



MONASH University

*Defining the role of protease-activated  
receptor 4 during thrombosis*

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## Abstract

**Background:** Thrombin-induced platelet activation is a major contributor to the formation of the arterial thrombi that cause two of the most lethal manifestations of cardiovascular disease – acute myocardial infarction and occlusive stroke. Thrombin activates human platelets via protease-activated receptors 1 (PAR1) and 4 (PAR4). Much is known about the function of the higher affinity receptor, PAR1, and a PAR1 antagonist is in clinical use for the prevention of myocardial infarction. However, PAR1 inhibition increases the rate of intracranial haemorrhage, suggesting there remains much to learn regarding how to best manipulate thrombin signalling in platelets for safe and effective antithrombotic therapy. As a result, there has been much recent interest in examining PAR4 as an antiplatelet drug target. However, in contrast to PAR1, far less is known about the function of PAR4 during platelet activation and thrombus formation.

**Aim:** To examine the function of PAR4 during thrombosis in order to determine whether targeting PAR4 represents a valid antithrombotic approach.

**Key Findings:** First, the function of PAR4 on human platelets during thrombosis was determined. To achieve this, a function-blocking PAR4 polyclonal antibody was developed and used to probe for platelet activation events dependent on PAR4. This approach determined that thrombin-induced production of procoagulant platelets is mediated predominantly by PAR4, and is largely independent of PAR1. This effect translated into impaired thrombin production and fibrin formation in an *ex vivo* human thrombosis model, rationalising PAR4 as a novel antithrombotic target. Second, the effects of targeting PAR4 were examined *in vivo* using mouse genetic models. Previous studies have shown that mice genetically deficient in PAR4 are protected against thrombosis but do not bleed spontaneously. However, here, PAR4-deficient mice exhibited a subtle but significant spontaneous bleeding phenotype in the

perinatal period, suggesting PAR4 inhibition in humans may cause unwanted bleeding effects. However, in contrast to human platelets, mouse platelets do not express PAR1 and so thrombin signalling in mouse platelets is reliant entirely on PAR4. These *in vivo* studies therefore highlighted the need for an animal model of platelet-PAR expression that would more accurately predict the impact of PAR4 inhibition in humans. Therefore, third, the production of a mouse that expresses human PAR1 on the platelet surface was attempted. This approach failed to provide platelet PAR1 expression and suggests alternative animal models will be required for preclinical evaluation of PAR4 antagonism. Finally, the first human monoclonal inhibitory antibody targeting PAR4 was developed as a potential therapeutic strategy in humans. This antibody specifically and effectively inhibited thrombin-induced cleavage and activation of PAR4 and provided marked antithrombotic effects in human blood. Furthermore, this antibody-based approach to inhibiting PAR4 was equally effective against a commonly expressed and clinically significant sequence variant of PAR4 that is resistant to an existing PAR4 antagonist.

**Conclusion:** The findings from this thesis suggest that targeting PAR4 is valid antithrombotic approach. These studies have uncovered a distinct role for PAR4 in the setting of human thrombosis, developed an antibody against PAR4 with therapeutic potential, and shown that antibody-based inhibition of PAR4 provides an antithrombotic effect distinct from all current antiplatelet drugs and that may have advanced clinical utility over other PAR4 antagonists. Together, these studies provide strong rationale for the pursuit of PAR4 antagonists for the prevention of arterial thrombosis.

## **Publications during enrolment**

**French SL** and Hamilton JR. (2017) Drugs targeting PAR4 improve the antithrombotic therapeutic window. *Ann Trans Med.* doi: 10.21037/atm.2017.09.10

**French SL** and Hamilton JR. (2017) Perinatal lethality of *Par4*<sup>-/-</sup> mice delivered by primiparous dams reveals spontaneous bleeding in mice without platelet thrombin receptor function. *Platelets.* doi: 10.1080/09537104.2017.1349310.

**French SL**, Paramitha AC, Moon MJ, Dickins RA and Hamilton JR. (2016) Humanizing the protease-activated receptor (PAR) expression profile in mouse platelets by knocking PAR1 into the Par3 locus reveals PAR1 expression is not tolerated in mouse platelets. *PLoS One.* 11 (10): e016556.

**French SL**, Arthur JF, Lee H, Nesbitt WS, Gardiner EE, Andrews RK and Hamilton JR. (2016) Inhibition of protease-activated receptor 4 impairs platelet procoagulant activity during thrombus formation in human blood. *J Thromb Haemost.* 14(8):1642-54

**French SL** and Hamilton JR (2016). Protease-activated receptor-4: from structure to function and back again. *Brit J Pharmacol.* 173(20):2952-65

**French SL**, Arthur JF, Tran HA and Hamilton JR (2015) Approval of the first protease-activated receptor antagonist: rationale, development, significance, and considerations of a novel anti-platelet agent. *Blood Reviews.* 29: 179-89.

## Declaration: Thesis including published works

I hereby declare that this thesis contains no material which has been accepted for the award of any other degree or diploma at any university or equivalent institution and that, to the best of my knowledge and belief, this thesis contains no material previously published or written by another person, except where due reference is made in the text of the thesis.

This thesis includes 3 original research papers and 3 reviews published in peer reviewed journals plus 1 submitted manuscript. The core theme of the thesis is 'Protease-activated receptor 4 during thrombosis'. The ideas, development and writing up of all the papers in the thesis were the principal responsibility of myself, the student, working within the Australian Centre for Blood Diseases under the supervision of Justin Hamilton. The inclusion of co-authors reflects the fact that the work came from active collaboration between researchers and acknowledges input into team-based research.

In the case of *Chapters 2-5* my contribution to the work involved the following:

Thesis Chapter	Publication Title	Status (published, in press, accepted or returned for revision, submitted)	Nature and % of student contribution	Co-author name(s) Nature and % of Co-author's contribution*	Co-author(s), Monash student Y/N*
2	Inhibition of protease-activated receptor 4 impairs platelet procoagulant activity during thrombus formation in human blood	Published	57%. Writing of manuscript, performed all experiments and data analysis for Figures 1, 3 and 4. Contributions to Figures 2 and 5.	1) Jane F Arthur, performed some experiments in Figure 5, 10%. 2) Hannah Lee, performed some experiments in Figure 2, 10%. 3) Warwick S Nesbitt, supervision, contribution of tools and reagents, 1%. 4) Robert K Andrews, supervision, contribution of tools and reagents, 1%. 5) Elizabeth E Gardiner, supervision, contribution of tools and reagents, 1%. 6) Justin R Hamilton, design of research, supervision and writing of manuscript, 20%	No  Yes  No  No  No  No
3	Perinatal lethality of Par4-/- mice delivered by primiparous dams reveals	Published	70%. Performed data collection and analysis and writing of manuscript.	1) Justin R Hamilton, design of research, supervision and writing of manuscript, 30%	No

	spontaneous bleeding in mice without platelet thrombin receptor function				
4	Humanizing the protease-activated receptor (PAR) expression profile in mouse platelets by knocking PAR1 into the Par3 locus reveals PAR1 expression is not tolerated in mouse platelets	Published	60%. Writing of manuscript, performed all experiments and data analysis for Figures 1, 3, 4, and 5. Contribution and supervision of experiments and data analysis for Figures 2, 6 and 7.	2) Antonia Christa Paramitha, performed some experiments in Figures 2, 6 and 7, 15%. 3) Mitchell J Moon, technical support, 2.5% 4) Ross A Dickens, supervision and contribution of reagents and tools, 2.5% 5) Justin R Hamilton, design of research, supervision and writing of manuscript, 20%	Yes  Yes  No  No
5	Inhibition of protease-activated receptor 4 (PAR4) with function blocking antibodies provides equivalent antithrombotic activity in the face of the hyper-reactive Thr120 PAR4 variant	Submitted	65%. Writing of manuscript, performed all experiments and data analysis with the exception of Figures 1B and 1C.	1) Claudia Thalmann, performed the experiments in Figure 1B and 1C, 5%. 2) Mark A Sleeman, supervision, contribution of reagents, 10%. 3) Justin R Hamilton, design of research, supervision and writing of manuscript, 20%	No  No  No

I have not renumbered sections of submitted or published papers in order to generate a consistent presentation within the thesis.

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The undersigned hereby certify that the above declaration correctly reflects the nature and extent of the student's and co-authors' contributions to this work. In instances where I am not the responsible author I have consulted with the responsible author to agree on the respective contributions of the authors.

