

The PIA1/A2 polymorphism of glycoprotein IIIa in relation to efficacy of antiplatelet drugs: a systematic review and meta-analysis

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WHAT IS ALREADY KNOWN ABOUT THIS SUBJECT

- Resistance to antiplatelet drugs is associated with adverse cardiovascular outcomes.
- The PIA1/A2 polymorphism of glycoprotein IIIa has been identified as a possible biomarker for antiplatelet resistance.
- There exists poor inter-study agreement in relation to the impact of carriage of the PIA2 allele.

WHAT THIS STUDY ADDS

- Carriage of the PIA2 allele does not appear to be a biomarker for resistance to aspirin, P2Y12 antagonists or glycoprotein IIb/IIIa inhibitors.
- Calculated odds ratios are dependent on the method used for defining resistance.
- Significant inter-study heterogeneity prevents detailed sub-group analysis and the need for further studies remains.

AIM

The PIA1/A2 polymorphism of glycoprotein IIIa (GPIIIa) has been associated with both antiplatelet drug resistance and increased cardiovascular events. The aim of this study was to conduct the first meta-analysis investigating the association between carriage of the PIA2 allele and resistance to currently licensed antiplatelet drugs.

METHODS

Electronic databases (MEDLINE and EMBASE) were searched for all articles evaluating genetic polymorphisms of GPIIIa. For studies where antiplatelet resistance was measured using validated techniques, pooled odds ratios (ORs) were calculated using fixed effects and random effects models.

RESULTS

Sixteen studies were eligible for statistical analysis and included 1650 PIA1 homozygous subjects and 668 carriers of the PIA2 allele. For carriers of the PIA2 allele, OR 0.924 ($n = 2318$; 95% CI 0.743, 1.151; $P = 0.481$) was observed for resistance to any antiplatelet drug, OR 0.862 ($n = 2085$; 95% CI 0.685, 1.086; $P = 0.208$) for resistance to aspirin and OR 1.429 ($n = 233$; 95% CI 0.791, 2.582; $P = 0.237$) for resistance to clopidogrel. In the aspirin cohort, sub-group analysis revealed no statistical association in either healthy subjects or those with cardiovascular disease. PIA2 carriage was marginally associated with aspirin sensitivity using the fixed effects model when identified by the PFA-100 assay ($n = 1151$; OR 0.743, 95% CI 0.558, 0.989; $P = 0.041$) but with significant heterogeneity ($I^2 = 55\%$; $P = 0.002$). Significance was lost with analysis using a random effects model.

CONCLUSIONS

The totality of published data does not support an association between carriage of the PIA2 allele and antiplatelet drug resistance. Significant heterogeneity indicates the need for larger studies using validated and standardized assays.

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Introduction

The central role of platelets is to terminate haemorrhage following vascular injury [1]. The fibrinogen receptor is integral to this process as it represents the final common pathway of platelet activation, adhesion and aggregation, and binds a combination of fibrinogen, von Willebrand factor (vWF) and fibronectin [2]. The fibrinogen receptor is the most abundant integrin on the platelet surface [3] and is formed from two subunits: glycoprotein IIb (GPIIb, integrin α_{IIb}) and glycoprotein IIIa (GPIIIa, integrin β_3). The GPIIIa subunit is polymorphic with single amino acid substitutions resulting in a number of stable allelic variants [4]. The PIA1/A2 diallelic antigen system is one of the more heavily studied due to its involvement in alloimmunity [4] and is subject to ongoing controversy relating to its possible association with cardiovascular disease and resistance to antiplatelet agents. The prevalence of the PIA2 allele is dependent on ethnicity, with a frequency of approximately 15 per 100 in Caucasian populations [5] falling to 1 per 100 in Oriental populations [6].

Lying within the extracellular domain, the PIA1 epitope is transformed into that of the PIA2 allelic variant by a single amino acid substitution of proline for leucine at position 33 [7]. The proximity of the epitope to the ligand binding site stimulated investigation into the impact of the polymorphism on the binding of fibrinogen and vWF, with results in static systems showing no significant differences in maximal binding (B_{max}) or dissociation constant (K_d) [8, 9]. In cell culture under conditions of shear stress, enhanced adhesion to both ligands was subsequently observed in the presence of the PIA2 allele, and this was attributed to enhancement of outside-in signalling [10, 11]. These disagreements between studies extend into observations regarding possible differences in basal level of platelet activation in the resting state and in the activation of platelets in response to various agonists [8, 12].

Despite this lack of clarity into the molecular impact of the single nucleotide polymorphism, carriage of the PIA2 allele showed early promise as a biomarker for cardiovascular risk when its prevalence was noted to be higher in those with acute coronary thrombosis, especially in patients with a primary event occurring under the age of 60 years [13]. Meta-analyses have since yielded inconsistent results, with no association found with myocardial infarction [14] but a positive association found with coronary artery disease more generally (odds ratio (OR) 1.10; 95% CI 1.03, 1.18) [15]. A possible mechanism for increased cardiovascular risk is resistance to the antiplatelet drug aspirin (acetylsalicylic acid), with a meta-analysis in 2008 finding a significant association between carriage of the PIA2 allele and aspirin resistance in healthy subjects (OR 2.36; 95% CI 1.24, 4.49; $P = 0.009$) [16]. However in the same study, no such difference was identified in subjects with cardiovascular disease (OR 0.92; 95% CI 0.65, 1.30; $P = 0.64$).

The heterogeneity in the findings observed in these meta-analyses are a reflection of the heterogeneity of the populations studied and the diverse methods used for assessing endpoints. There remains a need to assess definitively the PIA2 allele as a biomarker for cardiovascular disease and, if confirmed as a useful biomarker, to refine current hypotheses into the underlying mechanism. To this end, we present here the first systematic review and meta-analysis investigating the impact of carriage of the PIA2 allele on the efficacy of all currently licensed antiplatelet agents.

