Polymorphisms of VKORC1 and CYP2C9 are associated with warfarin sensitivity in Chinese population

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Objective: Warfarin is a commonly prescribed anticoagulant for prevention of thromboembolic events. Wide inter-individual dose variation, narrow therapeutic range and risk of serious bleeding result in difficulties in achieving the therapeutic effect. The present study was designed to clarify the real biological significance of the polymorphisms of VKORC1 and cytochrome P450 2C9 (CYP2C9) in warfarin metabolism.

Methods: A total of 214 patients with warfarin anticoagulant therapy were selected. During follow-up of anticoagulation, warfarin dosage and associated international normalized ratio values were recorded. Genetic polymorphisms of VKORC1 promoter and from exon 1 to exon 3 and CYP2C9 exon 4 sequence were detected by polymerase chain reaction and gene sequencing.

Results: Five polymorphisms were identified in this research, which were VKORC1 1173C>T (intron 1), 1542G>C (intron 2), 2255C>T (intron 2), 3730C>T (3’-downstream) and CYP2C9 exon 4 -65G>C. VKORC1 1173CT, 1542GC, 2255CT and 3730CT polymorphisms were detected in same patients, but CYP2C9 exon 4 -65GC carriers were different from them. VKORC1 1173CT, 1542GC, 2255CT, 3730CT carriers and CYP2C9 exon 4 -65GC carriers had significantly higher warfarin daily dosage than others (3.2±0.6 vs 3.1±1.1 vs 2.6±0.8 mg/day). Logistic regression analysis revealed VKORC1 1173CT, 1542GC, 2255CT, 3730CT carrier status (odds ratio [OR] =3.233, 95% confidence interval [CI]: 1.259–8.303, P=0.015) and obesity with body mass index >27 kg/m² (OR =1.223, 95% CI: 1.097–1.363, P<0.001) to have independent and statistically significant contributions to high warfarin dosage.

Conclusion: In general, in VKORC1 1173CT, 1542GC, 2255CT and 3730CT carriers and in obese patients, warfarin maintenance doses were significantly higher than in the others.

Keywords: warfarin, vitamin K, cytochrome P450, polymorphism

Introduction

Warfarin is the most commonly used oral anticoagulant drug in the prevention of thromboembolism. For patients with prosthetic heart valve replacement, the incidence of hemorrhage is inversely connected with the efficacy of anticoagulation and appears to be a major risk indicator for bleeding complications.¹² It has been reported that the Chinese populations are more sensitive to warfarin than western populations.¹³ To maintain the international normalized ratio (INR) of 2.0–3.0, the mean daily warfarin dosage is lower in Chinese than in Caucasians. This cannot be explained simply by a lower mean body weight among the Chinese.³

The vitamin K epoxide reductase complex 1 (VKORC1) recycles vitamin K 2, 3-epoxide back to active vitamin K hydroquinone, an important cofactor involved
in the activation of vitamin K-dependent clotting factors. VKORC1 is the target enzyme of inhibition of warfarin,\textsuperscript{4,5} and variants of the VKORC1 gene are known to be associated with warfarin resistance. Four different exonic mutations have been described in individuals with known warfarin resistance phenotype.\textsuperscript{6} It was notable that these mutations clustered in the luminal loops of the gene and were predicted to have functional effects.\textsuperscript{7} More recently, VKORC1 promoter polymorphisms that occur frequently have been discovered. They exhibit linkage disequilibrium, and are associated with higher warfarin dose requirements.\textsuperscript{8–10}

Cytochrome P450 2C9 (CYP2C9) is polymorphic in human and is principally responsible for the metabolism of warfarin. Six distinct single-base pair substitution polymorphisms of the CYP2C9 gene have been discovered and are designated *1 through *6.\textsuperscript{11,12} However, known CYP2C9 polymorphisms cannot entirely account for the low-dose requirement of warfarin seen in Chinese patients with mitral valve replacement.

Collectively, genetic polymorphisms are involved in both pharmacodynamic (VKORC1) and pharmacokinetic (CYP2C9) factors. They appear to interplay in the overall inter-individual variability of warfarin doses. In this context, we screened the promoter and the whole exons (exon 1 to 3) of VKORC1 and CYP2C9 exon 4 polymorphisms and investigated their roles in warfarin sensitivity in Chinese population.

Patients and methods
A total of 214 patients with warfarin anticoagulant therapy were selected. Average age was 62.1±10.5 years. They were receiving maintenance warfarin therapy with a stable, therapeutic INR between 2 and 2.5 for at least 3 weeks. Mean daily warfarin requirement was from 1.250 to 5.077 mg/day when therapeutic INR (2.0–2.5) was 24.7±3.8 kg/m\textsuperscript{2}. Standard clinical laboratory tests indicated that all of the patients had normal liver function and renal function. Patients with concurrent medications with potential to affect warfarin’s metabolism included amiodarone, non-steroidal anti-inflammatory drugs, cimetidine, thyroid hormone, or carbamazepine were excluded. The study was approved by the Ethics Committee of Tianjin Medical University General Hospital and Shenzhen Sun Yat-sen Cardiovascular Hospital. Written informed consent was obtained from each patient.

