

## **RESEARCH ARTICLE**

# Utility of Amplification Refractory Mutation System PCR for the detection of JAK2 mutation in Chronic Myeloproliferative Disorders.

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#### Abstract

Detection of the JAK2 V617F mutation is helpful in differential diagnosis, prognosis, and predication of therapeutic response in CMPD disorders. Because the mutation can be present in a small proportion of granulocytic populations in MPD patients, a highly sensitive detection method is required. The study was aimed to optimize ARMS-PCR for the detection of JAK2V617F mutations .This study was conducted at Prince Fahd Bin Sultan Research Chair .Fifty subjects were registered from the King Khaled Hospital of Tabuk -Northwestern part of Saudi Arabia among which 25 newly diagnosed MPD patients and 25 healthy controls. DNA was extracted done using DNA extraction Kit from Qiagen .ARMS was optimized in which a a primers were designed in such a way that it is able to discriminate among templates that differ by a single nucleotide residue in one tube reaction.Both cases and controls were tested for JAK2 mutations by ARMS-PCR. Demographic data, spleen size, hemoglobin level, white blood cell and platelet counts were recorded. Independent sample t-test was used to study the correlation of JAK2 mutations with age, haemoglobin, blood counts and spleen size. Fisher's exact test was applied to compare disease progression in mutation positive and negative cases. This study was conducted on 25 newly diagnosed MPD patients among which 17 were polycythemia vera,5 Essential thrombocytopenia and 3 primary myelofibrosis and 25 healthy controls. Both cases and controls were tested for JAK2 mutations by ARMS-PCR. The JAK2 mutation was detected in 17/25 samples among which 13 were Polycythemia Vera, 02 from Essential thrombocythemia, 01 from Primary myelofibrosis. The JAK2 mutation was not detected in any of the healthy controls by ARMS-PCR. The main advantage of ARMS was that the amplification step and the diagnostic steps were combined, in that the presence of an amplified product indicates the presence of a particular allele and vice versa. For routine diagnosis, this characteristic of ARMS means that it is a very time-efficient method.It was concluded that the ARMS-PCR assay optimized allows the discrimination between homozygous and heterozygous individuals with the JAK2 V617F mutation and has a key role in acting as a reliable screening test for the presence or absence of the mutation in individuals with MPDs. The test will be potentially useful to follow minimal residual disease when targeted therapies against JAK2 kinase become available. The test is therefore a simple, fast, and inexpensive procedure that does not entail any special equipment other than a thermocycler.

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## **Introduction:-**

Myeloproliferative neoplasms (MPNs) are multipotent hematopoietic stem cell disorders characterized by uncontrolled proliferation of maturing blood cells<sup>1</sup>. Among its types, Chronic myeloid leukemia (CML) is the most common MPN, followed by polycythemia vera (PV), essential thrombocythemia (ET), and idiopathic myelofibrosis (IMF). The pathogenesis of myeloproliferative disorders (MPD), is still largely unknown except the chronic myeloid leukemia (CML), where the presence of the t(9;22) Ph' chromosome and of the underlying molecular abnormality has been known for decades and ultimately led to the development of target-specific drugs, the molecular lesion(s) at the basis of the other MPD has remained elusive.<sup>2-4</sup> JAK2 is a cytoplasmic tyrosine kinase that plays an essential role in the signaling pathways of cytokines and growth factors. An acquired  $G \rightarrow T$  point mutation in exon 12 of the gene encoding the Janus tyrosine kinase 2 (JAK2), leading to amino acid substitution of phenylalanine for a highly conserved valine (V617F), in

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the JH2 pseudo-kinase domain, renders JAK2 kinase constitutively active and leads to cell proliferation in the absence of the growth factors<sup>5-6</sup>. In vivo expression of  $JAK2^{V617F}$  in a murine transplant model induced erythrocytosis<sup>7</sup>. the mechanism of activation of JAK2 kinase activity by mutations in the JAK2 signalling pathway as depicted in figure 1. The JAK2V617F and JAK2 exon 12 mutant kinases bind cytokine receptors and are phosphorylated in the absence of ligand, and lead to ligand-independent activation of downstream signaling pathways.



