

Original Research Article

Allelic frequency of PON1 Q192R, CYP2C19*2 and CYP2C19*17 among Jordanian patients taking clopidogrel

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Abstract

Purpose: To investigate the influence of allelic frequencies of PON1 Q192R, CYP2C19*2 and CYP2C19*17 genetic polymorphisms on the response to clopidogrel among Jordanian patients.

Methods: Polymorphisms in CYP2C19 were assessed among 148 patients using PCR-RFLP assay.

Results: The CYP2C19*2, CYP2C19*17, and PON1 Q192R allele frequencies were 9.8, 28.72 and 28.7 %, respectively. On the genotyping side, the frequencies of CYP2C19*1/1* and CYP2C19*1/2* were 80.4 and 19.6 %, respectively, but none of the patients had CYP2C19*2/2* genotype. The genotype frequencies CYP2C19*17 were 47.97, 46.62 and 5.41 % for wild-type C-C, heterozygote C-T, and the mutant T-T, respectively. PON1 genotype was 42.7 % for QQ, and 57.8 % for QR. None of the patients had RR genotype.

Conclusion: Relative to other populations, the observed allelic frequencies are consistent with the values reported for Caucasian and Middle Eastern populations.

Keywords: CYP2C9 polymorphisms, Clopidogrel, Genotype, Allele frequency, PON1 genes

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INTRODUCTION

Cytochrome 2C19 (CYP2C19) is responsible for the metabolism of many drugs such as s-mephenytoin, proton pump inhibitors (PPIs), clopidogrel and proguanil [1]. Genetic variants play a crucial role in inter-individual variations in response to drugs [1]. Recent studies have shown that CYP2C19 polymorphisms lead to differences in response to clopidogrel. The CYP2C19 gene is highly polymorphic in humans [2]. In the literature, more than 30

genetic variants of CYP2C19 have been described. The most frequent SNPs are CYP2C19*2 and CYP2C19*3 which cause complete absence of enzymatic activity *in vitro* and *in vivo* [3].

Another common alteration that results in multiple active alleles leading to increased activity of the enzyme is CYP2C19*17 [3]. Consequently, humans can be divided, according to the ability to metabolize CYP2C19 substrates, into poor,

intermediate, extensive, and ultra-rapid metabolizers. Poor metabolizers (PMs) have two non-functioning alleles, whereas the intermediate metabolizers (IMs) have one functioning allele. Extensive metabolizers (EMs) carry two functioning alleles, while the ultra-rapid metabolizers are characterized by having a gain-in-function variant (*CYP2C19*17*) that enhances *CYP2C19*'s activity [4].

Many studies have reported that PM patients who have *CYP2C19*2* loss-of-function allele are associated with marked decrease in platelet response to clopidogrel, and hence increased cardiovascular events [4]. On the other hand, patients carrying the *CYP2C19*17* allele with UM phenotype may have higher benefit from clopidogrel treatment after acute myocardial infarction [5]. The *CYP2C19*2* SNP has been reported to account for approximately 12 % of variations in responses to clopidogrel [5]. This has indeed attracted researchers to look for other genetic factors that may account for the remaining variabilities in response to clopidogrel. Subsequently, it was revealed that the enzyme activity of paraoxonase-1 (PON1) has a strong link with clopidogrel bio-activation [6]. Moreover, PON1 is the first enzyme that catalyzes the second stage in clopidogrel bio-activation process [6]. These observations are consistent with the observed decreases in plasma paraoxonase activity, the maximum concentrations of active clopidogrel metabolites, and the increases in percentage of platelet inhibition in patients with QQ genotype.

The present study was carried out to determine the allelic frequency of *PON1* Q192R, *CYP2C19*2*, and *CYP2C19*17* among Jordanian patients on clopidogrel therapy. The findings might help in predicting the frequency of patients at risk of inadequate treatment with clopidogrel and those who may need dose adjustments.