Methods

Data sources

Electronic databases (MEDLINE and EMBASE) were searched up until 1 April 2013 for all articles evaluating genetic polymorphisms in the platelet GpIIIa receptor. The Medical Subject Headings terms and text words used for the primary search were 'genetic', 'gene', 'polymorphism', 'mutation' and 'genotype' in combination with 'glycoprotein IIIa', 'glycoprotein IIb/IIIa', 'GP IIIa', 'GP IIb/IIIa', 'integrin beta3' and 'ITGB3'. All languages were searched and included. A secondary search was performed on all potentially relevant articles for any additional articles. The inclusion criteria for the systematic review were the use of a licensed antiplatelet drug (either *in vivo* or *in vitro*) and measurement of platelet function. Inclusion into the meta-analysis required the definition and measurement of antiplatelet resistance using validated laboratory techniques.

Statistical analysis

Data were analyzed using Comprehensive Meta-analysis software, version 2 (Biostat, USA). For each antiplatelet drug for which data were available for at least two studies, a meta-analysis was performed. For each antiplatelet drug, a pooled OR was calculated using fixed and random effects models, along with the 95% CI to measure the strength of association. Fixed effects summary ORs were calculated using the Mantel–Haenszel method [17] and the DerSimonian method was used to calculate random effects summary ORs [18]. For data where an individual was assessed for drug sensitivity by multiple methods, combined effects were calculated [19].

Tests for heterogeneity were performed for each meta-analysis, with significance set at $P < 0.05$ [20]. For assessment of publication bias, we utilized a funnel plot and the Egger regression asymmetry test [21]. In addition, the effect of individual studies on the summary OR was evaluated by re-estimating and plotting the summary OR in the absence of each study.

Results

The primary search yielded 2288 articles of which 45 were identified as being potentially relevant. Following the

Table 1

Summary of number of studies included in the systematic review and meta-analysis

Antiplatelet drug	Number of studies included in systematic review	Number of studies included in meta-analysis
Aspirin	28	14
Clopidogrel	8	3
Ticagrelor	1	0
Abciximab	5	0*
Tirofiban	3	0*
Eptifabide	4	0*

*There are no accepted definitions of GPIIb/IIIa inhibitor resistance and so this drug class was not eligible for inclusion in the meta-analysis.

addition of articles from the secondary search, a total of 36 articles met the inclusion criteria for the review, of which 16 contained suitable data for statistical analysis. Table 1 summarizes the number of studies identified based on individual antiplatelet drugs. Studies that were not eligible for inclusion in the statistical analysis are summarized in Tables 2–4.

The 16 studies analyzed statistically included 1650 subjects who were homozygous for PIA1 and 668 carriers of the PIA2 allele. An OR of 0.924 (95% CI 0.743, 1.151; $P = 0.481$) was observed for resistance to any antiplatelet drug in subjects carrying the PIA2 allele (Figure 1). Significant heterogeneity existed between studies ($P = 0.006$) and analysis using the random effects model did not lead to a significant association (OR 0.955, 95% CI 0.663, 1.376; $P =$

Table 2

Summary of studies investigating the association of the PIA1/A2 polymorphism and resistance to aspirin which could not be included in the statistical analysis

Study	Subject characteristics	Assessment methods	Comment
Andrioli <i>et al.</i> (2000) [36]	Healthy subjects ($n = 16$)	LTA (AA agonist)	PIA2 carriers demonstrated increased sensitivity to antiplatelet action of <i>in vitro</i> aspirin ($1\text{--}100 \mu\text{mol l}^{-1}$)
Boudoulas <i>et al.</i> (2001) [37]	Healthy subjects ($n = 30$)	LTA (epinephrine agonist)	PIA2 carriers more sensitive to antiplatelet action of low dose <i>in vitro</i> aspirin in response to epinephrine agonist
Cooke <i>et al.</i> (1998) [38]	Healthy subjects ($n = 26$)	LTA (ADP and epinephrine agonist)	PIA1 homozygotes associated with a reduced response to <i>in vitro</i> aspirin ($0.053\text{--}53 \mu\text{mol l}^{-1}$)
Cooke <i>et al.</i> (2006) [39]	Stable coronary artery disease ($n = 20$)	LTA (ADP, collagen and epinephrine agonists) GPIIb/IIIa activation and α -granule release measured by flow cytometry (ADP, collagen and epinephrine agonists)	PIA2 carriers significantly were less inhibited by <i>in vivo</i> aspirin (325 mg once daily for 10 days) when collagen used as agonist
Dropinski <i>et al.</i> (2007) [40]	Male coronary artery disease ($n = 28$)	Thrombin generation and bleeding time	PIA2 carriers associated with reduced response to <i>in vivo</i> aspirin (300 mg once daily for 2 weeks)
Isordia-Salas <i>et al.</i> (2012) [41]	Emergency percutaneous coronary intervention for acute coronary syndrome ($n = 60$)	LTA (AA agonist)	No association between PIA2 carriers and non-response to <i>in vivo</i> aspirin (300 mg)
Lepantalo <i>et al.</i> (2006) [42]	Elective percutaneous coronary intervention ($n = 101$)	LTA (AA agonist) PFA-100 (collagen/epinephrine cartridge)	No association between PIA2 carriers and response to <i>in vivo</i> aspirin (mean dose 100 mg once daily)
Lim <i>et al.</i> (2007) [43]	Postoperative period following coronary artery bypass grafting ($n = 63$)	LTA (ADP, collagen and epinephrine agonists)	PIA2 carriers had more impaired response to <i>in vivo</i> aspirin (325 mg or 100 mg once daily) but did not reach statistical significance
Michelson <i>et al.</i> (2000) [12]	Healthy subjects ($n = 56$)	LTA (epinephrine agonist)	PIA1/A2 more sensitive to <i>in vitro</i> aspirin than homozygous PIA1 and homozygous PIA2
Morawski <i>et al.</i> (2005) [45]	Patients undergoing coronary artery bypass grafting ($n = 51$)	PFA-100 (collagen/ADP and collagen/epinephrine cartridges) Bleeding time	PIA2 carriers more sensitive to <i>in vivo</i> aspirin (150 mg)
Stepien <i>et al.</i> (2007) [44]	Elective percutaneous coronary intervention ($n = 31$)	Thrombin generation Soluble CD40 ligand (sCD40L)	PIA2 carriers had statistically higher thrombin generation but no difference if sCD40L generation in response to <i>in vivo</i> aspirin
Szczeklik <i>et al.</i> (2000) [46]	Healthy male subjects ($n = 80$)	Bleeding time and thrombin generation	Carriers of PIA2 allele demonstrated significantly reduced antiplatelet effect following <i>in vivo</i> aspirin (300 mg)
Undas <i>et al.</i> (1999) [47]	Healthy male subjects ($n = 40$)	Thrombin generation	PIA2 carriers had significantly impaired depression of thrombin generation in response <i>in vivo</i> aspirin (75 mg once daily for 7 days)
Undas <i>et al.</i> (2001) [48]	Healthy male subjects ($n = 24$)	Various measures of thrombin generation in bleeding time blood	PIA2 carriers had significantly impaired antithrombotic action in response to <i>in vivo</i> aspirin (75 mg once daily for 7 days)