**Main Supervisor signature:**  **Date:** 24/10/2017

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## List of Abbreviations

AA	Arachidonic acid
ACS	Acute coronary syndrome
ADP	Adenosine diphosphate
AMI	Acute myocardial infarction
APTT	Activated partial thromboplastin time
Ca <sup>2+</sup>	Calcium
CHO	Chinese hamster ovary
COX	Cyclooxygenase
DAG	Diacylglycerol
DRG	Dorsal root ganglion
ERK	Extracellular signal-related kinase
FDA	Food and Drug Administration
GEF	Guanine exchange factors
GP	Glycoprotein
GPCR	G-protein coupled receptor
HEK	Human embryonic kidney
MAP	Mitogen-activated protein
MI	Myocardial infarction
MLC	Myosin light chain
MTSP1	Membrane type serine protease 1
PAD	Peripheral artery disease
PAR	Protease-activated receptor
PAR-AP	Protease-activated receptor activating peptide
PI3	Phosphoinositide 3-kinase
PLC	Phospholipase C
PS	Phosphatidylserine
PT	Prothrombin time
TBD	Thrombin binding domain
TIA	Transient ischaemic attack
TLS	Tethered ligand site

TMPRSS2	Transmembrane protease, serine 2
TP	Thromboxane receptors
TT	Thrombin time
TXA2	Thromboxane A2
VWF	von Willebrand factor



# **Chapter 1 - Introduction**

## **1.1 Cardiovascular disease and antithrombotic therapy**

### *1.1.1 There is a significant clinical need for improved antithrombotic therapy*

Cardiovascular disease, manifesting predominantly as ischaemic heart disease and ischaemic stroke, is by far the most common cause of death and disability in the world, accounting for approximately 30% of all deaths<sup>1</sup>. Strikingly, despite increased awareness and improved management, and in contrast to most other high-impact communicable and non-communicable diseases, the rates of cardiovascular disease rose over the past decade<sup>1</sup>. The increasing burden of conditions such as diabetes, obesity, and depression, as well as an overall ageing population, are likely to ensure that cardiovascular disease rates continue to rise into the foreseeable future, with each of these conditions having increased prevalence of cardiovascular-related morbidity and mortality.

Arterial thrombosis precipitates the most prevalent cardiovascular disease manifestations, most notably acute myocardial infarction (AMI), ischaemic stroke, and peripheral artery disease (PAD). Activated platelets are the main cellular component of arterial thrombi. Therefore, current therapies for the prevention of arterial thrombosis are predominantly antiplatelet agents, which prevent platelet activation. The current clinical recommendation for the prevention of primary or secondary cardiovascular events in patients with acute coronary syndrome (ACS) is antiplatelet therapy alone<sup>2</sup>. Yet current standard-of-care treatments have limited efficacy and are tempered by the attendant bleeding risk, meaning that improved antiplatelet approaches are sought.

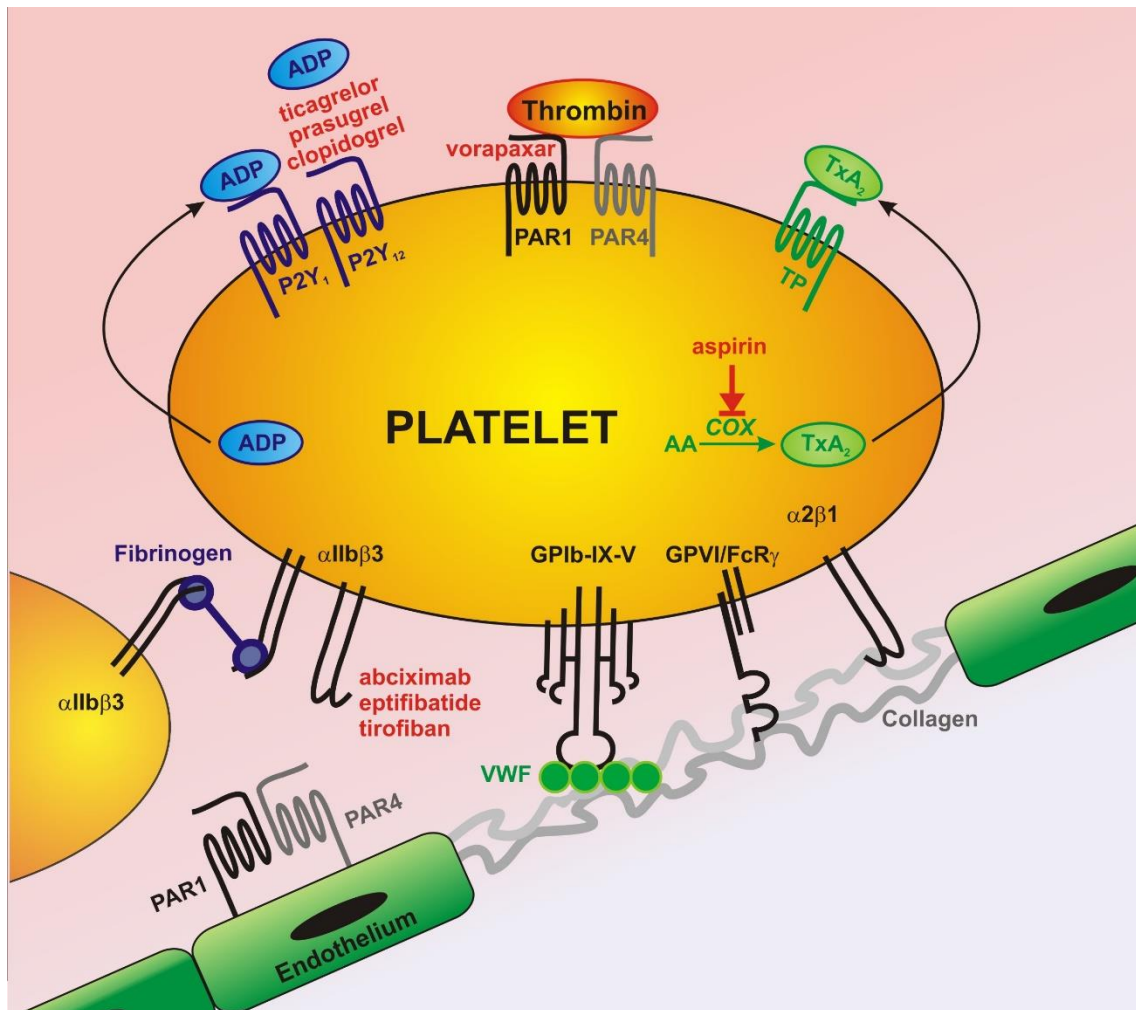
### *1.1.2 Current antiplatelet agents as antithrombotics*

The processes controlling platelet function in the setting of arterial thrombosis have been extensively studied for the purpose of rationalising the development of improved antiplatelet therapies. Platelet behaviour during thrombus formation is generally agreed to involve cell adhesion, activation, and aggregation<sup>3</sup> (Figure 1). Understanding these processes has been highly informative in predicting the success of antiplatelet approaches.

*i) Adhesion:* In response to vascular damage, such as atherosclerotic plaque rupture in the case of ACS, platelets adhere to the damaged vessel wall through a complex between sub-endothelial proteins and cognate receptors on the platelet surface. Rapid initial adhesion is mediated by von Willebrand factor (VWF) binding to the glycoprotein (GP) Ib-IX-V receptor complex<sup>4</sup>, although this interaction does not support stable adhesion and a second adhesive step between collagen and GPVI and/or the integrin  $\alpha_2\beta_1$ , as well as fibrinogen and the integrin  $\alpha_{IIb}\beta_3$ , is required for firm platelet adhesion to the vessel wall<sup>3</sup>.

*ii) Activation:* following adhesion, the captured platelets are activated and undergo a series of morphological and functional changes including shape change, spreading, release of granular contents, and the local generation of thrombin at the platelet surface as an endpoint of coagulation<sup>5</sup>. This platelet activation is primarily driven by the triumvirate of thrombin, ADP, and thromboxane A<sub>2</sub> (TxA<sub>2</sub>).

*iii) Aggregation:* regardless of the adhesion and activation mechanism, the final common result is the activation of the integrin  $\alpha_{IIb}\beta_3$ , which engages fibrinogen (also VWF and fibronectin) to mediate platelet aggregation and ultimately stable thrombus formation, via thrombin-induced fibrin production<sup>5</sup>.



**Figure 1. Activation and adhesion receptors on the platelet.**

In response to vascular damage, platelets rapidly adhere to the vessel wall via the interaction of subendothelial von Willebrand factor (VWF, green spheres) with the platelet glycoprotein (GP) Ib-IX-V complex, and collagen (grey strands) with GPVI and integrin  $\alpha_2\beta_1$ . Adherent platelets become activated, and this activation is enhanced by signalling pathways initiated by: thromboxane receptors (TP) activated by thromboxane A<sub>2</sub> (TxA<sub>2</sub>) generated from arachidonic acid (AA) by cyclooxygenase (COX); P2Y<sub>1</sub> and P2Y<sub>12</sub> receptors activated by ADP; and PAR1 and PAR4 activated by thrombin. Platelet activation culminates in the activation of integrin  $\alpha_{IIb}\beta_3$  which binds fibrinogen (blue dumbbell), VWF and fibronectin (not shown) to mediate platelet aggregation. Current antiplatelet agents (in red) include the P2Y<sub>12</sub> inhibitors clopidogrel, prasugrel and ticagrelor; the COX inhibitor, aspirin;  $\alpha_{IIb}\beta_3$  inhibitors, abciximab, eptifibatide and tirofiban; and the most recent addition, vorapaxar, which inhibits PAR1.

