

Bedside stratification of patients according to *CYP2C19**2 genotype

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Objective. The task was to identify the defective *CYP2C19**2 allele by using a novel fast-genotyping method in the patients having coronary artery disease who developed stent thrombosis or high on-treatment platelet reactivity.

Materials and methods. A total of 35 patients were included in the study. All the patients were screened for *CYP2C19**2 using Spartan RX *CYP2C19* Assay Package.

Results. *CYP2C19**1*1 variant was found in 63%, *1*2 in 34%, *2*2 in 3% of patients. The distribution of genotypes was similar in men and women.

Conclusions. By using the bedside technique, *CYP2C19**2, which is responsible for a poorer effect of clopidogrel, might be identified in less than an hour.

Keywords: stent thrombosis, *CYP2C19*, clopidogrel, gender, antiplatelets, genotype

INTRODUCTION

A considerable amount of progress has been made in stent implantation techniques since the introduction of this method into clinical practice. Nowadays stent thrombosis remains a major life-threatening event following stent implantation. Optimal anti-platelet therapy is one of the most important factors in the prevention of stent thrombosis (Vieira et al., 2013). Current European guidelines recommend dual antiplatelet therapy with a combination of aspirin and a new antiplatelet agent (ticagrelor or prasugrel) as first-line therapy to the patients who

underwent percutaneous coronary intervention (PCI) and stent implantation (Hamm et al., 2011). Clopidogrel, which was shown to be inferior to ticagrelor or prasugrel, is still used worldwide more often than the new oral antiplatelet agents (Serebrany, Fortmann, 2014).

Clopidogrel, a P2Y₁₂ antagonist, has been used since 2002 for treating acute coronary syndromes (ACS) and in the patients undergoing PCI. Different variability in response to the treatment with clopidogrel is observed worldwide. Usually up to 30% of patients of European descent are identified as non-responders to clopidogrel antiplatelet therapy (Liu et al., 2011). A lack of a therapeutic clopidogrel effect is associated both with genetic and non-genetic factors. Various studies identified

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CYP2C19 G681A (or *2 allele) to be associated with a reduced clopidogrel therapeutic effect. Carriers of at least one *2 (reduced-function) allele have high on-treatment platelet reactivity and a significantly increased risk of major adverse cardiovascular events (Mega et al., 2010). On 12 March 2010, the Food and Drug Administration (USA) announced a black box warning about Plavix (clopidogrel), which highlighted the impact of ineffective clopidogrel therapy on the carriers of a reduced *CYP2C19* enzyme activity (FDA, 2010).

The *CYP2C19**17 allele (gain-of-function allele), on the contrary, is responsible for an increased activity of this enzyme. It results in exaggerated bio-activation of clopidogrel and is related to a higher bleeding risk, with average multi-ethnic allele frequencies accounting from 3% to 21% (Grosdidier et al., 2011).

Among the non-genetic factors, ethnicity, sex, age, body weight, comorbidities, drug–drug interactions, and other factors might also significantly affect the response to clopidogrel – antiplatelet therapy (Xie et al., 2011).

Newer-generation drugs prasugrel and ticagrelor have a more predictable therapeutic effect and are more potent than clopidogrel (Sabouretand Tael-Sartral, 2014). Retrospective genetic studies have shown that *CYP2C19* did not have any significant effect on the therapeutic effect of newer-generation drugs (Grosdidier et al., 2013). The application of new-generation drugs in clinical practice further remains complicated. Ticagrelor and prasugrel were shown to cause bleeding and other side effects more often than clopidogrel does. It was also shown that clopidogrel was effective enough in the carriers of wild-type (*1*1) *CYP2C19*, which represents a normal metabolism rate of this drug (Reese et al., 2012). In addition, only up to one-fourth of the prasugrel-treated patients are discharged from hospital for antiplatelet treatment at home (Sandhu et al., 2013; Bagai et al., 2014).

The available evidence suggests that personalised antiplatelet treatment (drug prescription according to the patients' genotype) is cost-effective. It can provide more clinical

value than conventional treatment by reducing the rates of thrombosis or bleeding complications (Li et al., 2014).

As a rule, genotyping in a clinical laboratory is a robust and time-consuming method. Therefore it is hard to apply this method in emergency situations. We used a novel bedside *CYP2C19*-genotyping equipment which allowed us to determine a patient's *CYP2C19* genotype in an hour. By using a novel bedside method, our main task was to determine the defective *CYP2C19**2 allele in the sample of the patients with coronary artery disease (CAD) who, following stent implantation, developed stent thrombosis or high on-treatment platelet reactivity, and who required accurate personalized drug therapy. We also wanted to compare the distribution of *CYP2C19**2 to a sample of healthy Lithuanian population.