AA, arachidonic acid; ADP, adenosine diphosphate; GPIIb/IIIa, glycoprotein IIb/IIIa; LTA, light transmission aggregometry.

Table 3

Summary of studies investigating the association of the PIA1/A2 polymorphism and resistance to P2Y12 antagonists which could not be included in the statistical analysis

Study	Subject characteristics	Assessment methods	Comment
Angiolillo <i>et al.</i> (2004) [49]	Elective percutaneous coronary intervention (<i>n</i> = 38)	GPIIb/IIIa activation and P-selectin expression measured by flow cytometry (ADP agonist)	PIA2 carriers had significantly higher GPIIb/IIIa activation in response to <i>in vivo</i> clopidogrel (single 300 mg dose)
Cooke <i>et al.</i> (2006) [39]	Stable coronary artery disease (<i>n</i> = 20)	LTA (ADP, collagen and epinephrine agonists) GPIIb/IIIa activation and α -granule release measured by flow cytometry (ADP, collagen and epinephrine agonists)	PIA1/A2 genotype significantly more sensitive to ADP and trending to be less sensitive to collagen when compared to PIA1/A1 following <i>in vivo</i> clopidogrel (75 mg once daily for 10 days)
Dropinski <i>et al.</i> (2005) [50]	Male coronary artery disease or myocardial infarction >6 months previously (<i>n</i> = 48)	Bleeding time and thrombin generation PFA-100 (ADP/collagen cartridge) GPIIb/IIIa activation and P-selectin expression measured by flow cytometry (ADP agonist)	Greater increase in bleeding time and PFA-100 closure time for PIA2 carriers, with a greater reduction of P-selectin expression PIA2 carriers demonstrated superior antithrombotic response to <i>in vivo</i> clopidogrel (75 mg once daily for 14 days)
Fontana <i>et al.</i> (2006) [24]	Healthy males (<i>n</i> = 94)	LTA (ADP agonist)	No association between PIA2 carriers and response to <i>in vivo</i> clopidogrel (75 mg once daily for 1 week)
Isordia-Salas <i>et al.</i> (2012) [41]	Emergency percutaneous coronary intervention for acute coronary syndrome (<i>n</i> = 60)	LTA (ADP agonist)	No association between PIA2 carriers and non-response to <i>in vivo</i> clopidogrel (300 mg loading dose)
Storey <i>et al.</i> (2009) [53]	Stable atherosclerotic disease or non-ST segment elevation acute coronary syndrome (<i>n</i> = 147)	LTA (ADP agonist)	PIA2 allele did not influence response to <i>in vivo</i> ticagrelor (100 mg once daily for 28 days)

ADP, adenosine diphosphate; GPIIb/IIIa, glycoprotein IIb/IIIa; LTA, light transmission aggregometry.

Table 4

Summary of studies investigation the association of the PIA1/A2 polymorphism and resistance to GPIIb/IIIa inhibitors

Study	Subject characteristics	Drug exposure	Assessment methods	Comment
Aalto-Setälä <i>et al.</i> (2005) [54]	Healthy subjects (<i>n</i> = 28)	Abciximab <i>in vitro</i> (0–3 $\mu\text{g ml}^{-1}$) Tirofiban <i>in vitro</i> (0–70 ng ml^{-1}) Eptifabide <i>in vitro</i> (0–1 $\mu\text{g ml}^{-1}$)	PFA-100 (ADP/epinephrine cartridges)	PIA2 carriers more sensitive to antiplatelet effects of tirofiban with sodium citrate as anticoagulant Effect abolished if PPACK anticoagulant used
Michelson <i>et al.</i> (2000) [12]	Healthy subjects (<i>n</i> = 56)	Abciximab <i>in vitro</i> (0.5–2.5 $\mu\text{g ml}^{-1}$)	LTA (ADP agonist)	PIA2 allele conferred increased sensitivity
Sirotkina <i>et al.</i> (2007) [55]	Healthy subjects (<i>n</i> = 35)	Eptifabide <i>in vitro</i> (50–150 mg ml^{-1})	LTA (ADP agonist)	No significant difference in aggregation between genotypes
Verdoia <i>et al.</i> (2013) [56]	Elective percutaneous coronary intervention (<i>n</i> = 40) Elective percutaneous coronary intervention (<i>n</i> = 40)	Abciximab <i>in vivo</i> (0.25 mg kg^{-1} bolus, 0.125 $\mu\text{g kg}^{-1} \text{min}^{-1}$ infusion) Tirofiban <i>in vivo</i> (25 $\mu\text{g kg}^{-1}$ bolus, 0.1 $\mu\text{g kg}^{-1} \text{min}^{-1}$ infusion) or Eptifabide <i>in vivo</i> (double bolus 180 $\mu\text{g kg}^{-1}$, 2 $\mu\text{g kg}^{-1} \text{min}^{-1}$ infusion)	Impedance aggregometry (AA, ADP, collagen, prostaglandin E1 and TRAP agonists)	The PIA2 allele did not influence platelet response to either abciximab or the small molecule inhibitors
Weber <i>et al.</i> (2002) [57]	Healthy subjects (<i>n</i> = 62) and Coronary artery disease (<i>n</i> = 177)	Abciximab <i>in vitro</i> (0.03–3 $\mu\text{g ml}^{-1}$) Tirofiban <i>in vitro</i> (0.3–30 nmol l^{-1}) Eptifabide <i>in vitro</i> (0.01–1 $\mu\text{g ml}^{-1}$)	Fibrinogen binding measured by flow cytometry (ADP agonist)	The PIA2 allele did not significantly influence fibrinogen binding in either healthy subjects or patients with coronary artery disease
Wheeler <i>et al.</i> (2002) [58]	Patients undergoing percutaneous coronary intervention (<i>n</i> = 87)	Abciximab <i>in vivo</i> (0.25 mg kg^{-1} bolus, 10 $\mu\text{g min}^{-1}$ infusion)	LTA (ADP agonist) Ultegra rapid platelet function assay (TRAP agonist) Abciximab binding assay	Less completely inhibited at 1 h as assessed by LTA Less completely inhibited at 1 h and 24 h as assessed by Ultegra PIA2 carriers had less receptor occupancy at 24 h