Study protocol
Blood (5–10 mL) was obtained from 12 to 16 h after administration of the last dose of warfarin during a routine clinic visit. The average daily dose of warfarin was calculated from 1-week period, and latest INR was recorded.

Genotyping protocol
DNA was extracted with Gentra DNA Isolation Kit (Hilden, Germany). Sequence amplification was performed by polymerase chain reaction. Gene sequencing of VKORC1 promoter and exon 1 to exon 3 and CYP2C9 exon 4 was performed following methodology described previously.\textsuperscript{8}

Statistical analysis
Statistical analysis was performed using Statistical Package for the Social Sciences (SPSS) for Windows (version 15.0). Continuous data are presented as mean ± standard deviation and analyzed using independent-samples t-test, repeated measures analysis of variance (ANOVA), or one-way ANOVA, as appropriate. Categorical variables were expressed as frequencies and analyzed with \( \chi^2 \) test. Logistic regression was used to analyze the risk factors of outcome. P-values <0.05 were considered statistically significant.

Results
Average body mass index (BMI) of the 214 participants was 24.7±3.8 kg/m\textsuperscript{2}. Mean daily warfarin requirement was from 1.250 to 5.077 mg/day when therapeutic INR (2.0–2.5) levels were maintained. Patient demographics are presented in Table 1.

Five polymorphisms were identified in this study. They were VKORC1 1173C>T, 1542G>C, 2255C>T, 3730C>T and CYP2C9 exon 4 –65G>C. Genotype distributions of all polymorphisms were in Hardy-Weinberg equilibrium. These genotype frequencies are shown in Table 2. VKORC1 1173CT, 1542GC, 2255CT and 3730CT carrier status was in the same patients, and CYP2C9 exon 4 –65GC carriers

<table>
<thead>
<tr>
<th>Table 1 Patient characteristics</th>
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<tbody>
<tr>
<td>Parameters</td>
</tr>
<tr>
<td>Sex (M/F)</td>
</tr>
<tr>
<td>Age (years)</td>
</tr>
<tr>
<td>BMI (kg/m\textsuperscript{2})</td>
</tr>
<tr>
<td>Smoking (n, %)</td>
</tr>
<tr>
<td>Alcohol drinking (n, %)</td>
</tr>
<tr>
<td>Diabetes mellitus (n, %)</td>
</tr>
<tr>
<td>Kinds of disease</td>
</tr>
<tr>
<td>Atrial fibrillation (n, %)</td>
</tr>
<tr>
<td>AVR (n, %)</td>
</tr>
<tr>
<td>MVR (n, %)</td>
</tr>
<tr>
<td>AVR + MVR (n, %)</td>
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<tr>
<td>MVR + CABG (n, %)</td>
</tr>
</tbody>
</table>

Note: Data are presented as either mean ± SD or counts.

Abbreviations: M, male; F, female; BMI, body mass index; AVR, aortic valve replacement; MVR, mitral valve replacement; CABG, coronary artery bypass grafting; SD, standard deviation.
were absolutely different from them. The clinical characteristics of the VKORC1 1173CT, 1542GC, 2255CT, 3730CT carriers, CYP2C9 exon 4 –65GC carriers and the others are exhibited in Table 3. These patient groups did not differ significantly in relation to other parameters. However, VKORC1 1173CT, 1542GC, 2255CT, 3730CT carriers and CYP2C9 exon 4 –65GC carriers had significantly higher daily warfarin dosage than the others (3.2±0.6 vs 3.1±1.1 vs 2.6±0.8 mg/day) (Figure 1).

Logistic regression analysis included age, sex, BMI and carrier statuses as covariates and showed that VKORC1 1173CT, 1542GC, 2255CT, 3730CT carriers (odds ratio [OR] =3.233, 95% confidence interval [CI]: 1.259–8.303, \( P=0.015 \)) and obese individuals (with BMI >27 kg/m²) (OR =1.223, 95% CI: 1.097–1.363, \( P<0.001 \)) had independent and statistically significant contributions to high warfarin dosage (Table 4).

**Discussion**

Warfarin’s anticoagulant activity results from inhibition of hepatic vitamin K epoxide reductase that affects the synthesis of various coagulation factors. Recently, variants of the VKORC1 have been described to have potentially functional consequences.\(^\text{13-15}\) However, these polymorphic variants are not useful for testing in Chinese population because of their low prevalence. In the present research, all exons of VKORC1 (exon 1–exon 3) and the promoter were examined by gene sequencing. Four polymorphisms were identified, which were 1173C>T, 1542G>C, 2255C>T, 3730C>T. The results indicated that polymorphisms are found in absolute linkage disequilibrium since 1173CT, 1542GC, 2255CT, 3730CT are carried by same patients. VKORC1 1173CT could be used as the tagging SNP to represent the others for further research. Logistic analysis showed that 1173CT, 1542GC, 2255CT, 3730CT were one of the independent contributions of high warfarin dosage. This is consistent with previously published report, which indicates that 1173CT was associated with lower VKORC1 mRNA levels in human liver.\(^\text{16}\) This finding suggests that 1173CT may be associated with the lower levels of reduced form of vitamin K, thereby making patients with this variant more susceptible to the anticoagulation effect of warfarin.