Figure 1 The mechanism of activation of JAK2 kinase activity by mutations in the JAK2 signaling pathway

The JAK2 mutation has been detected in 65 to 97% of patients with polycythemia vera, 23 to 57% of those with essential thrombocythemia (ET), and 30 to 57% of idiopathic myelofibrosis.<sup>7-8</sup> JAK2 mutation is absent in normal individuals, in patients with secondary erythrocytosis and thrombocytosis, or in patients with chronic myelogenous leukemia; detection of the mutation thus aids in differential diagnosis in cases with similar clinicopathological features.<sup>9</sup>Methods for detection of JAK2<sup>V617F</sup> mutation based on the analysis of genomic DNA obtained from peripheral blood granulocytes have been devised and already entered the clinical practice. Because V617F is an acquired mutation that can be present in a small proportion of granulocytic populations in some cases, especially of ET, a highly sensitive detection method is essential<sup>10</sup> .Methodologies reported in the literature include direct sequencing,<sup>6</sup> PCR-restriction fragment length polymorphism,<sup>10</sup> allele-specific polymerase chain reaction (PCR),<sup>11</sup> pyrosequencing,<sup>12</sup> and amplification refractory mutation system (ARMS)-PCR.<sup>13</sup> The modified assay detects mutation with a highly reproducible analytic sensitivity of 0.05 to 0.1%, making it suitable for the detection of low levels of JAK2 mutation. The test will be potentially useful to follow minimal residual disease when targeted. The merits and demerits of the various methods used for JAK2 mutation detection have been by studied described for principles and potential clinical utilities of each assay<sup>14</sup>. Some have adapted the use of mRNA as the basis for testing for JAK2 mutations and have shown that RNA allows more sensitive detection of mutations than does DNA at early stages of disease The use of RNA rather than DNA provides the additional advantage of capturing abnormalities in platelets and detecting alternatively spliced transcripts<sup>15</sup>. The reported frequency of  $JAK2^{V_{617F}}$  mutation is around 80% (range, 65–100%) in PV, 40% (range, 23–57%) in ET and 55% (range, 35–95%) in IM patients<sup>16</sup> . $JAK2^{V_{617F}}$  mutation is found in either the heterozygote or the homozygote status, the latter arising from mitotic recombination; there are significant differences in the incidence of homozygosity, which involves about 30% of patients in PV and IM and less than 4% in ET.

There are indications that the clinical phenotype might be dependent on the proportion of mutated allele, as the expression levels of some aberrantly regulated genes in MPD, such as *PRV-1* or *NF-E2*, have been found higher in homozygote than in heterozygote patients.<sup>17-18</sup> In a large survey on ET patients, the presence of the mutation was associated to clinical and biological features suggestive of a PV-like disease, thus supporting the view that these two disorders represent a biological continuum; furthermore, according to the presence of the *JAK2*<sup>V617F</sup> mutation, ET patients could be divided into two distinct subpopulations characterized by different disease presentation, clinical complications and response to therapy . We aimed to optimize the diagnostic values of the ARMS-PCR method for detecting mutations in exon 14 of the *JAK2* gene to diagnose myeloproliferative disorders.

## Material and Methods:-

#### Patients and Controls:-

#### Selection criteria of patients:-

- The study was carried out at Prince Fahd Bin Sultan Research chair, Cancer Molecular Genetics Lab, Department of Medical Lab technology, University of Tabuk. The study was ethically approved from the Ethics committee, University of Tabuk.
- The subjects were recruited from the hematology OPD of King Khaled hospital, Tabuk and all the subjects obtained consent form before the sample collection.

#### Inclusion criteria for myeloproliferative disorders:-

- Myeloproliferative disorders with Ph negative .
- Clinically confirmed cases of polycythemia vera (PV), essential thrombocytosis (ET), myelofibrosis (MF).
- ✤ All participants were of Saudi origin.