EXPERIMENTAL

Sample collection

Whole venous blood (3 – 5 mL) was collected in EDTA tubes from 148 unrelated Jordanian patients who were taking clopidogrel (55 females and 95 males). The ethical committee of the University of Jordan gave approval for this study (IRB no. 2011-35-2) which complied to the guidelines of Declaration of Helsinki [7]. All patients signed informed consent before collecting blood samples.

DNA extraction

The extraction of DNA was done using Wizard DNA extraction kit (Promega, Madison, WI, USA) according to the manufacturer's instructions.

Genotyping method

The PCR-RFLP method was carried out to determine the genotype of *CYP2C19*2*, **17* and *PON1* as described previously [8-10], with minor modifications. In essence, the 50- μ L PCR reaction contained 200 ng DNA, 10 μ L Taq polymerase buffer, 0.2 mM dNTPs, 1.0 μ L MgCl₂, 10 picomole of the forward and reverse primers (Table 1), and 1 U of Taq polymerase. The reaction mixtures were incubated in PCR tubes at 94 °C for 8 min, followed by 30 cycles at 94 °C for 30 sec, and then at 63, 56 or 52 °C for 30 sec (*CYP2C19*17* **2*, **17* and *PON1* polymorphisms, respectively).

Restriction enzyme analysis

The PCR products for *CYP2C19*2*, **17* and *PON1* were digested with appropriate restriction enzymes (*Sma*I, *Alw*I and *Nsi*I, respectively), and the DNA fragments were separated onto 3.0 % agarose gel stained with ethidium bromide.

Statistical analysis

The *CYP2C19* allele frequencies were tested for Hardy–Weinberg equilibrium using Chi-square (χ^2) test. Levels were deemed significant at *p* values less than 0.05.

RESULTS

Table 2 shows the allele and genotype frequencies of *CYP2C19*2*, **17* and *PON1* Q192R among 148 unrelated Jordanian patients on clopidogrel therapy. The frequencies of *CYP2C19*2*, **17* and *PON1* Q192R variant alleles were 9.8, 28.72 and 28.7 %, respectively. For genotype frequency, *CYP2C19*2* scored 80.4 % for the **1/*1* wild-type, 16.6 % for the heterozygote **1/*2* and 0.0 % for the mutant **2/*2* genotype. The frequencies of *CYP2C19*17* genotype were 47.97, 46.62 and 5.41 % for wild-type C-C, heterozygote C-T, and mutant T-T, respectively.

For *PON1* Q192R, the genotype frequencies were 42.7, 57.3, 0.0 % for wild-type Q-Q, heterozygote Q-R, and mutant R-R, respectively. The homozygous mutant allele *CYP2C19*2/*2* and *PON1* R192R were not detected in any of 148 patients (Table 2).

Depending on the classification of CYP2C19 enzymatic activity [4], the subjects in this study were placed in 4 metabolic groups namely: poor, intermediate, extensive and ultra-rapid metabolizers (Table 3).

When compared with other populations, the observed frequencies were consistent with the values reported in Turkish, Italian, African-American, Saudi Arabian, German, and Danish populations, but they were profoundly different from the allelic frequencies in Chinese, Koreans, and Japanese populations (Table 4).

DISCUSSION

Clopidogrel is extensively used by Jordanians. Thus, it is important to determine the allelic frequency of the SNPs that may

play a role in the effectiveness of this pro-drug. This study was aimed at determination of the allelic frequency of *CYP2C19*2*, *CYP2C19*17*, and *PON1 Q192R* SNPs in Jordanian patients on clopidogrel therapy. The three SNPs have been shown to play major roles in the conversion of clopidogrel to its active form. Moreover, the allelic frequency of *CYP2C19*2* and *CYP2C19*17* may be important in patients taking other drugs that are metabolized by *CYP2C19* such as anti-ulcer medications, antidepressants, and β -adrenoceptor blockers [19].

Several studies have been carried out on *CYP2C19*2* and *CYP2C19*17* because they are the most common alleles that result in reduced or increased *CYP2C19* activity [19].

