation in Semarang MPNs patients.

METHODS

This is a cross-sectional study of 187 patients who were referred to Center for Biomedical Research (CEBIOR) for *JAK2* V617F mutation analysis. Myeloproliferative neoplasm's consisted of PV, ET, PMF and/or MPNs unclassified subtypes. The diagnosis was determined by the referring clinician according to clinical symptoms and signs, and other supporting data. The study period was November 2010 until November 2015.

DNA was extracted from peripheral blood using the salting out method at the Center for Biomedical Research (CEBIOR) Faculty of Medicine, Diponegoro University (FMDU). *JAK2* V617F mutation was obtained using the Amplification Refractory Mutation System (ARMS-PCR) according to Baxter *et al* with some minor modifications.¹² The principle of the method is using one common reverse primer and two forward primers. One forward primer would be specifically paired to the mutant *JAK2* (if the patient carry the mutation) and the other forward primer would be paired with the wild-type (if the patient did not carry the mutation). The third base from 3' end was intentionally mismatched to maximize the allele specificity. The appearance of two bands at 364 bp and 203 bp is determined as positive, and single band at 364 bp as negative. The 364 bp band also acted as internal control for every sample (See Figure 1). DNA Sequencing to ascertain the presence of G to T nucleotide substitution at position 1849 was done in The Agency for The Assessment and Application of Technology/*Badan Pengkajian dan Penerapan (BPPT)* in Jakarta (See Figure 2).

Patient's clinical data, the result of routine hematology examination, bone marrow smear analysis, and/or relevant findings were retrieved from hospital medical record and summarized in a case report form for each subject.

The data were tested for mutation status difference in proportions using chi-square test. Unpaired *t*-test was used to compare continuous variables between *JAK2* V617F positive and negative groups such as age, and blood count, except for leukocytes.

The ethical clearance was approved by the Health Research Ethical Committee of MFDU/Dr. Kariadi Hospital, Semarang and Dr. Kariadi Hospital Research

Review Board. Study subjects were provided with informed consent.

RESULTS

During the study period, there were 188 patients from around Central Java who were referred for *JAK2* V617F mutation. (Table 1). There were 59 patients who were diagnosed as unspecified MPNs, myeloproliferative disease (MPD) or myelodysplastic syndrome (MDS), while there were 70 patients who were diagnosed as PV or having high hemoglobin level, 53 patients diagnosed as ET or showing thrombocytosis, and 6 diagnosed as myelofibrosis (See Table 1). Among the four groups, only the ET/Thrombocytosis group showed lower proportion of *JAK2* V617F-positive mutation.

Table 1. Referral diagnosis for *JAK2* V617F analysis

MPNs Referral Diagnosis	<i>JAK2</i> V617F		P
	(+) n(%)	(-) n(%)	
MPN/MPD/MDS	35 (59.32)	24 (40.68)	0.39
PV	43 (61.43)	27 (38.57)	
ET/Thrombocytosis	25 (47.17)	28 (52.83)	
Myelofibrosis	4 (66.67)	2 (33.33)	
	107 (56.92)	81 (43.08)	

Other than diagnosis in MPNs, several of the study subjects had other diagnosis in hematology or vascular disease (See Table 2). These conditions showed a possibility that they also experienced vascular events, especially thrombosis that was related to their MPNs diagnosis and *JAK2* V617F mutational status. Diagnosis involving other systems showed that elevation of blood count (in PV and ET) might be an incidental finding that was found during the diagnosis work-up, not solely derived from symptoms and signs of MPNs

Table 2. Accompanying hematology and/or vascular event (thrombosis) diagnosis

Diagnosis	n	<i>JAK2</i> V617F	
		Positive	Negative
Aplastic anemia	1	1	0
Anemia	1	0	1
Trombocytopenia	3	2	1
Pancytopenia	1	0	1
Leukocytosis	5	4	1
Leukemia	2	2	0
Acute Myeloid Leukemia, Non Haemorrhagic Stroke	1	0	1
Combination of Anemia, Leukocytosis, Splenomegaly	3	2	1
CML	1	0	1
Splenomegaly	3	2	1
Non Haemorrhagic Stroke	1	1	0
Stroke	1	0	1
Iischemic Heart Disease	1	0	1
Acute Myocardial Infarction	1	1	0
Old Myocardial Infarction	2	2	0
Acute Coronary Syndrome	1	0	1
Pperipheral Artery Disease	1	0	1
Deep Vein Thrombosis	2	0	2
Thalassemia	1	0	1
Total	32	17	15