(Source: French et al., 2015, Appendix I)

The current guidelines for treatment of ACS and associated diseases are dual therapy with aspirin (an inhibitor of TxA<sub>2</sub> synthesis) and a P2Y<sub>12</sub> receptor antagonist, such as the thienopyridines, clopidogrel or prasugrel, or the cyclopentyl-triazolo-pyrimidine, ticagrelor (American guidelines tend to favour the use of clopidogrel, whereas European guidelines favour either ticagrelor or prasugrel, with clopidogrel as an alternative)<sup>2</sup>. However, aspirin and clopidogrel prevent just 15 and 17% of lethal cardiovascular events respectively and are only marginally more effective in combination<sup>6-8</sup>. In addition, an increasing number of patients are being reported as resistant to these agents<sup>9, 10</sup>. Large scale clinical trials assessing the efficacy of aspirin and clopidogrel (together and separately) in reducing thrombotic risk in cardiovascular disease (CAPRIE<sup>7</sup> and CHARISMA<sup>8</sup>) have ultimately determined greater efficacy when used in combination, with an associated minor increase in risk of bleeding complications. Antagonists of the major platelet integrin,  $\alpha_{IIb}\beta_3$ , such as tirofiban, eptifibatide, and abciximab, are by far the most potent platelet inhibitors as they inhibit platelet aggregation regardless of the activating pathway. However, these drugs all cause significant bleeding complications<sup>11, 12</sup> due to their disruption of the haemostatic function of platelets also dependent on this pathway. Therefore, existing antiplatelet therapies have limitations in one or both of safety and efficacy, with no current agent (or combination of agents) affording sufficiently potent, safe, and orally active prevention of arterial thrombosis. As a result, improved antiplatelet approaches are required to meet the significant clinical need for the safe and effective prevention of arterial thrombosis.

## **1.2 Targeting protease-activated receptors as an antiplatelet approach**

### *1.2.1 Thrombin-induced platelet activation is important during thrombosis*

Thrombin is a multifunctional allosteric serine protease which is by far the most critical non-cellular component of haemostasis. Thrombin interacts with a series of substrates via its

activate site, located in the centre of the enzyme, and two exosites (exosite I and exosite II) located on either side of the active site. Generation of thrombin occurs via the coagulation cascade – a series of serine protease activation steps which ultimately produce and further amplify thrombin generation. In flowing blood, activated platelets are the main cellular contributors to thrombin amplification, via providing a phospholipid membrane surface for coagulation factor assembly. Once generated, thrombin is able to mediate platelet activation, while also engaging in the proteolysis of fibrinogen to facilitate stable thrombus formation.

The rationale for targeting thrombin-induced platelet activation as an antithrombotic approach has long held appeal. First, thrombin functions at a time and place that is predicted to provide safe and effective antiplatelet activity. As outlined above, thrombin, in combination with ADP and TxA<sub>2</sub>, mediates the platelet activation required for thrombus growth that follows initial cell adhesion. Second, thrombin is the most potent endogenous activator of platelets. This suggests that targeting thrombin-induced platelet activation may provide greater efficacy over existing mechanisms that block the platelet activating functions of TxA<sub>2</sub> (aspirin) and ADP (clopidogrel and co.). Third, existing drugs that block the production (e.g. rivaroxaban) or activity (e.g. dabigatran) of thrombin are effective as antithrombotics. However these agents provide global inhibition of thrombin's actions and, consequently, incur a significant bleeding risk – particularly at the high doses used for the prevention of platelet-rich arterial thrombi. Specifically targeting thrombin-induced platelet activation whilst leaving the other functions of thrombin intact may mitigate the bleeding risk and provide a more selective effect for arterial thrombosis prevention. In combination, these factors provided the impetus to develop inhibitors of platelet thrombin receptors.

### *1.2.2 Thrombin activates platelets via protease-activated receptors*

Platelet responses to thrombin are predominately mediated by protease-activated receptors (PARs)<sup>13, 14</sup>. PARs are G-coupled protein receptors (GPCRs) which are expressed on the surface of numerous cell types. In the cardiovascular system this includes platelets<sup>14-17</sup>, but also leukocytes<sup>18</sup>, vascular endothelial and smooth muscle cells<sup>19, 20</sup>, cardiomyocytes<sup>21</sup>, and cardiac fibroblasts<sup>22</sup>. Humans express four PARs, with PAR1, PAR3 and PAR4 being activated by thrombin<sup>15-17</sup>, and PAR2 by trypsin, tryptase, coagulation factors VIIa and Xa, and membrane-bound serine proteases MTSP1 and TMPRSS2<sup>23-25</sup>. The cleavage-based PAR activation mechanism is unique. PARs are all activated the same way, whereby the protease agonist cleaves the amino-terminus of the receptor to reveal a cryptic neo-amino terminal sequence known as the “tethered ligand”<sup>26</sup>. The newly exposed tethered ligand then activates the receptor by binding intramolecularly to the second extracellular loop<sup>26</sup>. This self-activation prompts the conformational change of the receptor that allows interactions with G proteins of the G<sub>q</sub>, G<sub>12/13</sub> and G<sub>i/z</sub> families and consequent intracellular signalling events<sup>13, 27</sup>. The interactions with G<sub>12/13</sub> drive Rho-dependent cytoskeletal responses involved in platelet shape change<sup>28</sup>, whereas G<sub>q</sub>-mediated signalling facilitates the processes important for platelet granule release, activation of cell surface integrins, and platelet aggregation<sup>29</sup>. G<sub>i/z</sub> proteins are involved in inhibition of adenylyl cyclase, thereby removing a brake on platelet activation, while the G<sub>i</sub> protein family is involved in modifying activities of enzymes such as PI3 kinase<sup>27</sup>.

Human platelets express two thrombin-sensitive PARs, PAR1 and PAR4, and activation of either receptor is capable of inducing platelet activation<sup>14-16</sup>. Thrombin-induced platelet activation initiates platelet shape change, promotes platelet aggregation, and provides the procoagulant lipid surface that facilitates secondary coagulation reactions<sup>13, 30</sup>. Although often described as a dual platelet thrombin receptor system, some distinctions between PAR1 and PAR4 have been shown. PAR1 is a “high affinity” thrombin receptor, as it contains a high

affinity thrombin binding domain (TBD) (K<sup>51</sup>YEPF<sup>55</sup>) which binds exosite I of thrombin and aligns the active site with the specific cleavage site in the receptor<sup>31, 32</sup>. PAR1 is activated by trace amounts of thrombin (sub-nM range) and is responsible for the initial and rapid rise in intracellular calcium (Ca<sup>2+</sup>) induced in platelets by the coagulation protease<sup>33</sup>. However, the PAR1-induced signal is transient and requires additional support from other platelet agonists such as ADP<sup>34</sup>. In contrast to PAR1, the interaction between PAR4 and thrombin occurs primarily at the active site due to the absence of a high affinity TBD in the amino-terminal of PAR4. Despite the lack of interaction between PAR4 and exosite I of thrombin, the receptor does bind at the active site with high affinity due to two optimally-positioned proline residues immediately amino-terminal to the thrombin cleavage site<sup>35</sup>. This interaction facilitates slowed dissociation of thrombin from the receptor and results in a slower but more sustained intracellular signalling profile<sup>35</sup>. As a result of these distinct receptor activation and signalling kinetics, PAR4, in contrast to PAR1, is capable of inducing irreversible platelet aggregation in the absence of additional agonist activation<sup>34</sup>. It has therefore been suggested that PAR1 and PAR4 complement each other during thrombin-induced platelet aggregation. However, historically, the major clinical focus has been on inhibiting PAR1 for antithrombotic activity due to the significantly greater sensitivity of this receptor to thrombin.

### *1.2.3 Mice lacking platelet-thrombin receptor function are protected against thrombosis*

As is the case in many other systems, mouse genetic models and other small animal *in vivo* thrombosis experiments provided pre-clinical proof-of-concept studies. However the interpretation of these studies has always been hampered by the differing platelet PAR profile in these traditional model systems. Of the most commonly used animal models, only non-human primates appear to have an identical platelet PAR profile to humans (PAR1 and PAR4)<sup>36</sup>. In contrast, platelets from mice<sup>37</sup>, rats<sup>38</sup>, and rabbits<sup>39</sup> express PAR3 and PAR4, while platelets from guinea-pigs express PAR1, PAR3 and PAR4<sup>40</sup>. Despite these limitations,

significant insights have been gained from mouse genetic experiments. Although mouse PAR4 functions in an analogous manner to human PAR4, mouse PAR3 is incapable of mediating transmembrane signalling by itself, instead functioning as a cofactor that facilitates cleavage and activation of PAR4 at low thrombin concentrations<sup>41, 42</sup>. As this model predicts, platelets from PAR4<sup>-/-</sup> mice are unresponsive to thrombin<sup>37, 43</sup> and provide a clean genetic model to examine the overall contribution of thrombin-induced platelet activation in (patho)physiology.