AA, arachidonic acid; ADP, adenosine diphosphate; LTA, light transmission aggregometry; PPACK, Phe-Pro-Arg-chloromethylketone; TRAP, thrombin receptor activating peptide.

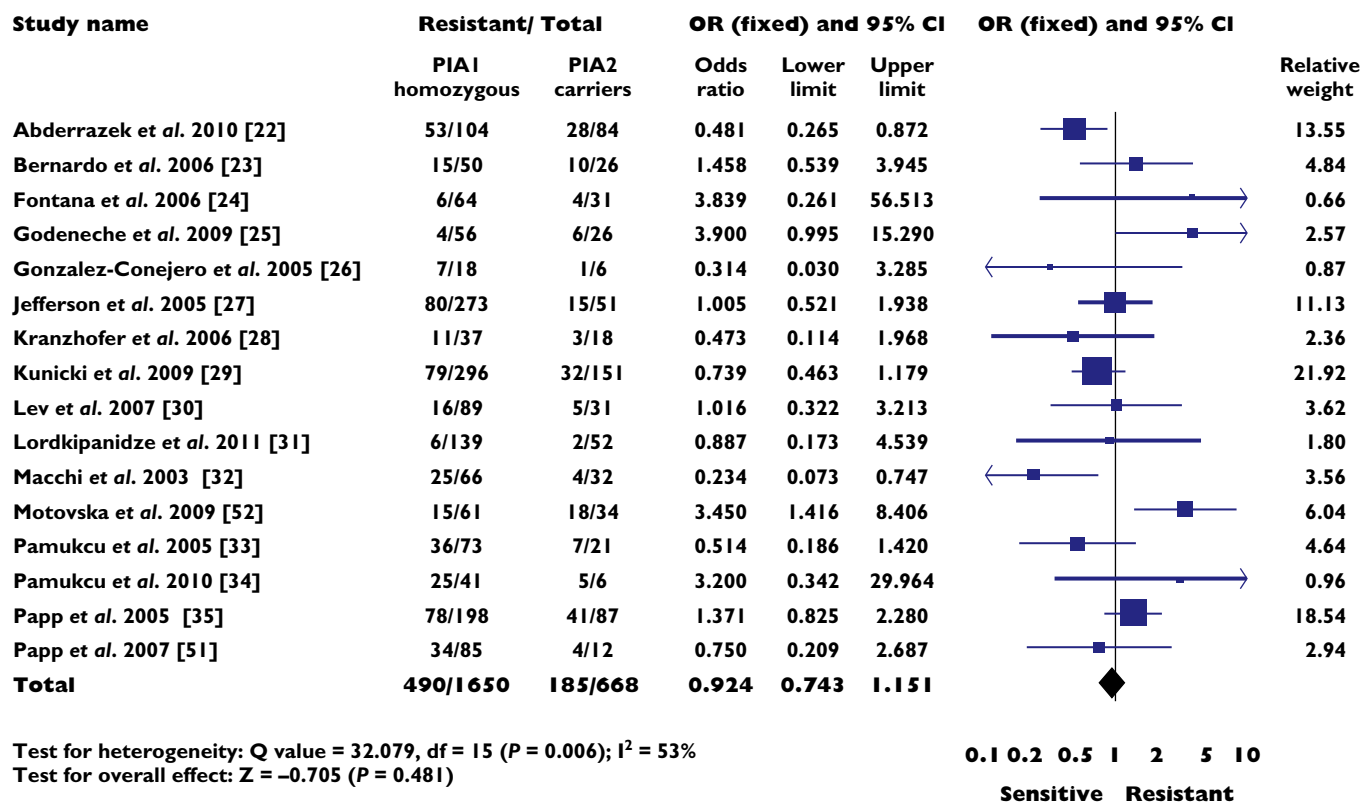


Figure 1

Association between the PIA1/A2 polymorphism and resistance to all antiplatelet drugs. OR, odds ratio; CI, confidence interval

0.804). The distribution of the log OR in relation to precision was symmetrical with a non-significant Egger regression test ($P = 0.675$), suggesting low probability of publication bias.

Aspirin

Fourteen of the 28 studies investigating aspirin resistance were eligible for inclusion in the statistical analysis and included 1463 subjects who were homozygous for the PIA1 allele and 622 who carried the PIA2 allele [22–35]. The remaining 14 studies are summarized in Table 2.