Table 1: Primer sequences used for amplification of *CYP2C19*2*, *CYP2C19*17*, and *PON1*

Primer name	Sequence (5'-3')
CYP2C19*2 -F	CAACCAGAGCTTGGCATATTG
CYP2C19*2 -R	CACAAATACGCAAGCAGTCAC
CYP2C19*17-F (PCR1)	GCCCTTAGCACCAAATTCTC
CYP2C19*17-R (PCR1)	ATTTAACCCCTAAAAAACACG
CYP2C19*17-F (PCR2)	AAATTTGTGTCTTCTGTTCTCAATG
CYP2C19*17-R (PCR2)	AGACCCTGGGAGAACAGGAC
PON1-F	TATTGTTGCTGTGGGACCTGAG
PON1-R	CCTGAGAATCTGAGTAAATCCACT

F: Forward, R: Reverse

Table 2: *CYP2C19* and *PON1* genotype and allele frequencies in a sample of Jordanian patients on clopidogrel therapy

Alleles	Number (n=148)	Genotype frequency (%)	Allele frequency (%) n=296	95 % CI
<u><i>CYP2C19*2</i></u>				
1-1	119	80.4		0.0501-0.1459
1-2	29	16.6	9.8(2)	
2-2	0	0		
<u><i>CYP2C19*17</i></u>				
C-C	71	47.97		0.2143-0.3601
C-T	69	46.62	28.72 (T)	
T-T	8	5.41		
<u><i>PON1</i></u>				
Q-Q	64	42.7		
Q-R	84	57.3	28.7 (R)	0.2141-0.3599
R-R	0	0		

Table 3: Predicted CYP2C19 phenotype based on *CYP2C19*2* and *CYP2C19*17* genotypes in Jordanian patients on clopidogrel therapy

Predicted phenotype	<i>CYP2C19</i> genotype	Number of individual (%)	95 % confidence interval
EM	<i>CYP2C19*1/*1</i> & <i>CYP2C19*2/*17</i>	64 (43.24 %)	0.3526-0.5122
Het EM/IM	<i>CYP2C19*1/*2</i>	18 (12.16 %)	0.0689-0.1743
PM	<i>CYP2C19*2/*2</i>	0 (0.00 %)
Het UM	<i>CYP2C19*1/*17</i>	58 (39.19 %)	0.3132-0.4706
UM	<i>CYP2C19*17/*17</i>	8 (5.41 %)	0.0177-0.0905

Table 4: *CYP2C19* and *PON1* alleles in Jordanians compared with other ethnic groups

Population	<i>CYP2C19*2</i> (Reference)	<i>CYP2C19*17</i> (Reference)	<i>PON1 Q192R</i> (Reference)
Chinese	24.9 [11]	1.2 [11]	64.8 [11]
Turkish	12.0 [12]	37.8 [12]
Italian	11.9 [13]	28.9 [13]
American-African	18.2 [14]	21.0 [14]	32.0 [14]
Saudi Arabians	11.2 [15]	25.7 [15]
Japanese	27.9 [16]	1.3 [16]	33.4 [16]
German	15.2 [17]	25.5 [17]	27.7 [17]
Danish	15.0 [18]	20.1 [18]
Jordanians (our study)	9.8	28.72	28.7

Inter-ethnic variations in *CYP2C19* polymorphism have been reported [20]. Among Jordanians, the allelic frequency of *CYP2C19*2* is within the range of 12.5 to 16 % [21,22]. These findings are not significantly different from the 9.8 % allelic frequency seen in the present study. Relative to other populations, the observed frequencies are in accordance with the values reported in Turkish, Italian, African-American, Saudi Arabian, German, and Danish populations. However, they are profoundly different from the allelic frequencies in Chinese, Korean, and Japanese populations. Interestingly, the allelic frequencies in different ethnic groups were relatively high. This may imply that this detrimental mutation is relatively old and occurred before the Black, Oriental, and Caucasian racial groups split [20].

At the phenotyping level, 29 out of the 148 patients (19.6 %) were found to carry one non-functional allele, and were classified as intermediate metabolizers, with intermediate metabolizer frequency of 12.16 %. These are in line with earlier report among Jordanians, where the frequency of intermediate metabolizers was in the range of 18 – 19.2 % [21, 22]. None of the subjects in this study had a poor metabolizer phenotype, in contrast to what was reported previously [21,22]. A possible explanation for this variation might be the differences in the methodologies used in these studies. Future studies with a larger sample should be conducted to investigate further the percentage of poor metabolizers among Jordanians.