Table 3. Accompanying diagnosis involving other systems

	<i>JAK2</i> V617F mutation		
	n	Positive	Negative
Grade II hypertension, Multiple Renal cysts, renal insufficiency, hypoalbumin, lung infiltrate, pyuria, hematuria	1	1	0
Grade II hypertension, chronic kidney disease stage IV, renal insufficiency, hypoalbumin, lung infiltrate, pyuria, hematuria	1	1	0
Right ovarian cyst	1	1	0
Wide stomatitis, esophageal candidiasis, pleuropneumonia, gastric ulcer	1	0	1
Melena	1	1	0
Gout Arthritis	1	1	0
Duplex rhinosinusitis maxillaris	1	0	1
Wound dehiscence	1	0	1
Post-cranectomy	1	1	0
Fever	2	1	1
Liver chirosis	1	1	0
Diabetic ulcer	1	0	1
Vertigo	1	0	1
Focal Nodular Hypertrophy	1	1	0
Post hematoma exploration	1	1	0
DM, Ikterus	1	1	0
Hemiparesis	1	1	0
Urine retention	1	1	0
Space occupying lesion	1	0	1
Post-Odenectomy	2	1	1
	22	14	8

Table 4 showed that positive cases were older than negative cases (54.43 ± 11.857 vs. 49.01 ± 16.351), although it was insignificant. In both sex groups, more positive cases were observed in the study subjects. Blood count at diagnosis showed that positive cases had higher hemoglobin and hematocrite values. In the contrary, leukocyte and platelet counts were lower in positive group. Four *JAK2* V617F mutation positive subjects (3 PV and 1 ET) was confirmed with sequencing, and showed the present of G to T nucleotide substitution (Figure 2).

Table 4. Demographic characteristics

	<i>JAK2</i> V617F mutation		P
	Positive	Negative	
Age (year)	54.43 ± 11.857	49.01 ± 16.351	0.09
Sex	62/113	45/75	0.49
Male	62 (54.86%)	51 (45.14)	
Female	45 (60%)	35 (40%)	

Table 5. Blood count

	<i>JAK2</i> V617F mutation				p
	N	Positive		Negative	
		mean±SD	n	mean±SD	
Hemoglobin (g%)	22	15.6 ± 5.20	11	12.0 ± 5.66	0.79
Hematocrite (%)	20	44.9 ± 18.03	11	34.4 ± 16.71	0.12
Leukocyte (/mm ³)	20	23.0 ± 13.16	9	49.3 ± 90.74	0.41
Platelet (10 ³ /mm ³)	20	798.4 ± 533.08	11	$1,038.6 \pm 483.65$	0.22

Comparison was tested using unpaired t-test, except for leukocyte (Fischer's exact test)

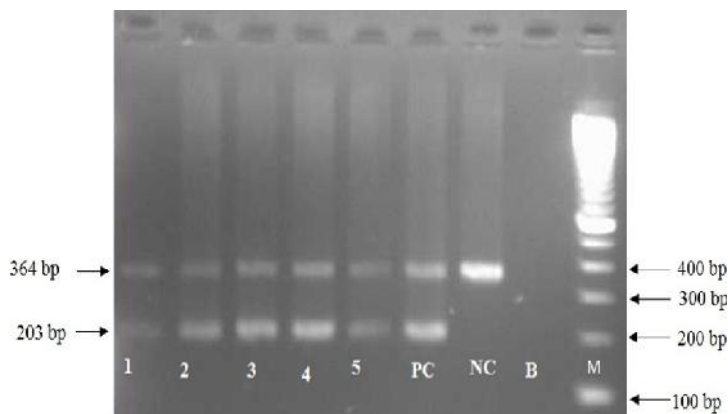


Figure 1. *JAK2* V617F mutation detection using ARMS-PCR. Lane 1 to 5 represented subjects number 12/F/11 to 16/F/11). Upper band at 364 bp were the internal PCR control which appeared in every sample tested. Lower band at 203 bp indicated the presence of mutation. Positive control (PC) was the positive result from previous PCR, negative control (NC) came from non-MPNs individual, (B) is blank, (M) 100 bp marker ladder.

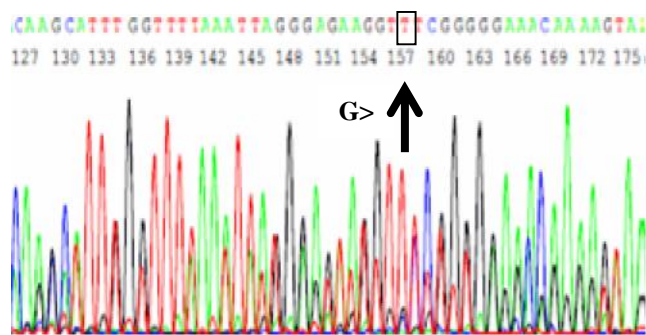


Figure 2. Sequencing profile of a subject (07/F/11) whose PCR positive for *JAK2* V617F mutation using forward primer. Arrow indicated the presence of mutation c.1849G>T in exon 14 of *JAK2* gene.