### Inclusion criteria for controls:-

- All controls specimens were timed around routine blood draws that are part of routine workout, and hence would not require additional phlebotomy.
- ✤ All participants were of Saudi origin

## **Exclusion Criteria:-**

- ✤ Patients with any other significant malignancy.
- \* Non-Arab Saudi or recently-naturalized Saudi Arabia were excluded

#### Sample Size:-

★ The study was conducted on 30 specimens .Twenty were healthy controls and 25 were clinically confirmed Myeloproliferative disorders.

#### **Experimental Design:-**

## Genomic DNA extraction from Blood samples of MPD patients and controls:-

The Blood samples were collected by venipuncture in EDTA tubes from MPD patients and healthy controls .The DNA extraction was done by using DNeasy Blood Kit cat 69506 from Qiagen as per the manufactures instructions. The QIAamp DNA Blood Mini Kit provides silica-membrane-based DNA purification. The QIAamp DNA Blood Mini Kit was designed for processing up to 200µl fresh or frozen human whole blood. QIAamp Mini spin columns can be easily processed in a centrifuge or on vacuum manifolds. Purification of DNA using the QIAamp DNA Blood Mini Kit was fully automated on the QIAcube .The DNA extracted was dissolved in nuclease-free water, and stored at 4°C until use. The DNA quality and yield was assessed using Nanodrop (optical density) and agarose gel electrophoresis

#### Amplification ARMS primers for the JAK2 mutation:-

ARMS-PCR was performed in a final volume of 5 uL containing 2uL of 50ng genomic DNA, 12.5 uL of PCR-master mix purchased from Epigentek (USA),0.25 uL of 25 pmol/L of each primer (Table 1) The Methylamp Taq 5x PCR Mix was premixed ready-to-use solution containing all reagents required for PCR (except template, primers and water). Methylamp Taq DNA polymerase ,5x Reaction Buffer: 1 0.4 M Tris-HCl, 0.1 M (NH4)2SO4, 0.1% w/v Tween-20 ,12.5 mM MgCl2: 1x PCR solution – 2.5 mM MgCl2 ,1mM dNTPs of each and remaining nuclease free ddH<sub>2</sub>O as depicted in Table 2. Finally the 2ul of DNA was added from each patient and control separately. The thermo cycling conditions for the PCR amplification of ARMS primers for JAK2 mutation were initial denaturation at 95<sup>o</sup>C for 10 minutes followed by 40 cycles 95 <sup>o</sup>C for 45 sec, 57.9 <sup>o</sup>C for 45 sec, and 72 <sup>o</sup>C for 45 sec followed by the final extension at 72 <sup>o</sup>C for 10 minutes.

Primer	Sequence	Annealing Temp	PCR product
Forward outer	5'- TCCTCAGAACGTTGATGGCAG- 3'	57.9	463bp
<b>Reverse</b> outer	5'- ATTGCTTTCCTTTTTCACAAGAT-3'		
Forward wt	5'-GCATTTGGTTTTAAATTATGGAGTATATG-3'		229bp
Reverse mut	5'-GTTTTACTTACTCTCGTCTCCACAAA-3'		279bp

#### Table 1:- Primers designed for ARMS PCR

Reagent	1x	5 x
PCR master mix	5ul	25ul
Forward outer	0.25 ul	1.25 ul
Reverse outer	0.25 ul	1.25 ul
Forward wild	0.25 ul	1.25 ul
Reverse mutant	0.25 ul	1.25 ul
Nuclease free water	17.0 ul	85ul
Total volume	23ul	110ul

Finally 2 ul DNA was added.

#### **Results and Discussion:-**

The purity of DNA extracted was checked by using Nanodrop1000 Spectrophotometer from Thermo .All DNA samples were screened for purity by measuring optical density (OD) at 260nm (OD<sub>260</sub>) and 280 nm (OD<sub>280</sub>). DNA concentration (ug/ml) was calculated based on the OD<sub>260</sub> reading .A ratio of ~1.8 was generally accepted as "pure" for DNA .The optimal rang of ratio obtained were 1.7-2 .The DNA 260/230: 1.7-2; Pure DNA must have a 260/280 ratio between 1.8 and 2. The quality of DNA was checked by running in 1% gel electrophoresis as shown in figure 2.