The first phenotype reported in PAR4<sup>-/-</sup> mice was protection against thrombosis, with mild associated bleeding<sup>37</sup>. Initially, it was shown that after a ferric chloride-induced injury of mouse mesenteric arterioles, the time to vessel occlusion was prolonged ~ 3 times in PAR4<sup>-/-</sup> compared to wild type mice<sup>37</sup>. Subsequent studies have shown a similar protection against thromboplastin-induced pulmonary embolism<sup>41</sup>, laser-induced endothelial cell ablation in mesenteric arterioles<sup>44</sup> as well as electrolytic<sup>45</sup> and trauma-induced<sup>46</sup> injury of the carotid artery in PAR4<sup>-/-</sup> mice. Bone marrow transplantation studies confirmed that the antithrombotic effects observed in PAR4<sup>-/-</sup> mice are due to deficiency of PAR4, and therefore thrombin signalling, specifically in platelets<sup>43</sup>. Intriguingly, it appears that the protection against thrombosis associated with PAR4 deficiency leaves other haemostatic responses intact<sup>44</sup>. PAR4 deficiency in mice is also associated with a mild bleeding phenotype. This haemostatic effect has been most commonly assessed via tail bleeding time, where PAR4<sup>-/-</sup> mice are consistently shown to have prolonged tail bleeding times compared with wild type mice<sup>37, 41</sup>. Despite this commonly-reported effect upon active limb trauma, there has been no evidence for spontaneous bleeding reported in PAR4<sup>-/-</sup> mice. PAR4 deficiency also has protective benefits in a mouse transient middle cerebral artery occlusion model of stroke, in which PAR4<sup>-/-</sup> mice exhibited lower cerebral infarct volume, improved neurologic and motor function, and reduced blood brain barrier disruption and cerebral oedema, compared with wild type animals<sup>47</sup>. Despite the limitations of using mice as a model of human platelet PAR function,



these early proof-of-concept studies provided insight into the relative importance of thrombin-induced platelet activation in the setting of *in vivo* thrombosis, and in large part drove the development of PAR antagonists as a novel antiplatelet approach.

#### *1.2.4 Considerations of the first clinically approved PAR1 antagonist*

In May 2014 the US FDA approved the first clinical PAR1 antagonist, vorapaxar, for use in the prevention of thrombotic cardiovascular events in patients with a history of myocardial infarction or peripheral artery disease. After showing promising results in phase 2 trials, vorapaxar was assessed in two large scale phase 3 trials in patients with ACS (TRACER<sup>48</sup>) and stable atherosclerosis (TRA 2°P – TIMI 50<sup>49</sup>). The overall findings of these trials showed that vorapaxar, when combined with standard-of-care therapy, reduced the risk of cardiovascular events at the cost of increased bleeding<sup>48, 49</sup>. However, several key issues and a narrow therapeutic window have limited the clinical utility of vorapaxar.

The major concerns associated with vorapaxar use were increased rates of intracranial bleeding and poor compatibility with standard-of-care antiplatelet agents. During the TRACER trial, intracranial haemorrhage increased more than 5-fold in the vorapaxar group<sup>48</sup>. This effect was especially noted in patients with a prior history of stroke or transient ischaemic attack (TIA) and increased incrementally over time<sup>48</sup>. As a result, patients with a history of stroke or TIA were removed from the TRAP 2°P – TIMI 50 trial, however intracranial haemorrhage was still increased two-fold in the vorapaxar group<sup>49</sup>. The trials showed that these bleeding rates increased in patients receiving vorapaxar in combination with a thienopyridine compared to those receiving a thienopyridine alone<sup>48</sup>. Further, in patients receiving high dose aspirin (> 300 mg daily at both baseline and time of discharge), a consistent, albeit non-statistically significant, trend toward higher bleeding and ischaemic outcomes was recorded<sup>50</sup>. It remains unknown whether the bleeding complications observed in these trials were due to PAR1 inhibition per se, the concurrent use of three antiplatelet drugs, or to a particular drug

combination. Regardless, overall, in patients with no history of stroke or TIA, and with a body weight above 60 kg, the data from TRA 2°P – TIMI 50 translated into 6 fewer cardiovascular deaths at the cost of two intracranial haemorrhages for every 1000 patients treated with vorapaxar<sup>51</sup>.

Despite these setbacks, PAR1 inhibition did demonstrate consistent efficacy in certain pathological conditions where thrombin-mediated platelet activation is known to play a significant role. For example, PAR1 inhibition demonstrated a consistent reduction in the rate of type 1 (spontaneous) MI in vorapaxar-treated patients across the phase 2 and phase 3 studies<sup>52</sup>. This is perhaps unsurprising given the well-known role of thrombin generation in acute MI, particularly in patients with a background of unstable angina and/or coronary artery disease<sup>53-55</sup>. Vorapaxar was also shown to reduce the complications associated with PAD, notably limb ischemia and the requirement for peripheral vascularisation<sup>56</sup>. One explanation for this effect is recent work which has shown that thrombin-dependent platelet activation (via PARs) is more prevalent at lower blood shear rates<sup>46</sup>, presumably due to limited assembly of blood borne coagulation factors on the surface of activated platelets at higher blood flow/shear rates. Understanding the relative contribution of thrombin-mediated platelet activation to thrombus formation in particular pathologies will likely be of significant use in predicting the clinical success of PAR inhibitors. Further, the limitations of PAR1 antagonism (i.e. increased bleeding, poor compatibility with other agents) have led researchers to revisit PAR4 antagonism as a novel antithrombotic approach.

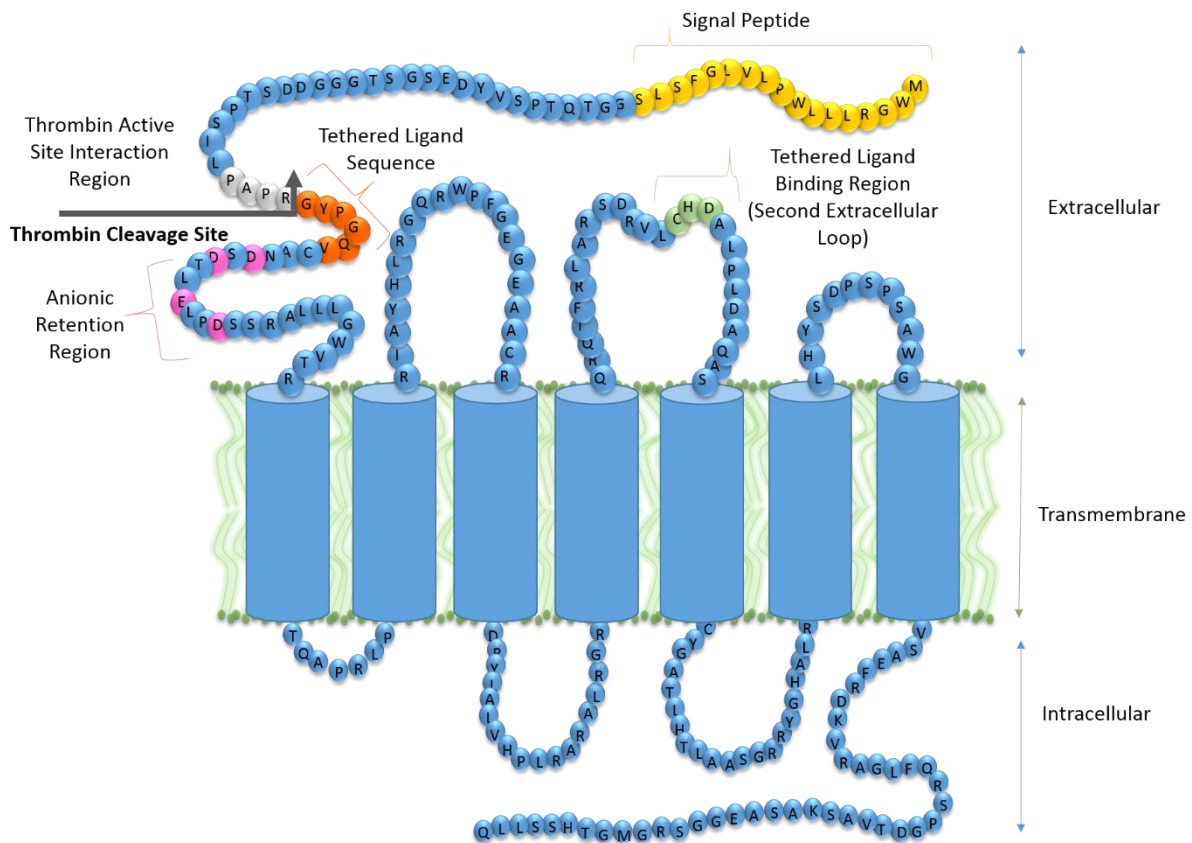
### **1.3 Is selectively targeting PAR4 a viable antiplatelet strategy?**

#### *1.3.1 PAR4 structure and biology*

Despite being traditionally thought of as a “back-up” receptor to PAR1, PAR4 has several structural and biologically distinct characteristics to the rest of the PAR family. PAR4 was first

cloned in 1998<sup>14, 16</sup> and is the most recently cloned member of the PAR family. It is a 300 amino acid seven transmembrane-spanning domain GPCR (Figure 2) that retains most of the core features of the other PARs despite its genetic departure from the family<sup>14, 16</sup>. The human PAR4 gene is remarkably smaller than those of the other PARs, and it resides at a distinct location on chromosome 19p12, with the genes for PARs 1-3 located in tandem on chromosome 5q13<sup>16, 57</sup>. This genetic divergence of PAR4 from the other PARs is thought to have arisen from a remote gene duplication and subsequent translocation event which gave rise to ancestral PAR4 and PAR1/2/3 genes<sup>57</sup>. More recently, two gene duplication events have occurred to separate PAR1 and PAR2 from PAR3; and then PAR1 from PAR2<sup>57</sup>. These genetic differences may underlie some of the key differences in receptor structure and function within the PAR family.