An OR of 0.862 (95% CI 0.685, 1.086; $P = 0.208$) was observed for aspirin resistance in subjects carrying the PIA2 allele (Figure 2). Significant heterogeneity was again present ($P = 0.033$) with a low probability of publication bias ($P = 0.663$).

Initial sub-group analysis of the aspirin cohort considered whether sensitivity to the drug in relation to the polymorphism was dependent on the presence or absence of cardiovascular disease, since such a dependence was previously reported [16]. No such association was identified either in patients with cardiovascular disease (OR 0.861, 95% CI 0.683, 1.087; $P = 0.208$) or in healthy subjects (OR 0.927, 95% CI 0.158, 5.434; $P = 0.933$) (Figure 3A).

A further sub-group analysis of the aspirin cohort considered how the methodology used to assess the presence of aspirin resistance might influence any association. The available data permitted the comparison of two methods: light transmission aggregometry (LTA) and the point-of-care assay PFA-100 (Figure 3B).

The use of LTA did not reveal a significant association between carriage of the PIA2 allele and aspirin resistance (OR 1.248, 95% CI 0.862, 1.809; $P = 0.241$), but did show significant homogeneity between studies ($P = 0.778$). This homogeneity is despite variation in agonists used to identify aspirin resistance, with three studies using the conventional cut-off value of 20% aggregation in response to arachidonic acid (AA) [24, 27, 31], and the remaining two studies assessing aggregation in response to two independently administered agonists [30, 35]. Sub-group analysis of the three studies using AA as the sole agonist did not increase homogeneity when compared with the entire aspirin LTA cohort ($P = 0.535$), nor result in a significant association with aspirin resistance (OR 1.053, 95% CI 0.579, 1.919; $P = 0.886$).

Use of the PFA-100 assay revealed a significant association between aspirin sensitivity and carriage of the PIA2 allele (OR 0.743, 95% CI 0.558, 0.989; $P = 0.041$), but with

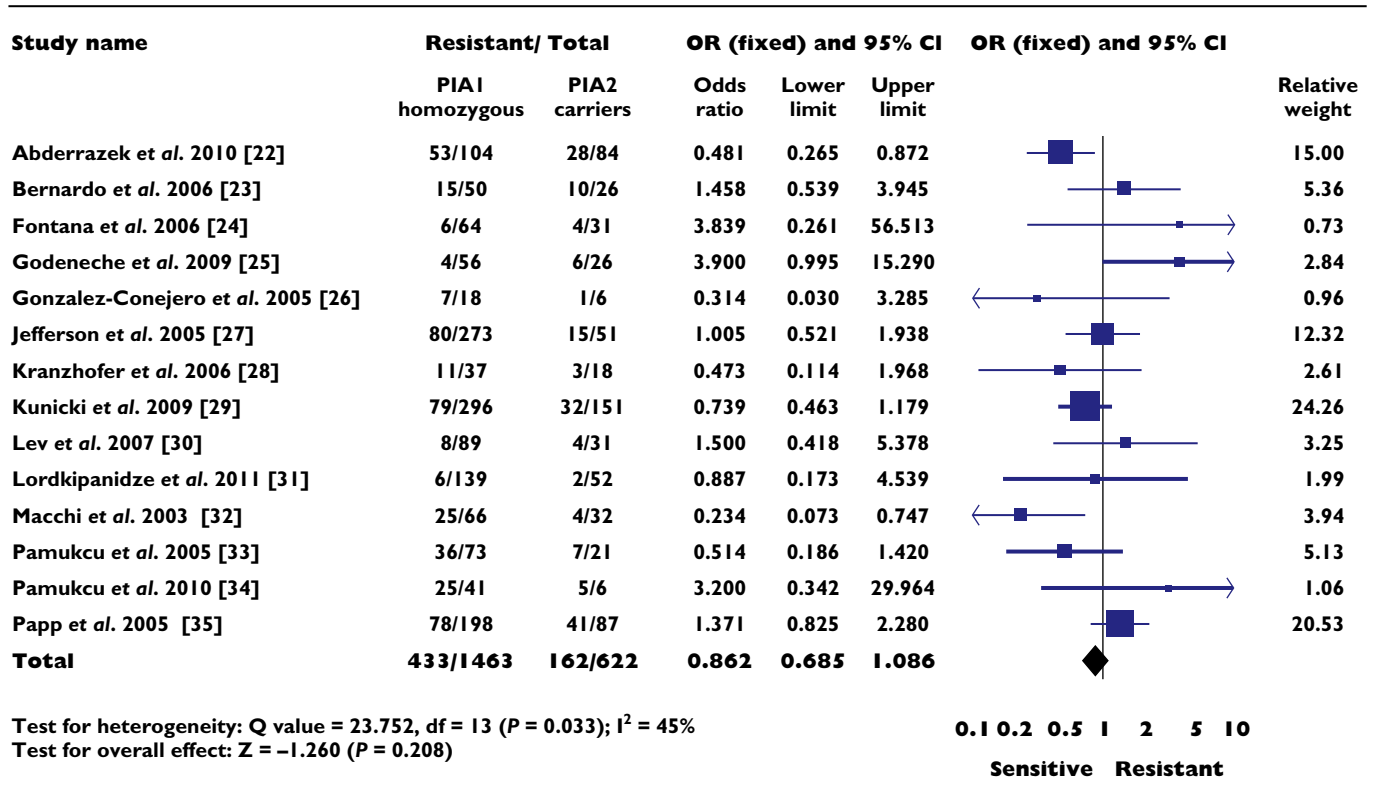


Figure 2

Association between the PIA1/A2 polymorphism and aspirin resistance. OR, odds ratio; CI, confidence interval

significant heterogeneity between studies ($P = 0.023$); significance was lost with use of the random effects model (OR 0.793, 95% CI 0.522, 1.204; $P = 0.277$). As with LTA, the PFA-100 assay enables the testing of different agonists, with the choice of either a collagen and epinephrine (CEPI) cartridge or a collagen and adenosine diphosphate (CADP) cartridge. In all studies, the CEPI cartridge was used to define aspirin resistance with a mean closure time of $<198 \pm 37$ s. In four of these studies, platelet reactivity in response to the CADP cartridge was compared between the CEPI-defined aspirin resistant and aspirin sensitive subjects. A significant ($P < 0.01$) reduction in CADP closure time was seen in those who had CEPI-defined aspirin resistance, suggesting that these patients had an increase in global platelet reactivity independent of changes related solely to the COX-1 pathway [23, 25, 29, 32].