MATERIALS AND METHODS

Study population

A total of 35 patients who were admitted to the hospital of the Lithuanian University of Health Sciences Kauno Klinikos at the Department of Cardiology from April 2013 to January 2014 due to CAD (coronary artery disease) and who underwent stent implantation and developed stent thrombosis 45.7% ($n = 16$) or high on-treatment platelet reactivity 54.3% ($n = 19$) were included into a further study. Platelet aggregation was measured by using the standard Born method (Born, 1962). Clinical and laboratory data were obtained from case histories. The research was approved by the local Bioethics Committee. All the participants signed informed consent.

Genotyping

All the patients were screened for the presence of *CYP2C19**2 by using Spartan RX™ *CYP2C19* Assay Package (Package Insert, 01001914_3.28; 08/2013, Spartan Bioscience Inc., Canada). The test was run on the Spartan RX™ platform which automatically extracts and genotypes the patients' DNA. The operation of the point-of-care genetic testing device consisted of the

following main separate steps intended to be made in less than 8 min:

1. acquiring a buccal swab;
2. inserting the swab into an assay cartridge;
3. pouring the reaction mix into a genetic testing device and analysing *CYP2C19**2.

The identification of *CYP2C19**2 carrier was made within 60 min. The results were as follows: homozygous for the wild-type *CYP2C19**1/*1, heterozygous *1/*2 or homozygous *2/*2.

Statistical analysis

Results are presented as mean \pm SD; categorical variables are presented in percentage. The Chi-square test was used for categorical variables and the unpaired Student t-test was used for continuous variables. The significance level was set at *p* value of 0.05.

RESULTS

The average age of the studied patients was 63.97 ± 11.67 years. About 2/3 (63%) of the patients studied were males. The patients with ST-elevation accounted for 37% of the sample. In total, 86% of the patients had systemic hypertension, more than a half of the patients studied had dyslipidaemia (57%), and 25% were patients with diabetes. Current smokers represented 29% of the sample. The baseline clinical characteristics of the patients are presented in the Table.

According to the *CYP2C19* genotype, a total of 22 patients had wild-type *CYP2C19**1*1 (63%), *1*2 was present in 12 patients (34%) and homozygous *2*2 was detected in one patient (3%).

According to the patients' sex, male patients ($n = 22$) were younger (61.49 ± 9.83 years) than female patients ($n = 13$) (69.30 ± 13.23 years), $p = 0.05$. *CYP2C19**1*2 was found in 36.4% ($n = 8$) of male patients. *CYP2C19**1*2 in female patients accounted for 31% ($n = 4$), one female carried *2*2 variant (2.9%).

The distribution of *CYP2C19* genotypes in a sample of Lithuanian healthy subjects was already shown in previous studies. The percentage of defective *1*2 and *2*2 variants in a sample of the healthy Lithuanian population

Table. **Baseline patient characteristics**

Men (%)	22 (62.9%)
Age (in years)	63.97 ± 11.67
Admission diagnosis	
STEMI	13 (37.1%)
NSTEMI	7 (20.0%)
UA	10 (28.6%)
SA/positive functional test	5 (14.3%)
Cardiovascular risk factors	
Diabetes mellitus	9 (25.7%)
Systemic hypertension	30 (85.7%)
Dyslipidaemia	20 (57.1%)
Current smoker	10 (28.6%)
Cardiovascular history	
Previous MI	17 (48.6%)
Previous CABG	4 (11.4%)
Angiographic characteristics	
One vessel treated	28 (80.0%)
Two vessels treated	6 (17.1%)
Drug-eluting stents	8 (22.9%)
Thrombotic lesion	16 (45.7%)
In-stent restenosis	10 (28.6%)

CABG – coronary artery bypass grafting; MI – myocardial infarction; NSTEMI – non-ST elevation myocardial infarction; PCI – percutaneous coronary intervention; STEMI – ST elevation myocardial infarction; UA – unstable angina; SA – stable angina.

($n = 279$) was lower (23.3% of intermediate *1*2 and 0.7% of poor metabolizer *2*2) ($p = 0.1$ for *1*2) as compared with the patient sample (Tatarūnas et al., 2012).