The N-terminus of PAR4 contains a hydrophobic signal peptide sequence, with a signal peptidase cleavage site present at S<sup>17</sup>/G<sup>18</sup>. The extracellular amino-terminus also contains a serine protease cleavage site at R<sup>47</sup>/G<sup>48</sup> that is essential for proteolytic receptor activation. Mutation of the serine protease cleavage site (R<sup>47</sup> → A) renders the receptor completely unresponsive to proteolytic activation. Site-specific receptor cleavage unmasks a cryptic tethered ligand sequence (GYPGQV)<sup>16</sup> which then binds intra-molecularly to a defined region in the second extracellular loop of the receptor (Figure 2), resulting in conformational change of the receptor and subsequent coupling to intracellular effectors, as per other GPCR family members. While this proteolytic activation mechanism is common amongst PARs, there are several notable differences in the structure of PAR4 versus the other receptors of the family. First, both the extracellular amino terminus and intracellular carboxy terminus have little sequence similarity to the corresponding regions of other PARs<sup>16</sup>. Second, the tethered ligand binding site of PAR4 contains only three core amino acids (CHD; Figure 2) of the consensus



**Figure 2. PAR4 structure.** Depicted is a cartoon of the proposed structure of the 300 amino acid GPCR, PAR4. Site-specific proteolytic cleavage of the receptor's amino-terminus (thrombin cleavage site, R<sup>47</sup>/G<sup>48</sup>; black arrow) reveals a neo-amino-terminus (tethered ligand sequence; G<sup>48</sup>YPGV; orange) which binds intra-molecularly to the second extracellular loop of the receptor (tethered ligand binding region; green). PAR4 contains two regions for enhanced thrombin interaction – the anionic retention region (pink) which interacts with thrombin's exosite I; and the PAPR sequence (grey) which binds at the active site of thrombin with high affinity.

(Source: French and Hamilton, 2016, Appendix II)

sequence conserved in PARs 1-3 (ITTCHDV)<sup>16</sup>. PAR4 also lacks the high affinity thrombin binding domain that is present in the other two thrombin receptors, PAR1 and PAR3<sup>14, 17, 58, 59</sup>.

PAR4 is broadly expressed, with gene expression by Northern blot most readily detected in the lungs, pancreas, thyroid, testis and small intestine<sup>14, 16</sup>. Moderate expression has also been detected in placenta, skeletal muscle, lymph nodes, adrenal gland, prostate, uterus and colon<sup>14, 16</sup>. In the nervous system, PAR4 protein and mRNA have been detected in rat dorsal root ganglion (DRG) non-neuronal cells (conversely, PARs 1-3 were detected in DRG neurons)<sup>60</sup>. Despite this vast expression profile of PAR4, most research has focussed on the physiological functions of this receptor in cardiovascular and inflammatory settings. In humans, key vascular cell types which express PAR4 include platelets<sup>14, 16</sup>, leukocytes<sup>61</sup>, endothelial cells<sup>62</sup>, and smooth muscle cells<sup>63-65</sup>. In addition to the well-known pro-thrombotic and pro-inflammatory actions of PAR4-activating proteases, perhaps one reason for this is that several important animal models, including mice, rats, guinea pigs, rabbits, dogs and monkeys have similar PAR4 expression to humans throughout the vasculature (platelets are one example of such conserved PAR4 expression). Despite this relatively conserved expression of PAR4 in vascular cells, there is some evidence to suggest that PAR4 may function differently between species. Specifically, it is known that PAR4 contributes to thrombin responses of mouse endothelial cells<sup>62</sup>, and, that PAR4 activation causes endothelium-dependent relaxation of rat aorta<sup>63</sup>. However, in human artery preparations as well as lung fibroblasts, these responses are only elicited when treated with additional inflammatory mediators<sup>64, 66</sup>. This distinct regulation of PAR4 expression and function between species remains an important consideration for interpretation of animal studies.

### *1.3.2 PAR4 activation and signalling: distinctions from PAR1*

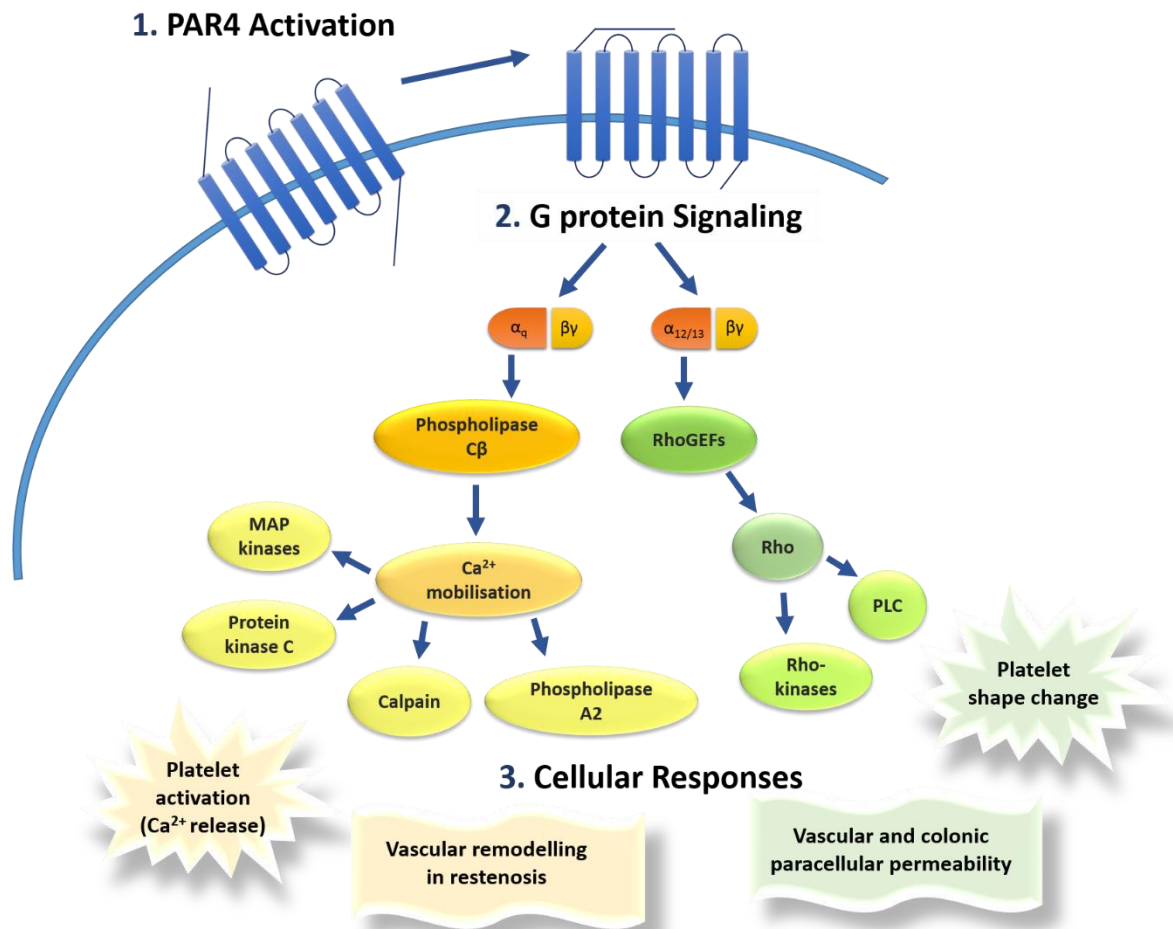
The setbacks of clinical PAR1 antagonists have sparked research interest into whether selective PAR4 antagonism may be an alternative approach to inhibiting thrombin-induced platelet

activation. Indeed, a significant line of evidence suggests that PAR4 signals differently and independently of PAR1 activation, and therefore potentially regulates distinct platelet activation mechanisms.

The serine proteases capable of cleaving PAR4 are generally key regulators of coagulation (e.g. thrombin and coagulation factor Xa) and/or inflammation (e.g. trypsin and cathepsin G released from neutrophils, and bacterial proteases such as gingipains)<sup>14, 16, 37, 67</sup>. Additional serine proteases that are specific for arginine or lysine cleavage include coagulation factors VIIa, IXa, Xa, XIa, urokinase, and plasmin; however, none of these have significant activity on PAR4, with the exception of factor Xa which showed small effects at non-physiological concentrations<sup>16</sup>. Initial studies indicated that PAR4 is activated by thrombin and trypsin at similar concentrations, with an EC<sub>50</sub> of ~ 5 nM each<sup>16</sup> – significantly higher than the EC<sub>50</sub> for thrombin at either PAR1 or PAR3 (~ 0.2 nM). The comparatively lower affinity of PAR4 for thrombin is a reflection of differences in the macromolecular association between the receptor and enzyme. As previously described, PAR1 is a high affinity thrombin receptor due to the presence of a thrombin binding domain (TBD) in its N-terminal exodomain<sup>59</sup>. The TBD sequence, K<sup>51</sup>YEPF<sup>55</sup>, interacts with exosite I of thrombin and upon binding causes significant allosteric effects essential for rapid association of thrombin<sup>33</sup>. PAR4 does not contain this TBD<sup>16</sup>. In fact, evidence indicates that PAR4 appears to have only limited interaction with exosite I of thrombin, since  $\gamma$ -thrombin (which lacks a functional exosite I) activates PAR4 as effectively as  $\alpha$ -thrombin<sup>16</sup> and mutations in the exosite I of thrombin have significantly less effect on cleavage of PAR4 than of PAR1<sup>68</sup>. Instead, PAR4 primarily interacts with the active site of thrombin via two optimally positioned proline residues (P<sup>44</sup> and P<sup>46</sup>) in the receptor<sup>35</sup>, just upstream of the thrombin cleavage site at R<sup>47</sup>/G<sup>48</sup>. A possible role for L<sup>43</sup> in the thrombin-PAR4 interaction has also been identified, and believed to be in facilitating high-affinity binding of the PAR4 amino-terminal to thrombin's active site<sup>35</sup>. PAR4 also contains an anionic