Studies that were not suitable for inclusion in the meta-analysis included 300 healthy subjects and 354 with cardiovascular disease (Table 2) [12, 36–48]. Conclusions from the studies were similarly heterogeneous, with half of the studies suggesting an association between carriage of the PIA2 allele and aspirin resistance [39, 40, 43, 44, 46–48].

P2Y12 receptor antagonists

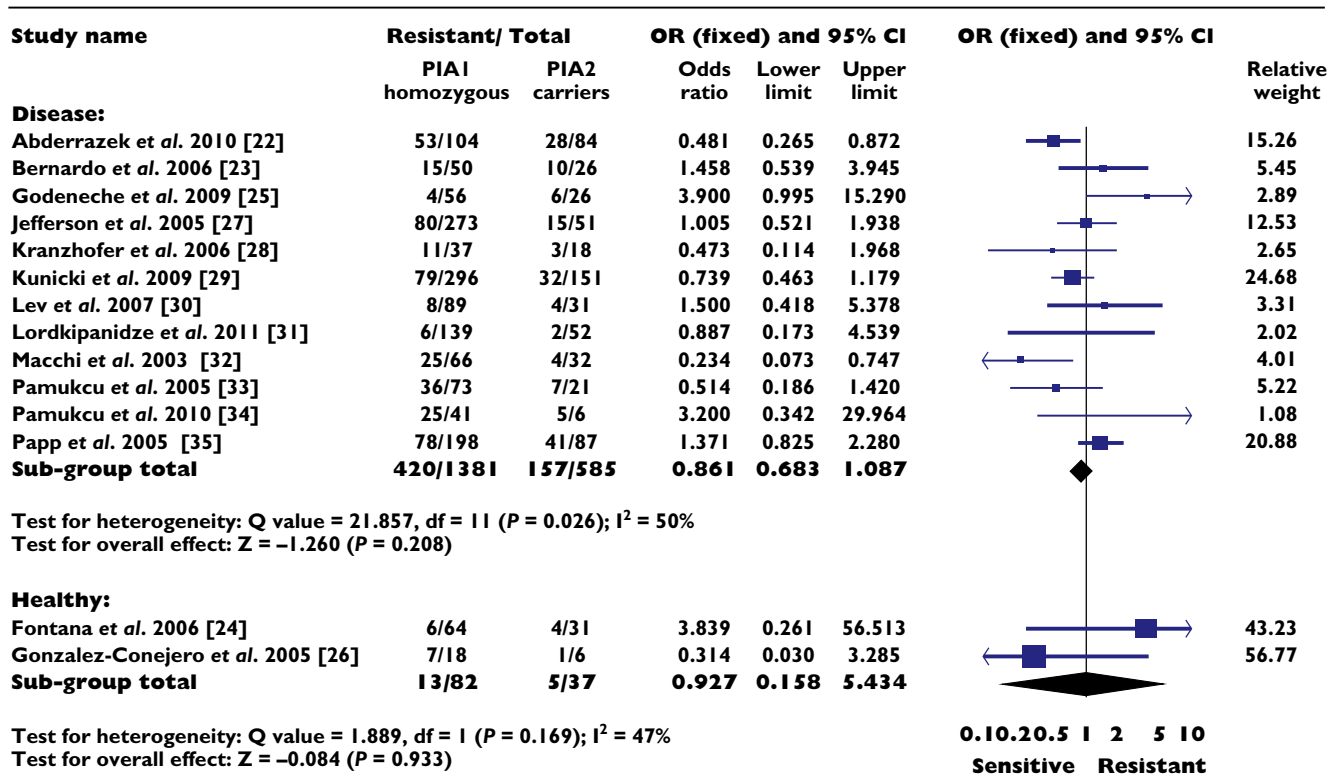
Nine studies were identified that investigated the impact of the PIA1/PIA2 polymorphism on the efficacy of P2Y12

antagonists [24, 30, 39, 41, 49–53]. All participants in these studies received *in vivo* clopidogrel except for a single study investigating *in vivo* ticagrelor [53]. No studies were identified that considered the efficacy of prasugrel in association with carriage of the PIA2 allele. Three studies were analyzed and included 100 clopidogrel-resistant and 212 clopidogrel-sensitive patients with coronary artery disease [30, 51, 52]. An OR of 1.429 (95% CI 0.791, 2.582; $P = 0.237$) was observed for clopidogrel resistance in subjects carrying the PIA2 allele (Figure 4). Significant heterogeneity existed between studies ($P = 0.034$). Despite the small number of studies, the distribution of the log OR in relation to precision was symmetrical with a non-significant Egger regression test ($P = 0.48$), suggesting low probability of publication bias.

Sub-group analysis of the two studies utilizing LTA decreased the OR to below unity, suggesting a possible association of clopidogrel sensitivity with PIA2 carriage, but this did not achieve significance (OR 0.712, 95% CI 0.322, 1.571; $P = 0.400$) [30, 51].