DISCUSSION

This study results supported the fact that *JAK2* V617F mutations were found in majority of MPNs patients (56.92 %). A study in India reported 68% of MPNs patients harbored this mutation.²⁰ The distribution of *JAK2* V617F mutation in the disease subtypes were similar with other reports in Caucasian,^{12,13,17} except for PV. This was possibly caused by different cut-off point that has been used by referring clinicians.

Although the routine blood count values were not significantly different between the two groups, the profile that was observed in our findings was similar with other studies. The conclusion of a meta-analysis measuring the effect of *JAK2* V617F on thrombotic risk in 492 studies suggested that *JAK2* V617F-positive patients were older at diagnosis, had higher hemoglobin, but lower thrombocyte counts.²¹ The most possible mechanism underlying these observations was that *JAK2* V617F mutation caused hypersensitivity to cytokine stimulation.¹³

A study which correlated *JAK2* mutation status, hemostatic risk factors and thrombophilic factors in ET patients stated that *JAK2* V617F mutation have been associated with older age at diagnosis (ET and PMF),¹⁵ higher hemoglobin level (ET and PMF),¹⁶ leukocytosis (ET and PMF),¹⁵ lower platelet count (ET)¹⁷. In addition, *JAK2* V617F mutational status predicts progression to large splenomegaly and leukemic transformation in primary myelofibrosis in PV, ET and PMF.¹⁷ Shorter survival in PMF patients have been associated with the *JAK2* V617F mutation, but less likely to require blood transfusion.¹⁸ High proportion of *JAK2* V617F mutation which was found in our study subjects should be carefully followed-up by the referring clinicians, in concern of the disease course related to positive results.

Among patients with hemoglobin level above normal cut-offs, the presence of *JAK2* V617F mutation allows the diagnosis of PV in >95% cases. Less than 2% of these patients might carry *JAK2* exon 12 mutations.²² Further study should elucidate the presence of other *JAK2* mutations in *JAK2* V617F negative patients.

The demonstration of *JAK2* V617F mutation in patients with thrombocyte count < 600.000/dL (according to 2001 WHO classification system),²³ supported the use of lower cut-off of thrombocyte count for ET diagnosis in the 2008 revision (changed to > 450.000/dL) and might help the exclusion of reactive thrombocytosis.^{16,24} However, *JAK2* V617F mutation was found only in 47.17% ET and 66.67% PMF cases. Thus, bone marrow smear analysis remains as an important diagnosis tool of ET, since bone marrow appearance in ET patients is distinct from typical PV or PMF, and *vice versa* for PMF.²⁵

The use of peripheral blood, as in our study, has been reported as objective and made the *JAK2* mutations screening become more accessible and practical in order to study patients suspected having MPNs, compare to invasive test like bone marrow examination or expertise-dependent test such as direct red cell mass measurement.²⁸ However, this molecular testing alone could not outweigh the important baseline information yielded from bone marrow histology and cytogenetic analysis.

Results of studies in familial MPNs about increased risk of first-degree relatives might improve the information that the *JAK2* V617F testing should be provided for families of MPNs patients.

However, there were some limitations of the study. First, only *JAK2* V617F mutation was studied. Despite the fact that the mutation is the commonest in MPNs, a large number of other mutations have been discovered and it has not been understood yet how is the hierarchy in this genetic complexity of MPNs. Second, the diagnosis of MPNs subtypes was made without incorporating red cell mass, serum Epo level or EEC (the complete 2008 WHO diagnostic criteria) although the remaining criteria were sufficient to establish the diagnosis. Third, the research setting was at only one center, given that the prevalence of MPNs is 2,1 per

100.000 individuals a multicenter study might yield higher number of participants.²

CONCLUSION

Supported by the high incidence of *JAK2* V617F mutation in our results, which were comparable with studies for in Indian and Caucasian we would recommend the integration of *JAK2* V617F mutation testing in the diagnosis and management of MPNs cases in Indonesia.

SUGGESTIONS

Further studies should be conducted to explore the association of cytogenetic and *JAK2* V617F mutation, clinical profiles, complications and prognosis in MPNs. As well as the exploration of other mutations that involved in the pathogenesis of MPNs. The main goal of the follow-up studies are to establish *JAK2* inhibitors for *JAK2* V617F-positive patients in clinical practice.

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