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J Phys Pharm Adv 2014, 4(7): 389-395

DOI: 10.5455/jppa.20140726032925



Molecular Identification of Mediterranean Mutation in Patients with Deficiency of Glucose-6-Phosphate Dehydrogenase (G6pd) in North West of Iran

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Abstract

Glucose-6-phosphate dehydrogenase (G6PD) is a highly polymorphic enzyme that is encoded by human X-linked gene. G6PD deficiency is the most common enzymopathy in human with approximately 400 million people affected worldwide. According to the previous studies, Mediterranean mutation (C563T) is one of the most common G6PD enzyme mutations in Iranian population with a prevalence range of %63 to %86/4. This study was conducted in order to determine Mediterranean mutation, through molecular method, in the population of the North-west of Iran. In this study, by Rapid Genomic DNA Extraction (RGDE) method, from 77 blood samples of unrelated male and female patients with genetic deficiency of G6PD, DNA was extracted and after digestion by *MboII* enzymes, in order to search for Mediterranean mutation, they were analyzed by means of PCR-RFLP and sequencing methods. Of the 77 patients, 48 patients (37 male, 11 female) had a Mediterranean mutation frequency of 62/3 %. 29 patients (37/7%) had mutations other than Mediterranean type. This study revealed that G6PD is the most common mutation in Iranian North-west population. Further studies are recommended to identify the mutation type of other varieties.

Keywords: Glucose-6-phosphate dehydrogenase, mediterranean mutation, north- west of Iran.

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Received on: 20 Jun 2014

Revised on: 24 Jul 2014

Accepted on: 26 Jul 2014

Online Published on: 30 Jul 2014

Introduction

The enzyme G6PD (Glucose-6-phosphate Dehydrogenase) is one of the most important enzymes in human body, which different cells in human body including red blood cells contain varying amounts of the enzyme (Vulliamy *et al.*, 1992). G6PD catalyzes the first and pace-making reaction of the pentose phosphate cycle (PPC) by which glucose-6-phosphate (G6P) is oxidized to 6-phosphogluconolactone with production of NADPH (Grimes, 1980; Beutler, 2001; Cappellini *et al.*, 2008; Arese *et al.*, 1990). PPC converts glucose into pentose phosphates and produces reducing power in the form of NADPH. In RBCs NADPH is essential for the protection against oxidative stress (Pandolfi *et al.*, 1995). A major role of NADPH is to maintain reduced glutathione (GSH) at a ratio greater than 500:1 over the oxidized form GSSG (Cappellini *et al.*, 2008; Arese *et al.*, 1990). GSH plays a vital role in antioxidant defense by reacting with H₂O₂ and organic peroxides as well as by maintaining thiol groups of Hb, other proteins and enzymes in the reduced state. H₂O₂ can be also detoxified by catalase which is also stabilized by tightly bound NADPH (Gaetani *et al.*, 1996; Scott *et al.*, 1993). The major clinical manifestations of G6PD deficiency are Acute Hemolytic Anemia, Neonatal Jaundice, Mental Retardation, Chronic Renal Failure, Chronic nonspherocytic Hemolytic Anemia. in children or adults with G6PD deficiency using of some substances such as Antimalarial drugs, oxidizing materials, or fava beans (*Vicia faba*) causes severe hemolytic attacks which can be life-threatening (Beutler *et al.*, 1994; Malluh *et al.*, 1992).

The gene encoding G6PD is located on the long arm of the X chromosome (Xq28) and consists of 13 exons and 12 introns, spanning approximately 18 kb. The coding sequencing of the gene is 1,545 bp long, coding for the 515 amino acids of the G6PD primary sequence (Arese *et al.*, 2012).

Due to the location of genes, heredity of G6PD is a X-linked pattern (Mortazavi *et al.*, 2006). Hence, the defective gene in men is fully out broken. In females due to having two X chromosomes, different forms of homozygote and heterozygote are visible (Beutler, 1994).