DISCUSSION

The *CYP2C19**2 allele (loss-of-function) is the most common type among the reduced-function genes and is associated with a reduced antiplatelet effect of clopidogrel and the increased risk for adverse cardiovascular events (Grosdidier et al., 2011). About 15% of the Caucasians, 29–35% of the Asians, and 25% of the African Americans have at least one *CYP2C19**2 allele (Rosemary, Adithan, 2007; Shetkar et al., 2014). Other *CYP2C19* variants

(*3, *4, *5, *6, *7, and *8 alleles) are also identified as having an impact on clopidogrel metabolism and therapeutic activity. Their frequency in the Caucasian population is relatively low (<1%) (Siller-Matula et al., 2013). According to our data, intermediate (*1*2) and poor metabolizer (*2*2) *CYP2C19* genotypes were observed in 34% and in 3% of our patients, respectively. We also compared the distribution of these genotypes in a sample of healthy Lithuanian subjects. The percentage of defective *1*2 and *2*2 variants in a sample of the Lithuanian population is lower than that in the sample of the patients (Tatarūnas et al., 2012).

The male patients studied were younger than the female patients. The frequency of *1*2 was similar in male and female patients. Sex differences have been described to have an impact on the response to antiplatelet therapy. Hobson and his colleagues demonstrated that there were both an elevated baseline clotting tendency and a reduced response to clopidogrel in young healthy females as compared to that in males. No such differences among males and females were observed in an older or postmenopausal female (Hobson et al., 2009). By contrast, males had a higher risk of developing CAD as compared with females, with the highest relative risk in young adults. The postmenopausal women had the same incidence of CAD as men. This might be accounted by a lack of estrogen, which can also inhibit the platelet function or have a positive effect on endothelium (Meyer et al., 2011; Kytö et al., 2015). There are other scientific data available suggesting that females have a higher metabolic activity of *CYP2C19* and a higher level of active clopidogrel metabolite than males (Xie et al., 2014). Nevertheless, clopidogrel reduces the risk of cardiovascular events in both women and men (Berger et al., 2009).

The latest European guidelines emphasized that platelet function testing or genetic testing should not be recommended in routine clinical practice on account of insufficient prospective data (Hamm et al., 2011). More and more data, however, are available about the benefit of phenotyping and genotyping to selecting antiplatelet treatment following PCI (Montalescot et al.,

2013; Kolh et al., 2014). A rapid gene trial showed a successful validation and clinical application of point-of-care genetic testing in selecting effective treatment following PCI (Roberts et al., 2012). Stimpfle and his colleagues also demonstrated that *CYP2C19* loss-of-function point-of-care genotyping identified the *CYP2C19**2 allele carriers and enabled those patients to be identified who did not respond well to conventional treatment and allowed effective treatment to be prescribed on the basis of the established genotype (Stimpfle et al., 2014). The method, which was used by our research group, allows a bedside identification of the patient's *CYP2C19* genotype in an hour. Thus, this might help differentiate the patients who respond to treatment with clopidogrel from those for whom treatment with clopidogrel is ineffective.

Fast and accurate genotyping and personalized antiplatelet can be more cost-effective and may provide fewer adverse outcomes as compared with the empirical drug dosage.

CONCLUSIONS

By using the bedside technique, *CYP2C19**2, which is responsible for a poorer effect of clopidogrel, can be identified in less than an hour. The *CYP2C19**1*2 was more frequent among patients than in the healthy population sample.

CONFLICT OF INTEREST

None declared.

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IŠEMINE ŠIRDIES LIGA SERGANČIŲ PACIENTŲ IDENTIFIKAVIMAS PAGAL CYP2C19*2 GENOTIPĄ

Santrauka

Išemine širdies liga sergantiems pacientams, kuriems išsivysto stento trombozė arba gydant antiagregantais nustatomas didelis trombocitų reaktivumas, siekėme nustatyti pataloginį CYP2C19*2 alelį, taikydami naują greitai genotipą nustatantį metodą. Iš viso tyrime dalyvavo 35 pacientai. Visi pacientai buvo ištirti dėl CYP2C19*2 naudojant *Spartan RX CYP2C19* tyrimų paketą. CYP2C19*1*1 genetinis variantas nustatytas 63 %, *1*2 – 34 %, *2*2 – 2 % tirtų pacientų. Vyrų ir moterų grupėse skirtingi genotipai pasiskirstė panašiai. CYP2C19*2 alelį, lemiantį blogesnę klopido-grelio antitrombocitinį efektą, greitu genotipo identifikacijos metodu galima nustatyti greičiau nei per valandą.

Raktažodžiai: stento trombozė, CYP2C19 variantas, klopido-greliis, lytis, antiagregantai