Figure 2:Genomic DNA preparation from peripheral blood of myeloproliferative patients and healthy controls



b). Genomic DNA preparation from peripheral blood of healthy controls

#### Amplification Refractory Mutation System (ARMS-PCR) for JAK2V617F mutation:-

Tetra-primer ARMS-PCR was established for the analysis of single nucleotide polymorphisms (SNP), which employs two primer pairs to amplify, respectively, the two different alleles of an SNP in a single PCR reaction. Two forward and two reverse primers are used in different combinations to generate three potential PCR products. Primers FO and RO flank the exon 14 of the *JAK2* gene, resulting in a band of 463bp to control for DNA quality and quantity. Primers Fwt and RO amplify a wild-type allele, generating a band of 229bp, and primers FO and Rmt generate a band of 279 bp from the mutant allele as shown in figure 3.



FIGURE 3. PCR amplification ARMS-PCR primers for the detection of JAK2V617F Mutation

JAK2 (V617F) mutation analysis by ARMS PCR in leukemia patients: L= 100bp Ladder. P1,P5,P6 and P7 = Mutation Negative. P2,P4,P4,P8 and P9= Mutation Positive.

The intended mismatches in Fwt and Rmt are shown in lowercase, and genotype-specific nucleotides are underlined as shown in figure 4. The amplification Refractory Mutation Screening (ARMS) PCR permits a single base change to be detected under ideal PCR conditions. This is ideal for detection of the single base  $G \rightarrow T$  transversion associated with the JAK2 mutation in question. The ARMS-PCR technique uses 4 primers as shown in figure 4 as follows; a forward outer primer, a reverse outer primer ,a forward inner wild type specific primer and a reverse inner mutant specific primer. It requires two temperature programs during the PCR reaction. Due to the positioning of the outer primers at varying distance from the site of the mutation there is the generation of three fragments in this example in a heterozygote: two small allele specific fragments and a large control PCR product. DNA Fragments can be distinguished via electrophoresis on agarose gel.





Performing a dilution series indicates the level of sensitivity of ARMS-PCR to be 1-2 %. The assay allows discrimination between homozygous and heterozygous individuals with the JAK2 V617F mutation and has a key role in acting as a reliable screening test for the presence or absence of the mutation in individuals with MPDs. An acquired mutation in the *JAK2* gene is a cytoplasmic tyrosine kinase that plays an essential role in the signaling pathways of cytokines and growth factors. The mutation 1849 G>T, which leads to amino acid substitution of phenylalanine for a highly conserved valine (V617F), (as shown in figure 5) renders JAK2 kinase constitutively active and leads to cell proliferation in the absence of the growth factors



Figure 5. Primer designing for ARMS-PCR assay of JAK2V617F Mutation

The study was conducted on 50 specimens 25 were controls and 25 were MPD cases. The 25 newly diagnosed MPD patients among which 17 were polycythemia vera, 5 Essential thrombocytopenia and 3 primary myelofibrosis and 25 healthy controls depicted in figure 6. Both cases and controls were tested for JAK2 mutations by ARMS-PCR. The JAK2 mutation was detected in 17/25 samples among which 13 were Polycythemia Vera, 02 from Essential thrombocythemia, 01 from Primary myelofibrosis as depicted in figure 7. The JAK2 mutation was not detected in any of the healthy controls by ARMS-PCR.