Glucose-6-phosphate dehydrogenase enzyme deficiency is one of the most frequent human genetic disorders so that more than 400 million people are affected worldwide (Ruwende *et al.*, 1995). However; its prevalence in different populations varies from 1 to 65% (Mason *et al.*, 2007).

Although Iranian population consists of different ethnic groups, but the overall incidence of G6PD deficiency in Iranian population is estimated around 10%-14.9% (Noori-Dalooi *et al.*, 2007). Some studies were carried out on the molecular basis of G6PD deficiency in Iran and showed that the Mediterranean mutation have the highest frequency in Gilan, Mazandaran, Golestan (Noori-Dalooi *et al.*, 2006b). Khorasan, Hormozgan, Sistan&Baluchestan, Yazd and Kerman (Noori-Dalooi *et al.*, 2007).

In most cases displacement of a base at the DNA level, causes the displacement of one amino acid with another amino acid. Normal type G6PD is known type B. About 20 percent of black African have G6PD type (A +) which is normal in terms of activity but is different with type B in terms of one amino acid and can be identified by electrophoresis (Luzzatto, 1973). Some African descents with G6PD type (A-) have unstable G6PD because of mutations in the DNA.

Type Mediterranean G6PD is the most common type of G6PD and its symptoms are more severe than of type (A-). This variant can be found in the Mediterranean regions (Italy, Greece, Sardinia), the Jewish Kurds and Arabs, India and in Southeast of Asia (Luzzatto, 1973; Oppenheim *et al.*, 1993).

According to previous studies, the most prevalent mutations in Iran are Mediterranean, Chatham and Cosenza. The purpose of this study is to investigate the frequency rate of the Mediterranean mutation in north-west of Iran.

Materials and Methods

In this cross-sectional study, 77 Peripheral blood samples (2-5 ml in 300 μ EDTA 0.5 M) from unrelated patients with G6PD deficiency were collected from the hospitals of the north-west of Iran (including Ardebil, Tabriz and Urmia provinces). The blood samples collection process

was performed with the ethics committee approval and the patients' informed consent through telephone correspondences. and until the DNA extraction, the falcons were held in the temperature of -20°C. Qualitative measurement of the enzyme activity was performed using Fluorescent Spot Test and Saba laboratory kit. The basis of this method is the catalytic activity of G6PD enzyme in conversion of G6PD Glucose 6-phosphate to 6-Phosphogluconate and simultaneous revival of NADP to NADPH₂. The produced NADPH₂ has fluorescent specifications under UV (365nm). The fluorescence intensity in the blood of healthy individuals is positive (strong) and in the blood of patients with G6PD deficiency is low or negative. Extraction and purification of the samples' DNA Genomic, from Peripheral blood leukocytes, was performed by Rapid Genomic DNA Extraction (RGDE) method.

RGDE method was performed as follow (Saremi *et al.*, 2010):

Pouring 500 μl or 0.5 gr of blood sample into a 1.5 ml microfuge tube and 1000 μl of Cell Lysis buffer.

Shaking microfuge tube gently, and then centrifuging it for 2 minutes at 6000 rpm.

Removing and discarding supernatant and repeating steps 1-3 two or three.

Adding 300 μl of Nuclei Lysis buffer to the microfuge tube and keeping the tube in room temperature for 2 minutes to prevent clot formation.

Adding 100 μl of saturated NaCl and 600 μl of Chloroform to the microfuge tube, shaking it gently then centrifuging it for 2 minutes at 6000 rpm.

Transferring 300-400 μl of supernatant to a new 1.5ml microfuge tube.

Adding 600 μl of cold Isopropanol to it; shaking it gently then quickly.

Centrifuging the microfuge tube for one minute at 13000 rpm to precipitate, and then removing some of supernatant and letting completely dried in

room temperature and keeping it for later uses.