cluster, D<sup>57</sup>...D<sup>59</sup>...E<sup>62</sup>...D<sup>65</sup>, just C-terminal of the thrombin cleavage site that is thought to interact with cationic residues that border exosite I of thrombin<sup>68</sup> and slow the dissociation rate of thrombin from PAR4<sup>35</sup>. This is reflected in the more sustained intracellular signals elicited by PAR4 activation compared with the more transient signals in response to PAR1 activation<sup>69</sup>. These differences in receptor structure and activation kinetics between PAR1 and PAR4 indicate the two receptors have the potential to initiate distinct intracellular signalling events.

Cleavage and activation of PARs prompts a conformational change in the receptor that allows G protein coupling and initiates multiple intracellular signalling events. G<sub>q</sub>, G<sub>i</sub>, and G<sub>12/13</sub> have all been demonstrated to mediate signals in response to activation of PARs, with PAR4 specifically being shown to couple to both G<sub>q</sub> and G<sub>12/13</sub> family proteins<sup>13, 70</sup>, but not G<sub>i/o</sub><sup>71</sup> (Figure 3). PAR4 coupling to G<sub>12/13</sub> initiates binding of RhoGEFs (guanine-nucleotide exchange factors that activate Rho) to the  $\alpha$ -subunit. Consequent Rho activation induces a series of Rho-dependent cytoskeletal responses and PLC activation (Figure 3). For example, in platelets, activation of Rho-kinase dependent cytoskeletal responses via such G<sub>12/13</sub>-mediated PAR4 signalling triggers platelet shape change<sup>13, 72, 73</sup> – one of the key initial events in platelet aggregation during thrombosis. Evidence also exists for PAR4 activation eliciting a prolonged signal via G<sub>12/13</sub>, which is thought to involve the regulation of MLC phosphorylation and RhoA<sup>74, 75</sup>. PAR4 coupling to G<sub>q</sub> causes intracellular Ca<sup>2+</sup> mobilisation via activation of phospholipase C $\beta$  and consequent phosphoinositide hydrolysis (Figure 3). PAR4-induced Ca<sup>2+</sup> mobilisation promotes the activity of several Ca<sup>2+</sup>-regulated kinases and phosphatases (e.g. MAP kinases, protein kinase Cs, phospholipase A2, and calpain)<sup>13</sup>, with wide-ranging effects. For example, PAR4 activation mediates several key platelet responses that are induced by G<sub>q</sub>-dependent Ca<sup>2+</sup> mobilisation, including the secretion of platelet storage granules and integrin activation – both of which are critical for effective platelet aggregation<sup>13, 76</sup>.



**Figure 3. PAR4 signalling.** Following cleavage and activation of the receptor, PAR4 signals via coupling to G<sub>q</sub> and/or G<sub>12/13</sub> family members. The most well characterised signalling downstream of G<sub>q</sub> is via phospholipase C $\beta$ -mediated phosphoinositide hydrolysis and resultant intracellular Ca<sup>2+</sup> mobilisation. This promotes the activity of several Ca<sup>2+</sup>-regulated kinases and phosphatases (e.g. MAP kinases, protein kinase C, phospholipase A2, and calpain) which underlie PAR4-induced cellular responses such as platelet activation and vascular remodelling. The most well characterised signalling downstream of PAR4-G<sub>12/13</sub> coupling is via RhoGEF and consequent Rho activation, which underlie PAR4-induced cellular responses such as cytoskeletal responses in platelets and vascular smooth muscle and endothelial cells.

(Source: French and Hamilton, 2016, Appendix II)



As indicated above, one key difference between PAR4 and PAR1 is the kinetics of receptor activation and signalling. Regarding G<sub>q</sub>-mediated signalling, it is known that PAR1 activation stimulates a rapid burst of intracellular Ca<sup>2+</sup> mobilisation, whereas PAR4 activation elicits a slower rise which is much more sustained over time<sup>69</sup>. This is likely a result of the slower cleavage of PAR4 allowing prolonged G protein signalling. Given that mice deficient in G<sub>q</sub> lack thrombin-dependent IP<sub>3</sub> and Ca<sup>2+</sup> responses<sup>29, 77</sup> and that PAR4 activation stimulates ongoing IP<sub>3</sub> and DAG, differences in Ca<sup>2+</sup> mobilisation are likely a result of prolonged G<sub>q</sub> signalling through PAR4<sup>34, 78</sup>. It is thought that this may be important for ongoing cell signalling under conditions of prolonged agonist exposure. In platelets, PAR1-mediated Ca<sup>2+</sup> signalling undergoes rapid desensitisation but can be rescued by subsequent PAR4 activation<sup>79</sup>. Receptor desensitisation is of particular importance for PARs due to the irreversible, cleavage-based, endogenous activation of these receptors, with continued responsiveness to agonists requiring new receptors to be trafficked to the cell surface. After receptor cleavage and activation, PARs are rapidly internalised on a phosphorylation signal in regions of the C-terminus of the receptor<sup>80</sup>. Yet PAR4 has a shorter C-terminus than PAR1 and does not have many of the phosphorylation sites shown to be necessary for desensitisation of these receptors<sup>81</sup>. As a result, agonist-triggered phosphorylation and consequent receptor internalisation is significantly slower for PAR4 than for PAR1<sup>81</sup>, providing a further mechanism for the comparatively prolonged intracellular signalling downstream of PAR4 activation.

### *1.3.3 PAR4 antagonists*

The unique structure and activation mechanism of PARs has long posed a problem for the development of inhibitors. In particular, activation by proteolytic cleavage is a highly efficient and irreversible system, while the tethered ligand-based activation mechanism requires antagonism of the binding of an agonist intrinsic to the receptor and presumably with

considerable steric advantage. These issues have hindered the development of PAR4 antagonists, although recent efforts have begun to overcome these challenges, with several distinct antagonist classes emerging (Table 1). This diverse set of PAR4 antagonists developed over recent years have been invaluable for investigations into the physiological roles of PAR4, and have also served to promote the idea of targeting this receptor as a novel therapeutic approach. However, there remains a lack of PAR4 antagonists that are able to demonstrate the necessary levels of specificity and efficacy required for clinical evaluation.

The earlier classes of PAR4 antagonists include peptidomimetics, pepducins and function-blocking antibodies. These inhibitors have been predominately research-focused. The first approach taken to identify potential inhibitors of PAR4, was the peptidomimetic approach which resulted in the generation of numerous peptide analogues based on the tethered ligand sequences of human, mouse, and rat PAR4. The resulting compound, tc-YPGKF-NH<sub>2</sub>, appeared to bind but not activate PAR4 and was shown to abolish PAR4-activating peptide (PAR4-AP)-induced aggregation of rat platelets and significantly reduce thrombin-induced platelet aggregation, at very high concentrations (400  $\mu$ M)<sup>38</sup>. tc-YPGKF-NH<sub>2</sub> has also been reported to inhibit thrombin-induced platelet aggregation in human platelets<sup>82</sup>, although there is limited other evidence of this agent inhibiting human PAR4<sup>83</sup>. Pepducins are a distinct class of PAR4 inhibitors that mimic the region of the receptor that binds G proteins and essentially work by absorbing the interactions between the receptor and effector G protein<sup>84</sup>. The anti-PAR4 pepducins, P4pal-10 (pal-SGRRYGHALR-NH<sub>2</sub>) and P4pal-i1<sup>85</sup>, have been shown to inhibit ~85% of thrombin-induced aggregation of both human and mouse platelets. However, the specificity of these remain debatable, as they reportedly display a level of cross-reactivity with PAR1-AP-induced platelet activation<sup>92</sup> as well as collagen and TxA<sub>2</sub><sup>95</sup>. Several function-blocking anti-PAR4 antibodies have also been developed and proven to be useful PAR4 antagonists. These include rabbit polyclonal antibodies against the thrombin cleavage site of