Since the percentage of intermediate metabolizers is not negligible in Jordanians,

some patients receiving clopidogrel (75 mg dose) may be at higher risk for stent thrombosis. This conclusion is supported by other studies carried out on different populations which indicated that *CYP2C19*2* carriers exhibited higher platelet index in a multivariate analysis, and had significantly higher cumulative 30-day incidence of stent thrombosis, when compared with wild-type *CYP2C19* carriers [19,23].

Regarding *CYP2C19*17*, this study is the first reported attempt to determine *CYP2C19*17* allele frequency in Jordanians. The allelic frequency falls within those of the African Americans and Saudi Arabians, and is profoundly higher than that of the Chinese. The *CYP2C19*17* allele and genotype frequencies reported in this study indicate that a substantial percentage of Jordanians have increased activity of the *CYP2C19* enzyme. The functional and clinical implications of *CYP2C19*17* on various *CYP2C19* substrates (omeprazole, pantoprazole, escitalopram, sertraline, voriconazole, tamoxifen, and clopidogrel) have been reported [24].

The functional effects of *CYP2C19*17* are unlikely to be clinically significant except for drugs with very narrow therapeutic windows. Out of the drugs reviewed, only clopidogrel and tamoxifen may be of concern. For clopidogrel, only *CYP2C19*17* homozygotes are likely to be at significantly increased risk of experiencing the effects of excessive inhibition of platelet aggregation. More recently, it was reported that *CYP2C19*2* and **17* alleles have significant influences on the response to clopidogrel response [25], which indicates that *CYP2C19*17* SNP is of interest in clopidogrel users as it may

increase the risk of bleeding especially for the 5.41 % that carry the homozygous activating alleles.

On *PON1* Q192R, the observed allelic frequency is consistent with the Caucasian populations including German, Italian, African-American and Japanese populations. However, its allelic frequency is significantly different from that of the Chinese. These findings indicate that there is a high degree of inter-ethnic variations in *PON1*Q192R, and like *CYP2C19**2, this SNP is relatively old and probably occurred before the racial groups were split.

With respect to *PON1* Q192R phenotype and clopidogrel metabolism, none of the study subjects was an extensive metabolizer. However, 86 patients had intermediate metabolizer phenotype. Moreover, an appreciable number of subjects (12.2 %) combined both *CYP2C19**2 and *PON1*Q192 SNPs. It is speculated that this may put them at higher risk for thrombosis than those with only one non-functioning allele of either *CYP2C19* or *PON1*.

Although clopidogrel is activated mainly by *PON1* enzyme [6], subsequent studies reported contradictory outcomes by excluding any significant role for *PON1* enzyme in the activation of clopidogrel [8]. Accordingly, a prospective study should be conducted to clarify the involvement of *PON1* in the activation of clopidogrel. Considering the fact that the variation in *CYP2C19**2 frequencies accounted for only 12 % of the observed variability in clopidogrel response, these studies are direly needed [2].

CONCLUSION

Compared with other populations, the observed *CYP2C19**2, *17 and *PON1* allelic frequencies are in accordance with the values reported for Caucasian and Middle Eastern populations. Further intensive studies are needed to unravel the influence of *CYP2C19**2, *17 and *PON1* polymorphisms on clinical clopidogrel response.

DECLARATIONS

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Conflict of interest

The authors declare that no conflict of interest is associated with this work.

Contribution of authors

We declare that this work was done by the authors named in this article and all liabilities pertaining to claims relating to the content of this article will be borne by the authors. Malek Rjoub did most of the laboratory work, with some assistance from Amer Imraish. He also wrote manuscript. Akram Saleh collected the blood samples from University of Jordan Hospital. Nancy Hakooz. Malek Zihlif supervised the work and reviewed the manuscript. Yazun Jarrar did the technical review.

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