Amplification of exon 6 from the G6PD gene was performed by primers F-Med [5'CCC CGA AGA GGA ATT CAA GGG GGT 3'] and R-Med [5'GAA GAG TAG CCC TCG AGG GTG ACT 3'] 92 in thermal cycler (Senso Quest-Germany). For each subject patient, 3 μl of the mentioned primers, DFS-Taqmaster mix 12/5 μl (Bioron-Germany) was added to the PCR tube. Then 2 μl of the DNA was added to the tube and until the tube's final volume 25 μl distilled water (dH₂O) was added to it. PCR conditions were as follows: Initial denaturation of DNA at 95 °C for 5 min. 35 cycles each consisting of three steps of denaturation at 94 °C for 1 min, annealing at 59 °C for 1 min, extension at 72 °C for 1 minute and final extension at 72°C for 5 minutes. PCR Products were electrophoresed on 1.5% agarose gel at presence of DNA Ladder 50-700 bp (Bioron-Germany) and negative control samples. (fig. 1) Then 583bp fragment, the amplified product, was digested by enzymes *MboII*.

For digestion of 10 μl PCR products, 2 μl of 10X Enzyme buffer (B⁺) and 1 μl *MboII* Enzyme (Fermentas Co.) were mixed in a tube and until the tube's final volume, 31 μl dH₂O was added to it. The Mixture was transferred to the Thermal Cycler device and for each of the cycle, the temperature of 65°C for 2 hours and the temperature of 37°C for 25 minutes were adjusted and digested products were electrophoresed on 1.5-2% agarose gel at presence of DNA Ladder and negative control samples. In case of the absence of Mediterranean mutation and the enzymatic digestion, 379 bp, 120 bp, 60 bp, 24 bp bands are visible. If there is a mutation, 379bp band is cut and turns to 276bp and 103 bp. (fig. 2). To study the G6PD genetic changes (exon 6) accurately, DNA Sequencing was performed in a randomly selected sample of Mediterranean Mutant. To do so the PCR product was sent to Fazapazhouh Company (USA) and the response was analyzed by means of Chromas Lite version 2.33 software (fig. 3).

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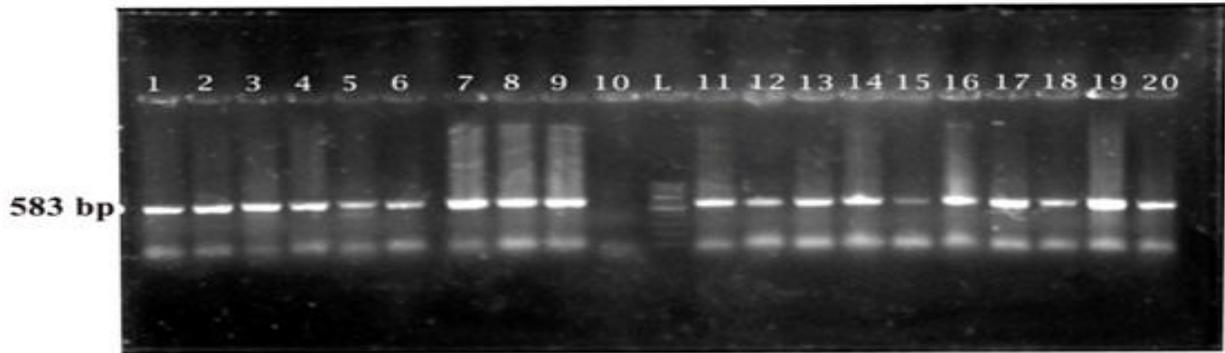


Fig. 1: The results of amplification of nucleotide sequence containing Mediterranean mutation. Lanes 1-20: PCR product. Lane 10: negative control. L: 50 bp plus DNA Ladder.