**Table 1. Properties of experimental PAR4 antagonists.**

<i>Class</i>	<b>Compound</b>	<b>Target</b>	<b>IC<sub>50</sub><sup>†</sup></b>	<b>Species</b>	<b>Cell Type</b>	<b>Reference</b>
<i>Peptidomimetics</i>	tc-YPGKF-NH <sub>2</sub>	TLS binding site	100 µM	Rat, human	Platelets	Hollenberg, 2001 <sup>38</sup> Hollenberg, 2004 <sup>83</sup> Ma, 2005 <sup>82</sup>
	YD-3	TLS binding site	28 µM	Rabbit, rat, human	Platelets, smooth muscle cells	Wu, 2002 <sup>86</sup> Wu, 2003 <sup>87</sup> Peng, 2004 <sup>88</sup>
<i>Small molecules</i>	19, 25, 31	TLS binding site	>36 µM	Human	Platelets	Huang, 2006 <sup>89</sup>
	ML354	TLS binding site	140 nM <sup>#</sup>	Human	Platelets	Young, 2013 <sup>90</sup> Wen, 2014 <sup>91</sup>
<i>Pepducins</i>	P4pal-10	Third-intracellular loop	1 µM	Mouse, human	Platelets	Covic, 2002a <sup>84</sup> Covic, 2002b <sup>92</sup>
	P4pal-i1	First-intracellular loop	5 µmol/L*	Guinea pigs, human	Platelets	Leger, 2006 <sup>85</sup>
<i>Function-blocking antibodies</i>	Rabbit polyclonal	Thrombin cleavage site	1 mg/mL*	Human, rat	Platelets, fibroblasts	Kahn, 1999 <sup>30</sup>
	Rabbit polyclonal (Can12)	Anionic region	10 ng/mL <sup>-1</sup>	Mouse, human	Platelets	Mumaw, 2014 <sup>93</sup>
	Mouse monoclonal (14H6, 5F10)	Thrombin cleavage site	50 µg/mL* partial inhibition	Human	Platelets, HEK-293s	Mumaw, 2015 <sup>94</sup>

TLS = tethered ligand binding site

<sup>†</sup> IC<sub>50</sub> values correspond to inhibition of thrombin-induced PAR4 activation

<sup>#</sup> IC<sub>50</sub> values correspond to inhibition of PAR4 agonist peptide-induced activation

\*Values are concentration reported to inhibit thrombin-induced PAR4 activation

(Source: Adapted from French and Hamilton, 2017, Appendix III)

PAR4<sup>30</sup> and anionic region of the receptor (C<sup>54</sup>ANDSDTLTLPD), just downstream of the thrombin cleavage site<sup>93</sup>. However, despite being valuable research tools, these antibodies have also struggled to demonstrate efficacy<sup>30</sup> and specificity<sup>93</sup>.

To date, the most widely used and desired inhibitor class are the small molecule PAR4 antagonists which have been developed via extensive screening of various heterocyclic structures. The first compound identified to selectively inhibit PAR4 was the indazole derivative YD-3 [1-benzyl-3(ethoxycarbonylphenyl)-indazole]<sup>86, 96</sup>. YD-3 has been shown to inhibit thrombin-induced platelet aggregation in rabbits with an IC<sub>50</sub> of 28  $\mu$ M; however in humans, it was shown to only partially inhibit platelet aggregation in response to thrombin concentrations lower than 0.5 nM<sup>86, 87</sup>. Several studies have aimed to increase the efficacy of YD-3, and have produced several derivatives<sup>89, 97</sup>, including the *N*<sup>2</sup>-(substituted benzyl)-3-(4-methylphenyl)-2*H*-indazoles, compounds 19, 25, and 31<sup>89</sup>, and ML354, a substituted indole derivative<sup>90, 91</sup>. However, these compounds have been shown to inhibit PAR1-AP induced activation with an IC<sub>50</sub> of 10  $\mu$ M, and demonstrate limited efficacy in inhibiting thrombin-induced PAR4 activation. Despite the challenges associated with developing suitable compounds for both research and therapeutic use, there remains a strong interest in continuing development of PAR4 antagonists.

### 1.3.3 Targeting PAR4 as an antithrombotic approach

During the course of this thesis, a number of major advances were made regarding the clinical significance of platelet PAR4 and the utility of PAR4 antagonists during arterial thrombosis. Firstly, the pharmaceutical sector developed a clinically viable small molecule PAR4 antagonist. In a recent study by Wong and colleagues<sup>98</sup>, a library of over 1 million compounds was screened to identify a lead candidate that was then subject to iterative rounds of medicinal chemistry and testing to result in BMS-986120 – a potent and selective PAR4 antagonist with impressive oral bioavailability and antithrombotic efficacy (Table 2). In contrast to previously

generated antagonists, BMS-986120 demonstrated specificity in human platelets, and effectively suppressed platelet aggregation in response to thrombin with concomitant PAR1 inhibition. Further, binding studies revealed that BMS-986120 is a high affinity and reversible binder of PAR4, which translated to normalised aggregation 24 h after a single dose of 0.2 mg/kg. Most importantly though, BMS-986120 appeared to provide an impressive therapeutic window, with a single oral dose of BMS-986120 providing marked antithrombotic effects and a low bleeding profile in a series of *in vivo* models in the cynomolgus monkey. Importantly, when compared directly with clopidogrel, at doses of these two agents that caused equivalent antithrombotic effects, markedly more bleeding was observed with clopidogrel compared with BMS-986120. This work indicates that PAR4 antagonism exhibits a markedly improved therapeutic window over at least one of the standard antiplatelet drugs.

The second major advance to occur during this thesis regarding the clinical utility of targeting PAR4 for arterial thrombosis prevention was the identification of a sequence variant in PAR4 that affects the receptor's pharmacology<sup>99-101</sup>. Specifically, platelets from healthy North American black subjects were shown to be hyper-responsive to PAR4 activation when compared with platelets from non-blacks. This increased sensitivity persisted in the face of PAR1 antagonism (vorapaxar) or aspirin treatment and led to the identification of a genetic variant of PAR4, Ala<sup>120</sup>Thr, which is associated with greater PAR4 reactivity and occurs at a much higher frequency in blacks versus non-blacks. This Thr<sup>120</sup> PAR4 variant, expressed in 20 – 80% of people depending on the population, markedly impacts PAR4 pharmacology by rendering the receptor hyper-sensitive to agonists and hypo-sensitive to antagonists<sup>100, 101</sup>. The mechanism behind this change in PAR4 pharmacology remains unknown, as does whether all PAR4 antagonists will be similarly affected and whether this sequence variant is associated with poorer cardiovascular outcomes. Studies directly addressing these points will be critical

**Table 2. Pharmacological characteristics of a leading PAR4 antagonist, BMS-986120.**

	<i>Cell</i>	<i>Parameter measured</i>	<i>BMS-986120</i>
<i>Binding studies</i>	PAR4-HEK293T membranes	K <sub>D</sub>	0.098 nM
		K <sub>on</sub>	0.12 nM
		K <sub>off</sub>	0.0082 nM
<i>Selectivity/specificity studies</i>	HEK293-PAR4 + PAR4-AP	Ca <sup>2+</sup> mobilisation (PAR4-AP)	0.56 nM
	HEK293-PAR1 + PAR1-AP	Ca <sup>2+</sup> mobilisation (PAR1-AP)	>5000 nM
	CHO-PAR2 + PAR2-AP	Ca <sup>2+</sup> mobilisation (PAR2-AP)	>42000 nM
<i>Inhibition of PAR4 signaling</i>	PAR4-HEK293T	G-protein activation	Gα11 – 3.4 nM Gαq – 3.9 nM Gα14 – 31 nM
		β-arrestin 2 recruitment	7.2 nM
		ERK 1/2 activation	47 nM
<i>Platelet aggregation</i>	Platelet-rich plasma (human)	γ-thrombin	7.3 nM
	Whole blood (human)	PAR4-AP	9.5 nM
	Whole blood (monkey; <i>ex vivo</i> )	PAR4-AP	2.1 nM
		PAR1-AP	No effect (1 mg/kg)
		Collagen	No effect (1 mg/kg)
		ADP	No effect (1 mg/kg)
<i>Coagulation studies</i>	Whole blood (monkey; <i>ex vivo</i> )	APTT	No effect (0.2-1 mg/kg)
		PT	No effect (0.2-1 mg/kg)
		TT	No effect (0.2-1 mg/kg)
<i>In vivo thrombosis</i>	Whole blood (monkey)	Occlusion time	0.2mg/kg - ~75 min 0.5mg/kg - ~75 min 1mg/kg – did not occlude
		Thrombus weight (% reduction from vehicle)	0.2mg/kg - -36% 0.5mg/kg - -50% 1mg/kg – -82%
<i>In vivo haemostasis</i>	Whole blood (monkey)	Kidney bleeding time (fold above vehicle control)	0.2mg/kg – 1.4X 0.5mg/kg – 1.9X 1mg/kg – 2.2X
		Mesenteric bleeding time	0.2mg/kg – 1.4X 0.5mg/kg – 1.7X 1mg/kg – 1.8X

PAR-AP = protease-activated receptor-activating peptide; HEK = human embryonic kidney; CHO = Chinese hamster ovary; ERK = Extracellular signal-related kinase; ADP = adenosine diphosphate; APTT = activated partial thromboplastin time; prothrombin time; TT = thrombin time.

(Source: Adapted from French and Hamilton, 2017, Appendix III)

in determining the approach to PAR4 antagonism that will afford consistent antithrombotic benefit across the population.