Six studies did not contain adequate data for statistical analysis and are summarized in Table 3 [24, 39, 41, 49, 50, 53]. Of these studies, one reported a positive association between carriage of the PIA2 allele and clopidogrel resistance [49], one reported a negative association [50] and the remainder failed to identify any association. The single

A



B

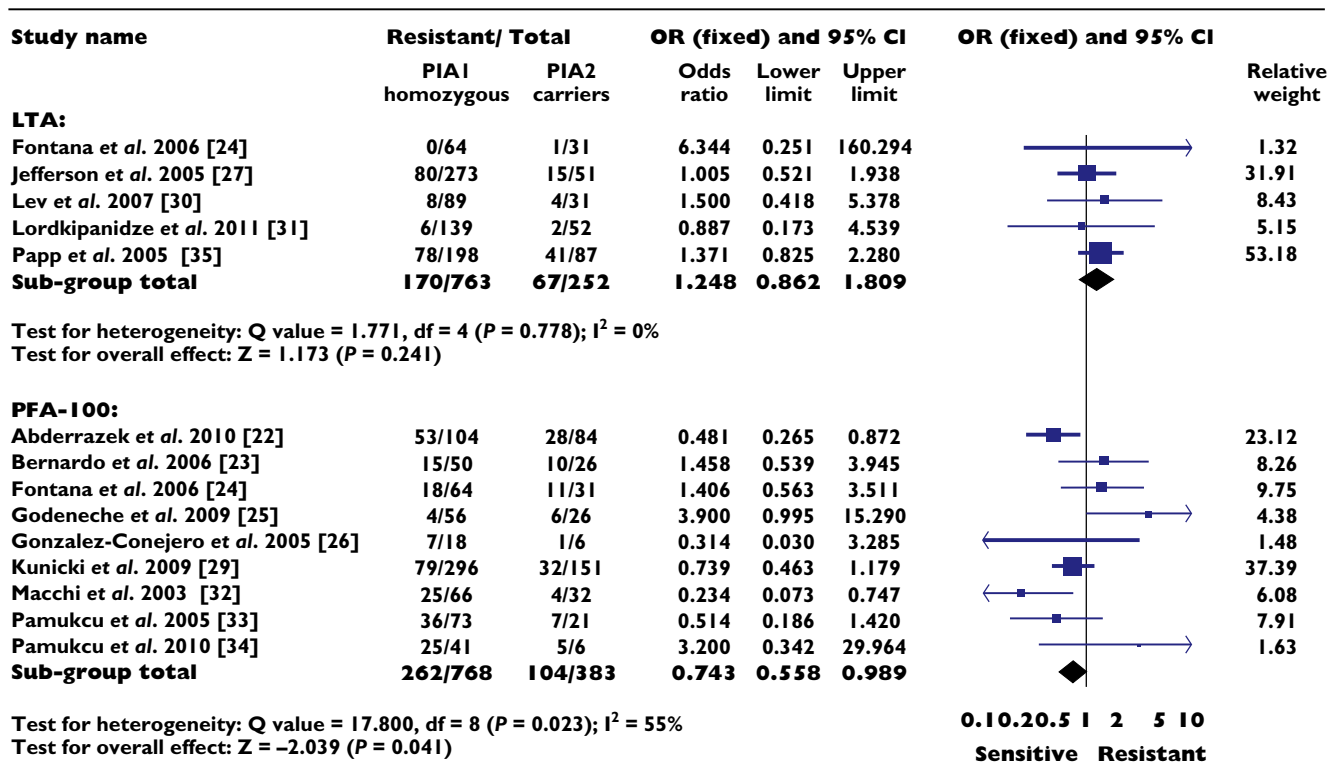


Figure 3

Association between the PIA1/A2 polymorphism and aspirin resistance, sub-group analyses. (A) Cardiovascular disease vs. healthy subjects. (B) Light transmission aggregometry vs. PFA-100. OR, odds ratio; CI, confidence interval

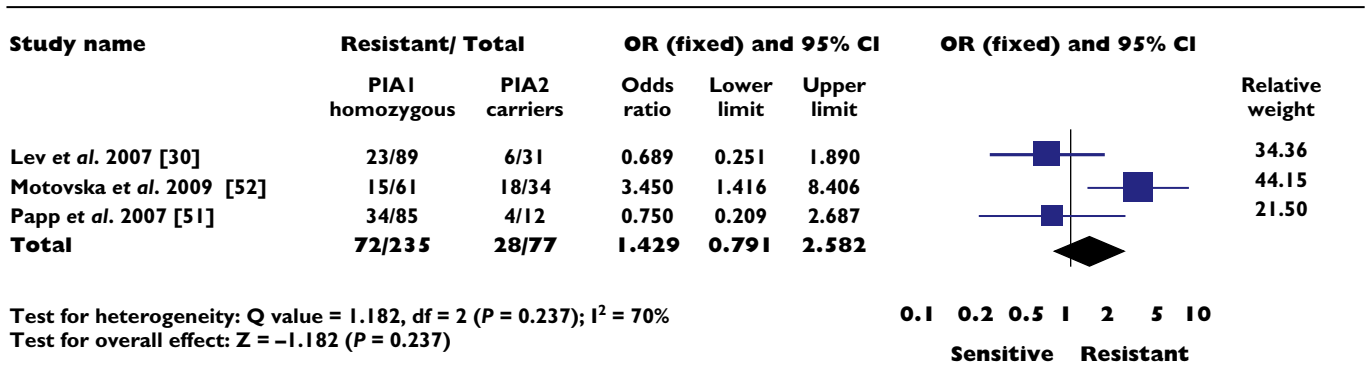


Figure 4

Association between the PIA1/A2 polymorphism and clopidogrel resistance. OR, odds ratio; CI, confidence interval

study considering the influence that carriage of the PIA2 allele might have on the efficacy of *in vivo* ticagrelor found no significant association in patients with cardiovascular disease [53]. Within all of the above studies no trend in response was identified in relation to subject characteristics, assessment methods or drug dosing schedules.

Glycoprotein IIb/IIIa inhibitors

Six studies were identified which examined the three licensed GPIIb/IIIa inhibitors (GPIs) abciximab, tirofiban and eptifabide [12, 54–58]. None of these studies was suitable for inclusion into the meta-analysis due to the lack of existing criteria defining resistance within this drug class. Table 4 summarizes the findings of these studies.