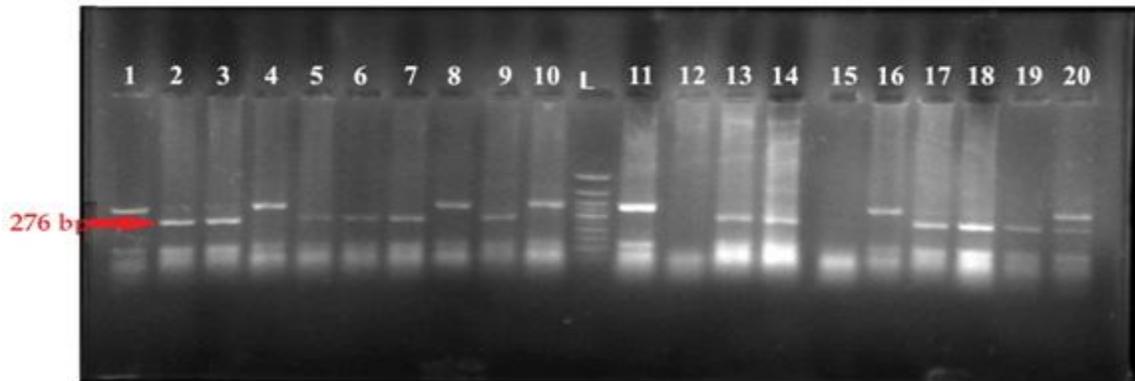
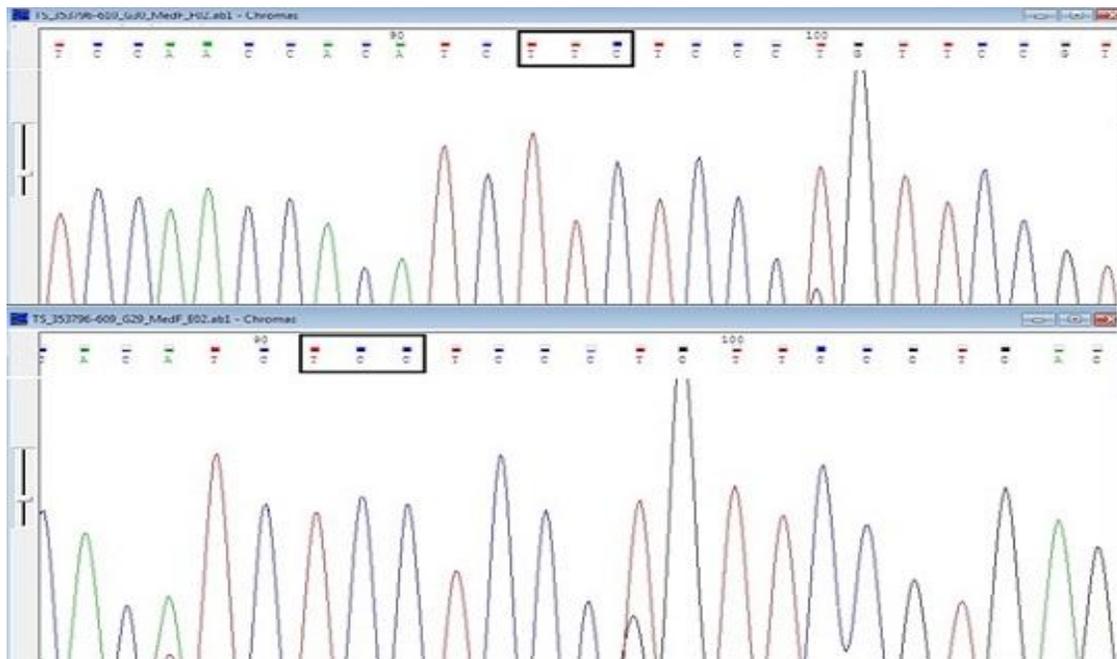


Fig. 2: Restriction digestion analysis of PCR products related to G6PD Mediterranean mutation with MboII. Lanes: 1, 4, 8, 10, 11, 16: normal samples. Lane 2: G6PD-Med Positive control. Lanes: 3, 5, 6, 7, 9, 13, 14, 17, 18, 19: G6PD Mediterranean mutation. Lanes: 12, 15: negative control. Lane 20: A heterozygote sample.



The results of DNA Sequencing for a sample of patient with 563 mutation (C-T) (top) and healthy individual (bottom).

As it was shown in the rectangular, in the patient sample C base has been replaced with T base.

