

## Are Platelet Membrane Glycoprotein Polymorphisms Predictive of Arterial Thrombosis?

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Arterial thrombotic syndromes are the leading cause of morbidity and mortality in the western world. Since platelets play a pivotal role in the formation of arterial thrombi, it has been suggested that markers of platelet activation would be potentially useful to identify patients at increased risk for arterial thrombosis.

The characterization of platelet glycoprotein polymorphisms paved the way for evaluation of their prevalence in populations of patients with arterial thrombotic syndromes, including myocardial infarction and stroke. Platelet membrane glycoproteins are highly polymorphic, and the allelic variations encoding the major platelet glycoproteins –  $\alpha 2\beta 3$  (GPIIb/IIIa), GPIa-IIa and GPIb-IX – are very common.

The platelet GPIIb/IIIa is a receptor for fibrinogen and von Willebrand factor.  $PI^A$  is the alloantigen most frequently implicated in immune mediated platelet destruction. The  $PI^{A2}$  polymorphism (proline at position 33 of the  $\beta$ -3 subunit) was initially suggested to be more common in coronary artery disease patients. However, of 18 studies only 8 demonstrated a positive association, while 10 showed no association between  $PI^{A2}$  and coronary artery disease [1]. The possibility that  $PI^{A2}$  is a risk factor for coronary atherosclerosis but not for arterial thrombosis may explain the discrepancies between some of these studies [2]. The risk conferred by  $PI^{A2}$  is particularly increased in young smokers.

A number of studies focused on the platelet collagen receptor to evaluate the correlation between inheritance of the integrin  $\alpha$ -2 allele-1 coding for increased platelet  $\alpha$ -2  $\beta$ -1 density and risks of adverse outcome in patients with thrombotic disorders. In patients with myocardial infarction, three studies [3–5] demonstrated a positive correlation, with odds ratio of 2.2–3.3, while three others showed no difference [6–8]. In addition, two studies in coronary artery disease patients and one in patients who had restenosis after coronary stenting were negative [4,9,10]. In stroke patients, three studies [11–13] showed a positive association, with odds ratio of 2.0–3.0, while another study was negative [9]. Furthermore, three studies in venous thrombosis patients were negative [9,14,15]. Thus, the bulk of evidence suggests that this platelet collagen receptor polymorphism is not helpful in predicting thrombotic risk.

In this issue of IMAJ, Rosenberg et al. [16] report the results of a case-control study, in young males with a first episode of acute myocardial infarction and healthy controls, carefully designed to evaluate the risks exerted by a number of platelet glycoprotein

polymorphisms in the GPIIIa, GPIa and the GPIb alpha genes. The study groups were different for smoking and other acquired metabolic risk factors, suggesting that their size was sufficiently large to detect significant clinically relevant differences. However, the prevalence of the studied platelet glycoprotein polymorphisms, including HPA-1 a/b in GPIIIa, 807C/T in GPIa, HPA-2a/b, VNTR and Kozak-C/T in GPIb alpha, were not significantly different in patients compared to controls.

Evaluation of platelet glycoprotein polymorphisms is elaborate and costly. Since studies of currently available glycoproteins by and large do not add information to trivial metabolic risk factors, the role of any of these polymorphisms as predictors of arterial thrombosis remains controversial. Furthermore, the confounding influences of age, gender, environment and metabolic risk factors should be rigorously assessed by studies of platelet glycoprotein polymorphisms in thrombosis patients and controls.

If indeed platelet glycoprotein polymorphisms will prove non-predictive for arterial thrombosis, the information gathered in these studies could still be potentially useful. For example, some of these glycoproteins may serve as receptors for anti-platelet agents. Single nucleotide polymorphisms attract considerable interest in the rapidly evolving field of pharmacogenomics. It is possible that these single nucleotide polymorphisms on the platelet surface may differentiate between variable responses to anti-platelet agents. Functional studies in patients from the Framingham Offspring Study demonstrated increased aggregation of  $PI^{A2}$  platelets [17]. Furthermore, it was suggested that  $PI^{A1A2}$  platelets may be more sensitive to inhibition of aggregation by aspirin or abciximab [18]. The clinical relevance of these findings on drug interactions has yet to be determined by epidemiologic studies.

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