

**GENETIC FACTORS AFFECTING THE DEVELOPEMENT OF ACUTE  
CORONARY SYNDROMES AND THE EFFECTIVENESS OF  
ANTIPLATELET DRUGS**

**Ph.D. thesis**

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## **INTRODUCTION**

The leading causes of mortality in the western countries are still the cardiovascular diseases, especially the ischemic heart disease (IHD) and the acute coronary syndromes (ACS). Approximately sixteen million patients died because of cardiovascular events in 2001. Besides the classic cardiovascular risk factors genetic mutations are thought to have a role in the development of the ACS. The positive family history and the ACS in the young underlies the importance of the genetics in the background of the ACS.

During the past decades acetylsalicylic acid, statins, ACE inhibitors and beta receptor blockers were shown to decrease mortality in the secondary prevention of IHD. Despite the more aggressive and effective therapy the cardiovascular mortality increased. New therapeutic strategies were introduced to stop the increase of mortality. The new antiplatelet drug clopidogrel was shown to reduce mortality and morbidity in patients with ACS, however, several studies reported ineffective antiplatelet therapy with either ASA or clopidogrel. In the background of the resistance to antiplatelet therapy the role of genetic factors were thought besides drug interactions, non-compliance etc. My study focuses on the following genetic factors affecting the development of ACS and the effectiveness of antiplatelet drugs:

### **1. Angiotensin converting enzyme (ACE) insertion/deletion (I/D) polymorphism**

There is an insertion (I)/deletion (D) polymorphism in the 16 intron of the ACE gene, which affects the plasma ACE level. DD genotype represents a 2 fold increase in ACE level compared to II genotype. Controversial results have been reported in the setting of the role of ACE polymorphism in the development of ACS.

### **2. Methylene tetrahydrofolate reductase (MTHFR) polymorphism**

The patients homozygous to C-T mutation in the 677 position of the MTHFR gene have elevated homocysteine level, which is an independent risk factor of IHD, stroke and peripheral arterial disease. Some studies showed a correlation between the C-T mutation and IHD.

### **3. Glycoproteine (GP) IIIa platelet receptor PIA (A1/A2) polymorfism**

PIA is a part of the GP IIb/IIIa membrane receptor family. The T-C mutation in the 1565 position of 2 exon of glycoprotein IIIa gene leads to PIA1 or A2 alleles. Recent data suggest that PIA2 allele could be associated to the development of ACS.

### **4. Mutations of lymfotoksin alfa (LTA) gene**

Lymfotoksin alfa is an inflammatory cytokine playing a key role in the initiation of the local inflammatory processes. As during the ACS local inflammation can cause the rupture of the unstable coronary plaque, the LTA can play a role in the initiation of ACS. Two mutations (LTA 1 and LTA3) were shown to be related to the development of ACS.

### **AIMS**

The aims of our studies were to determine the possible role of the above mentioned mutations in the development of ACS. We also studied the association between the PIA polymorphism and the resistance to ASA or clopidogrel.

## **RESULTS**

### **1. PIA, ACE, MTHFR, LTA1 and LTA3 mutations in ACS patients**

117 patients were PIA1/A1 (64 %) homozygous, 63 patients were A2 carriers. A2/A2 genotype was detected in 9 cases. There were significantly less A2 homozygous people or A2 carriers among healthy controls.

There was no significant difference in the distribution of the ACE genotypes in the two populations.

MTHFR mutation appeared significantly more frequently in patients with ACS compared to control population.

The prevalence of LTA1 és LTA3 polymorphisms was not significantly different in the two populations examined.

## **2. The possible association between PIA polymorphism and ASA resistance**

ASA resistance was observed in 119 subjects (65 ACS, 30 ischemic stroke and 24 high-risk patients), while 166 (93 ACS, 39 stroke and 34 high-risk patients) showed an appropriate response to the ASA therapy.  $PI^{A2/A2}$  genotype occurred in ten subjects (8.4 %) with ASA resistance. We should note, that none of the patients with adequate ASA response was homozygous to the  $PI^{A2}$  allele. The frequency of  $PI^{A2}$  allele was significantly higher in subjects with ASA resistance than in the population showing normal response (0.21 vs. 0.14,  $p < 0.05$ ) to the drug.  $PI^A$  genotypes were in Hardy-Weinberg equilibrium. Results of the adjusted logistic regression analysis show that genotypes containing A2 allele ( $A1/A2 + A2/A2$ ) are not an independent risk for ASA resistance (Odds ratio (95 % CI) of 1.04 (0.46-2.215),  $p = 0.99$ ).

## **3. Does glycoprotein IIIa gene ( $PI^A$ ) polymorphism influence the clopidogrel resistance**

Clopidogrel insensitive platelet aggregation was observed in 38 subjects (19 with ACS, 14 with ischemic stroke and 5 with PAD), while 59 (18 with ACS, 23 with stroke and 18 with PAD) showed an appropriate response to the clopidogrel therapy. No significant differences could be found regarding the duration of the therapy between the responder and resistant patient groups (55 days and 59 days prior to the platelet aggregometry, respectively).  $PI^{A2/A2}$  genotype was revealed in one subject without clopidogrel resistance (2 %), in contrast none of the patients with clopidogrel resistance was homozygous to the  $PI^{A2}$  allele. No significant difference in the prevalence of  $PI^{A2}$  allele and the allele frequencies ( $PI^{A2}$ : 0.09 vs 0.13) was detected, not even after adjusting for the concomitant medication and risk factors.  $PI^A$  genotypes were in Hardy-Weinberg equilibrium. Our results show that the genotype containing an A2 allele

(A1/A2 or A2/A2) does not represent an independent risk factor for clopidogrel resistance.

## **SUMMARY**

Our results support the hypothesis that some of genetic mutations could play a role in the development of ACS. The Hungarian gene stock is different from other European populations, thus mutations described in the international literature should be examined in domestic population, as well, to prove their role in Hungary. The presence of  $PI^{A2}$  allele (and especially the  $PI^{A2}$  homozygosity) could affect the effect of ASA on platelet aggregation, but not the effect of clopidogrel. Our preliminary results suggest that patients with  $PI^{A2}$  homozygosity might benefit from the administration of an ADP receptor antagonist (e.g., clopidogrel) rather than applying ASA.

## **PRACTICAL IMPLICATIONS**

1. This is the first study confirming the role of  $PI^{A2}$  allele in the development of ACS in Hungarian population.
2. The MTHFR polymorphism could play a role in the formation of ACS.
3. We could not find any connection between ACS and ACE polymorphism.
4. There wasn't any relation between LTA1, LTA3 polymorphisms and the ACS.
5. Our results suggest that in case of ACS PIA and MTHFR polymorphisms should be considered instead of ACE, LTA1 and LTA3 mutations, in Hungarian population.
6. We found relationship between the occurrence of  $PI^{A2}$  allele and ASA resistance.
7. There wasn't any correlation between  $PI^{A2}$  allele and clopidogrel resistance.
8. Further investigations are needed to be conducted to examine the genetic background of the clopidogrel resistance.
9. Our studies suggest, that the examination of PIA polymorphism should be considered in case of ACS, and clopidogrel should be administered instead of ASA in case of  $PI^{A2}$  homozygosity.