Of the six studies, a significant difference in drug efficacy related to PIA2 carriage was observed in two studies but with opposite effects. Carriage of the PIA2 allele was observed to exhibit increased sensitivity to *in vitro* abciximab in 56 healthy subjects, but a statistically significant difference was seen only between PIA1/A2 vs. PIA2/A2 genotypes (P = 0.046) [12]. The converse effect was observed *in vivo* where, in the presence of the PIA2 allele, aggregation was found to be less completely inhibited following a bolus of abciximab during percutaneous coronary intervention [58]. In this clinical study, the observed effect was assay-dependent as significance was no longer present at 24 h when assessed by LTA, but remained present when assessed by the Ultegra rapid platelet function assay. Using impedance aggregometry neither of these findings has been replicated *in vivo* or *in vitro* [56, 57].

The findings of studies investigating the small molecule GPIs (eptifabide and tirofiban) concur in their results, irrespective of subject cohort or method of drug exposure. In four studies comprising a total of 125 healthy subjects and 217 patients, no differences were detected between PIA2 carriers and non-carriers using a variety of assessment methods [54–57].

Discussion

In this systematic review and meta-analysis of 36 studies totalling 3539 subjects we have found no consistent association between the PIA1/A2 polymorphism and resistance to antiplatelet drugs, either overall or for individual drugs. Given the heterogeneous nature of the studies in relation to subject recruitment, antiplatelet drug used and methods of measurement, sub-group analyses were performed according to the parameters described above.

Results from analysis of the aspirin cohort broadly concur with those of a meta-analysis into the genetics of aspirin resistance performed in 2007 [16]. However, in healthy subjects, Goodman *et al.* observed that carriage of the PIA2 allele was significantly associated with aspirin resistance (OR 2.26, 95% CI 1.24, 4.49; P = 0.009) as opposed to the non-significant result presented here (OR 1.102, 95% CI 0.480, 2.533; P = 0.229). Our exclusion of two studies based on a lack of appropriate definition of aspirin resistance and non-standard measures of platelet function accounts for this discrepancy [46, 47].

LTA in response to AA and PFA-100 (using the CEPI cartridge) are the assays most commonly used to define aspirin resistance, as they are designed to challenge the COX-1 pathway of platelet activation which aspirin specifically inhibits. Despite both assays targeting the same pathway, they gave rise to apparently diverging results, with the 'gold standard' assay LTA favouring resistance albeit non-significantly (OR 1.248, 95% CI 0.862, 1.809; P = 0.241) and the point-of-care test PFA-100 favouring sensitivity (OR 0.743, 95% CI 0.558, 0.989; P = 0.041) to aspirin in the presence of the PIA2 allele. The discrepancy in the results of the two assays is perhaps unsurprising given the significant intra- and inter-individual variability associated with platelet function testing. In fact, the PFA-100 correlates especially poorly with LTA, as demonstrated by the two techniques identifying a prevalence of aspirin resistance of 59.6% and 4.0%, respectively, in a cohort of 201

patients with stable coronary artery disease [59]. This variation in assay sensitivity and specificity is also clearly highlighted by the range of intra-study results that are dependent on method of assessment [24, 26, 39, 44, 58]. Additionally, whilst the criteria for aspirin resistance by LTA were relatively uniform, the closure time identifying aspirin resistance by PFA-100 varied greatly between studies ($<198 \pm 37$ s).

The choice of assay with which to measure aspirin resistance is generally guided by the wish to investigate COX-1 dependent responses, with the assumption that failure to suppress COX-1 activity by aspirin will give rise to increased cardiovascular events. Indeed, a correlation between residual COX-1 activity and cardiovascular events has been demonstrated in a number of studies [60, 61], but should not be assumed to be the sole determinant of platelet aggregation in aspirin-treated patients. In a study of 700 aspirin-treated patients undergoing cardiac catheterization, it was response as assessed by the PFA-100 CADP cartridge rather than the CEPI cartridge that was significantly associated with major adverse cardiovascular outcomes [62]. It is interesting to note that, in the aspirin cohort, all who were investigated using the PFA-100 CADP cartridge demonstrated a significantly shortened closure time if defined aspirin resistant by the CEPI cartridge. This observation suggests an increased underlying level of platelet reactivity in such patients, independent of COX-1.

One limitation of this systematic review is the paucity of studies considering drugs other than aspirin, with clopidogrel being the only other drug eligible for statistical analysis. The weight of studies not included in the meta-analysis do however support the statistical results in demonstrating no significant association between the PIA1/A2 polymorphism and clopidogrel resistance.

Similarly, there were few studies considering the impact of the PIA2 allele on GPI efficacy, with the additional limitation that, due to there being no defined criteria for resistance to these drugs, statistical analysis was not possible. Once again however, no consistent pattern was seen between PIA2 carriage and response to the three licensed GPIs, *in vitro* or *in vivo*.

These technical issues and study limitations likely contribute to the significant heterogeneity observed within our analyses. Publication bias was not identified to be a contributing factor. Such heterogeneity indicates that, if more data were available in the future, more refined subgroup analyses considering parameters such as gender, age, co-morbidities and biochemical indices may reveal a possible effect of the PIA1/A2 polymorphism on antiplatelet drug resistance.

In conclusion, the data presented do not support an association between carriage of the PIA2 allele of GPIIIa and resistance to antiplatelet drugs, in either healthy subjects or those with cardiovascular disease, irrespective of the method of assessment. Significant heterogeneity was observed throughout the analyses and underlines the

necessity for larger studies using validated and standardized assays to assess accurately any potential impact of this polymorphism.

Competing Interests

Both authors have completed the Unified Competing Interest form at http://www.icmje.org/coi_disclosure.pdf (available on request from the corresponding author) and declare no support from any organization for the submitted work, no financial relationships with any organizations that might have an interest in the submitted work in the previous 3 years no other relationships or activities that could appear to have influenced the submitted work.

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