Importantly, VKORC1 1173C>T genotypes are significantly different among Asians, Caucasians and African Americans. The frequencies of VKORC1 1173C allele in Asians, Caucasians and African Americans were 11%, 58% and 91%, respectively.\(^\text{17-19}\) In the present result, the frequency of 1173CT is approximately 14%. Furthermore, multiple regression analysis showed that the VKORC1 1173C>T variant was an important covariate with respect to the inter-individual variability in warfarin dosage. Patients carrying the C allele at the position of 1173 of the VKORC1 gene are associated with a significantly higher dose requirement of warfarin. So, these results suggest that the lower dose requirements in Chinese population may possibly reflect the lower frequency of the VKORC1 1173C allele compared to western population.

**Table 2** Genotype distributions

<table>
<thead>
<tr>
<th>Polymorphisms</th>
<th>Genotype</th>
<th>Allele</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>VKORC1</td>
<td>TT(184)</td>
<td>CT(30)</td>
<td>C(30)</td>
</tr>
<tr>
<td>1173C&gt;T</td>
<td>CC(184)</td>
<td>GC(30)</td>
<td>G(30)</td>
</tr>
<tr>
<td>VKORC1</td>
<td>TT(184)</td>
<td>CT(30)</td>
<td>T(398)</td>
</tr>
<tr>
<td>1542G&gt;C</td>
<td>CC(184)</td>
<td>GG(30)</td>
<td>G(30)</td>
</tr>
<tr>
<td>2255C&gt;T</td>
<td>TT(184)</td>
<td>CT(30)</td>
<td>T(398)</td>
</tr>
<tr>
<td>VKORC1</td>
<td>GG(184)</td>
<td>GA(30)</td>
<td>A(30)</td>
</tr>
<tr>
<td>3730C&gt;T</td>
<td>GG(184)</td>
<td>GA(30)</td>
<td>A(30)</td>
</tr>
<tr>
<td>CYP2C9 exon 4</td>
<td>GG(193)</td>
<td>GC(21)</td>
<td>C(21)</td>
</tr>
</tbody>
</table>

**Note:** Data in parentheses are numbers of each genotype.

**Table 3** Clinical characteristics in three groups of genotypes

<table>
<thead>
<tr>
<th>Parameters</th>
<th>VKORC1 1173CT, 1542GC, 2255CT, 3730CT carriers (n=30)</th>
<th>CYP2C9 exon 4 –65GC carriers (n=21)</th>
<th>Others (n=163)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sex (M/F)</td>
<td>16/14</td>
<td>7/14</td>
<td>88/75</td>
<td>0.344</td>
</tr>
<tr>
<td>Age (years)</td>
<td>56.4±8.1</td>
<td>56.2±16.7</td>
<td>59.5±10.5</td>
<td>0.294</td>
</tr>
<tr>
<td>BMI (kg/m²)</td>
<td>24.5±3.1</td>
<td>25.2±5.0</td>
<td>24.0±3.4</td>
<td>0.423</td>
</tr>
<tr>
<td>Smoking (n, %)</td>
<td>10 (33%)</td>
<td>5 (25%)</td>
<td>58 (36%)</td>
<td>0.523</td>
</tr>
<tr>
<td>Alcohol drinking (n, %)</td>
<td>4 (14%)</td>
<td>4 (17%)</td>
<td>21 (13%)</td>
<td>0.871</td>
</tr>
<tr>
<td>Diabetes mellitus (n, %)</td>
<td>4 (14%)</td>
<td>2 (8%)</td>
<td>19 (12%)</td>
<td>0.512</td>
</tr>
</tbody>
</table>

**Note:** Data are presented as either mean ± SD or counts.

**Abbreviations:** M, male; F, female; BMI, body mass index; SD, standard deviation.
investigation is necessary to investigate the clinical impact of genetic variance of CYP2C9 on warfarin management. In addition, warfarin patients usually visit the clinic every 3–6 months depending on their INR levels. We included only patients into the study if they could return to the clinic for INR testing and warfarin dose follow-up. This led to the relative small sample size of the study.

**Conclusion**

The warfarin maintenance doses of VKORC1 1173CT, 1542GC, 2255CT and 3730CT carriers and in obese patients were significantly higher than in the others. Our results indicate that the ethnic origin of the warfarin-treated population must be considered as an influencing factor affecting the dose response. In Chinese population, both obesity and the genetic polymorphism of genes affecting the warfarin metabolism seem to affect considerably the clinical response to warfarin.

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**Disclosure**

The authors report no conflicts of interest in this work.

**References**