(TCC 188 TTC) → (Ser 188 Phe)

Results

This study was conducted on 77 individuals with deficient G6PD enzyme activity. Individuals' ages ranged from one month to 21 years. G6PD Mediterranean mutation (563 C-T) was observed in 48 (37 males and 9 females) cases, which represents the prevalence rate of 62/3%. 29 patients (37/7%) weren't affected by the Mediterranean-type mutation. Mediterranean defect prevalence rate was 77/1% in men and 22/9% in women and it shows the high incidence of this disease in males and X-Linked recessive pattern of inheritance. Among women, 9 cases (81/8%) were heterozygote females and 3 cases (18/2%) were homozygote.

Discussion

Using RFLP-PCR, this study showed that 62/3% of individuals with Mediterranean type enzyme deficiency has mutations in nucleotide 563

(C-T) of G6PD gene. In these individuals, at position 188, phenylalanine amino acid is replaced by serine. G6PD is one of the most common genetic diseases that many people in the world suffer from it. This disease has caused many damages on child development and it has taken into consideration by the World Health Organization. On the basis of different studies, Mediterranean mutation prevalence is different around the world and Iran. G6PD Mediterranean mutation prevalence is 79% in turkey (Keskin *et al.*, 2002), 80% in Saudi Arabia (Niazi *et al.*, 1996), 74% in Oman (Darr *et al.*, 1996), 55% in UAE (Bayoumi *et al.*, 1996), and 76% in Pakistan (Saha *et al.*, 1994). Noori-Dalooi *et al.*, studies conducted in northern, central and southern provinces of Iran (table 1) (Noori-Dalooi *et al.*, 207, 2006b), showed that the lowest and the highest prevalence of Mediterranean mutation in patients with G6PD deficiency belongs to Kerman 63% and Gilan 86.5% provinces, respectively.

Table 1: Prevalence of Mediterranean mutation in north, center and south of Iran.

Northern provinces	Area's group	
	Central provinces	Southern provinces
Gilan 86.4 %	Yazd 64 %	Sistan and Baluchestan 80%
Mazandaran 66.2 %	Kerman 63%	Hormozgan 79%
Golestan 69 %		
Khorasan 66%		

The Studies of Karimi *et al.*, in Fars Province (Karimni t al., 2003), Miri Moghaddam *et al.*, in Sistan and Baluchistan province (Mirimoghaddam *et al.*, 2002) and Mortazavi *et al.*, in Zanzan province (Mortazavi *et al.*, 2006) have reported 83.8%, 85.1% and 72.7%, respectively, prevalence in these provinces. the present study, prevalence rate of the mutation is 62.3% which, compared to the reported mutations, represents the lowest prevalence rate. This study and the previous studies, conducted in the provinces, reveal that Mediterranean mutation type is the dominant mutation in these areas it means that in Iran, G6PD dominant mutation is Mediterranean type. That this mutation has happened thousands of years ago and the mutant

genes have been transferred to other territories through migration or population displacement due to war and etc.

Several studies have shown that patients with G6PD deficiency are resistant against catching malaria (Guido *et al.*, 2007). Because the prevalence rate of Malaria in the northwest of the country is lower than the South and Southeast regions, so the prevalence of deficiency for this region is justifiable.

Although, due to the inheritance of X-linked recessive, it is expected that this deficiency should be observed only in males but based on the Lyon's hypothesis (random inactivation of sex chromosomes), although less prevalent, it is likely

to occur in females. In the present study, like other studies, a significant percentage of females (23%) had this deficiency.

29 patients (37.7%) of the subjects had no mutations. since more than 60 different mutations have been identified at the molecular level (Mortazavi *et al.*, 2006). So identification of mutations in these patients needs further molecular studies and it can be done through using SSCP and ARMS-PCR methods and sequencing the G6PD gene.

Acknowledgement

I would like to appreciate the respected authorities of the Hematology Laboratory of Koodakan-E-Tabriz (children hospital of Tabriz), Bu Ali Ardabil and shahid Motahari hospitals to give samples and also I would like to appreciate authorities of the Islamic Azad University of Marand to allow for the use of equipment and facilities of Genetics Laboratory.

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