JAK2 V617F - DNA Reference Standards (Horizon diagnostics) for wild as well as mutant allele were used along with ARMS-PCR assays. ARMS assay proved to be highly sensitive technique and can be used to monitor residual disease after treatment. The JAK2 mutation was discovered using direct sequencing after PCR. However, the method only detects the mutation present in more than 20 to 30% of total DNA. The JAK2 mutation has become an important diagnostic criterion for Philadelphia chromosome-negative myeloproliferative neoplasms (Ph-neg MPN), especially in polycythemia vera (PV).<sup>19</sup> accordingly, it has been included in the World Health Organization diagnostic criteria for Ph-neg MPN<sup>20</sup>. In Jak2 gene new exon 12 mutations involved in the pathogenesis of PV and idiopathic erythrocytosis have been described. These mutations are also important for the diagnosis of PV, as approximately 3% of PV cases have mutations in this exon. <sup>21</sup>In addition, other *JAK2* exon 14 mutations that are distinct from JAK2 V617F, such as C616Y, D620E, and C618R, have been detected in patients with myoproliferative neoplasms (MPN). <sup>22</sup> Despite this, the described *JAK2* mutations are neither specific for this group of diseases, nor are they present in all patients with classical Ph-neg MPN (especially in essential thrombocythemia [ET] and primary myelofibrosis). In addition, infrequent occurrence of the JAK2 V617F mutation has been reported in chronic myelomonocytic leukemia, atypical MPN, myelodysplastic syndromes, systemic mastocytosis, and acute myeloid leukemia. <sup>23</sup>Detection of the mutation is helpful in differential diagnosis, prognosis, and predication of therapeutic response.

Amplification Refractory Mutation System (ARMS-PCR) has also been termed allele-specific PCR or PCR amplification of specific alleles (PASA). The amplification refractory mutation system (ARMS) is an amplification strategy in which a polymerase chain reaction (PCR) primer is designed in such a way that it is able to discriminate among templates that differ by a single nucleotide residue. Thus, an ARMS primer can be designed to amplify a specific member of a multi-allelic system while remaining refractory to amplification of another allele that may differ by as little as a single base from the former.

Several laboratory techniques have been developed for JAK2 V617F genotypic analysis. Because JAK2 mutation is an acquired mutation that can be present in a small proportion of granulocytic populations in some cases, especially of ET, a highly sensitive detection method is essential. Different sensitivity of various assay methods partially accounts for the wide range of mutation frequencies reported in the literature.<sup>23</sup> As a matter of fact, re-evaluation of the same cases with a more sensitive technique has increased the detection rate from 73 to 97% in patients with polycythemia vera.<sup>24</sup> In addition, a highly sensitive technique is potentially useful to monitor residual disease after treatment. Methodologies reported in the literature include direct sequencing,<sup>24</sup> allele-specific polymerase chain reaction (PCR),<sup>24</sup> PCR-restriction fragment length polymorphism,<sup>25</sup> pyrosequencing,<sup>26</sup> and amplification refractory mutation system (ARMS)-PCR<sup>26</sup>..The main advantage of ARMS is that the amplification step and the diagnostic steps are combined, in that the presence of an amplified product indicates the presence of a particular allele and vice versa. For routine diagnosis, this characteristic of ARMS means that it is a very time-efficient method. However, this combination of the amplification and diagnostic steps are separated, e.g., PCR followed by restriction enzyme analysis. Moreover, the data show that direct sequencing is not an adequate technique for exon 12 mutation identification; therefore, appropriate methodology should be considered for using this molecular marker in the process of diagnosis. It has been have demonstrated that this high sensitivity is not

accompanied by any significant compromise in the diagnostic specificity of the assay. In addition, the ARMS-PCR assay shows different patterns for normal, mutant, or mixed genotypes, minimizing the chance of misinterpretation. With simply one tube of PCR reactions and obviation for special equipment and reagents, ARMS-PCR represents a time- and labor-saving as well as cost-efficient method that is readily and broadly applicable in clinical molecular diagnostic laboratories.

#### Conclusion:-.

It was concluded that the ARMS-PCR assay is an appropriate methodology should be considered for using JAK2 molecular marker in the process of MPD diagnosis and allows discrimination between homozygous and heterozygous individuals with the JAK2 (V617F and in exon 12) mutation and has a key role in acting as a reliable screening test for the presence or absence of the mutation in individuals with MPDs. The test is therefore a simple, fast, and inexpensive procedure that does not entail any special equipment other than a thermocycler.

#### **Competing interests:-**

The authors declare that they have no competing interests.

#### Authors' contributions:-

All authors read and approved the final manuscript. Thanks to all authors for their support and help in this study.

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