#### **1.4 Aims of this thesis**

As discussed above, thrombin-induced platelet activation is a key driver in the formation of the arterial thrombi that cause two of the most lethal manifestations of cardiovascular disease – acute myocardial infarction and occlusive stroke. However, there remains much to learn regarding how to best manipulate thrombin signalling in platelets for safe and effective antithrombotic therapy. Despite known differences in receptor structure and signalling, the specific function of PAR4 during platelet activation and thrombus formation in humans has gone largely unstudied, in large part due to several crucial limitations hindering research into PARs. The purpose of this project was to examine whether targeting PAR4 is a valid antithrombotic approach by examining the relative roles of platelet PAR1 vs PAR4 in the setting of thrombosis, and determining whether PAR4 antagonism is an effective *and* safe strategy for the prevention of arterial thrombosis. Specifically, this thesis aimed to:

- 1: Determine the relative roles of PAR1 and PAR4 on human platelets in the setting of thrombosis (Chapter 2).
- 2: Assess the safety and efficacy of PAR4 inhibition *in vivo* using animal models (Chapters 3 and 4).
- 3: Examine the utility of PAR4 antagonism for the prevention of human thrombosis (Chapter 5).

Sections of this chapter have been published as:

**French SL**, Arthur JF, Tran HA, Hamilton JR. Approval of the first protease-activated receptor antagonist: Rationale, development, significance, and considerations of a novel anti-platelet agent. *Blood Reviews*, 2015, 29 (3):179-89. (See Appendix I)

**French SL**, Hamilton JR. Protease-activated receptor 4: from structure to function and back again. *British Journal of Pharmacology*, 2016, 173:2952-65. (See Appendix II)

**French SL** and Hamilton JR. (2017) Drugs targeting PAR4 improve the antithrombotic therapeutic window. *Annals of Translational Medicine*, doi: 10.21037/atm.2017.09.10 (See Appendix III)



## **Chapter 2 – Inhibition of protease-activated receptor 4 impairs platelet procoagulant activity during thrombus formation in human blood**

### **2.1 Introduction**

The overall goal of this thesis was to examine the validity and utility of blocking PAR4 as a novel antithrombotic approach. As a first step, the specific functions of PAR4 on human platelets during thrombosis were determined. At the time of these studies, suitably specific and effective inhibitors of PAR4 were not freely available. Therefore this thesis began by developing a function-blocking anti-PAR4 antibody and then using it to probe for platelet activation events reliant on PAR4.

Defining the relative functions of the two platelet PARs is of patent interest for any rationalisation of PAR4 as an antiplatelet drug target. Whether PAR4 performs distinct functions to PAR1 on human platelets or simply acts in a redundant manner as a ‘back-up’ receptor has long been debated, although recent studies suggest the former alternative is more likely. As outlined in Chapter 1, one major distinction between the two platelet PARs relates to the kinetics of intracellular signalling, where PAR4 activation induces a slower and more sustained intracellular  $\text{Ca}^{2+}$  signal compared with PAR1 activation. This temporal difference in  $\text{Ca}^{2+}$  signalling was recently shown to prolong the thrombin-PAR4 interaction and facilitate persistent receptor activation. However, the cellular consequences of such sustained platelet activation downstream of PAR4 have not yet been completely characterised – most likely due to the limited availability of appropriate PAR4 antagonists required for such studies. The studies in this Chapter developed such an antagonist and then went on to directly address this question.

The findings from this Chapter demonstrate that thrombin-dependent platelet procoagulant activity – and consequent thrombin generation and fibrin formation – is mediated

predominantly by PAR4 and is largely independent of PAR1 in the setting of thrombus formation.

These findings suggest that PAR1 and PAR4 have distinct functions in platelets, and that the function of PAR4 may be important for thrombus formation and/or stabilisation. These findings suggest blocking PAR4 function may have utility in the prevention of arterial thrombosis and help rationalise the development of PAR4 antagonists for this purpose.

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## IN FOCUS

# Inhibition of protease-activated receptor 4 impairs platelet procoagulant activity during thrombus formation in human blood

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See also Lindahl TL, Macwan AS, Ramström S. PAR4 is more important than PAR1 for the thrombin-induced procoagulant effect on platelets. This issue, pp 1639–41.

## Essentials

- The platelet thrombin receptor, PAR4, is an emerging anti-thrombotic drug target.
- We examined the anti-platelet & anti-thrombotic effects of PAR4 inhibition in human blood.
- PAR4 inhibition impaired platelet procoagulant activity in isolated cells and during thrombosis.
- Our study shows PAR4 is required for platelet procoagulant function & thrombosis in human blood.

**Summary.** *Background:* Thrombin-induced platelet activation is important for arterial thrombosis. Thrombin activates human platelets predominantly via protease-activated receptor (PAR)1 and PAR4. PAR1 has higher affinity for thrombin, and the first PAR1 antagonist, vorapaxar, was recently approved for use as an antiplatelet agent. However, vorapaxar is contraindicated in a significant number of patients, owing to adverse bleeding events. Consequently, there is renewed interest in the role of platelet PAR4 in the setting of thrombus formation. *Objectives:* To determine the specific antiplatelet effects of inhibiting PAR4 function during thrombus formation in human whole blood. *Methods and Results:* We developed a rabbit polyclonal antibody against the thrombin cleavage site of PAR4, and showed it to be a highly specific

inhibitor of PAR4-mediated platelet function. This function-blocking anti-PAR4 antibody was used to probe for PAR4-dependent platelet functions in human isolated platelets in the absence and presence of concomitant PAR1 inhibition. The anti-PAR4 antibody alone was sufficient to abolish the sustained elevation of cytosolic calcium level and consequent phosphatidylserine exposure induced by thrombin, but did not significantly inhibit integrin  $\alpha_{IIb}\beta_3$  activation,  $\alpha$ -granule secretion, or aggregation. In accord with these *in vitro* experiments on isolated platelets, selective inhibition of PAR4, but not of PAR1, impaired thrombin activity (fluorescence resonance energy transfer-based thrombin sensor) and fibrin formation (anti-fibrin antibody) in an *ex vivo* whole blood flow thrombosis assay. *Conclusions:* These findings demonstrate that PAR4 is required for platelet procoagulant function during thrombus formation in human blood, and suggest PAR4 inhibition as a potential target for the prevention of arterial thrombosis.

**Keywords:** antiplatelet drugs; platelets; protease-activated receptors; thrombin; thrombosis.

## Introduction

Activated platelets are the key cellular components of arterial thrombosis – the most common cause of death and disability in the world [1,2]. Platelets are activated by a combination of endogenous agonists that trigger platelet aggregation and the promotion of coagulation, which together facilitate pathologic thrombus formation. Antiplatelet drugs therefore constitute the main pharmacotherapy for the prevention of arterial thrombosis. However, despite an array of such agents, limitations in safety and/or efficacy necessitate the rationalization of new antiplatelet drug targets.

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Thrombin is the most potent known activator of human platelets, and is also the key effector protease of the coagulation cascade. Thrombin activates platelets predominantly via protease-activated receptors (PARs) [3–9]. Mice lacking all platelet PAR function (PAR4<sup>-/-</sup>) are protected against thrombosis without showing spontaneous bleeding [10–15], indicating the potential of targeting these receptors for antithrombotic therapy. There are two PARs on human platelets, PAR1 and PAR4. Of these, PAR1 is the higher-affinity thrombin receptor and has been the focus of antiplatelet drug development with two PAR1 antagonists, atropaxar (E5555) [16] and vorapaxar [17,18], having been evaluated in clinical trials. Vorapaxar was recently granted Food and Drug Administration approval for use in the USA for the prevention of thrombotic events in patients with a history of myocardial infarction or peripheral artery disease [19]. However, vorapaxar in combination with single or dual antiplatelet therapy was also associated with significantly increased rates of intracranial bleeding, particularly in patients with a history of stroke or other predisposing factors [17,18]. These observations suggest that there remains much to learn regarding how to best manipulate thrombin signaling in platelets for safe and effective antithrombotic therapy. With this in mind, there has been renewed interest in the function of PAR4 on human platelets in the setting of thrombus formation.

Whether PAR4 performs distinct functions from PAR1 on human platelets or simply acts in a redundant manner as a ‘back-up’ receptor has long been debated, although recent studies suggest that the former alternative is more likely [20–25]. One major distinction between the two platelet PARs relates to the kinetics of intracellular signaling [20,21,26–28]. Both PAR1 and PAR4 signal via Gq to mobilize intracellular calcium and drive platelet functions, including integrin activation, granule secretion, and phosphatidylserine (PS) exposure. However, PAR4 activation induces a slower and more sustained intracellular calcium signal than PAR1 activation [26,29]. This temporal difference in calcium signaling may be attributable, in part, to an anionic sequence C-terminal to the PAR4 cleavage site that was recently shown to prolong the thrombin–PAR4 interaction and facilitate persistent receptor activation [29]. The cellular consequences of such sustained platelet activation downstream of PAR4 have not yet been completely characterized, but may involve sustained platelet-secretion kinetics [30] and platelet procoagulant function, given the reliance of these phenomena on sustained, elevated, intracellular calcium levels [31,32]. This hypothesis is supported by studies showing that selective activation of PAR4, but not of PAR1, results in the release of coagulation factor V from  $\alpha$ -granules [22], microparticle shedding [22], and partially sustained Akt phosphorylation [33].

Despite this previous work, there have been no studies examining the contribution of PAR4 to procoagulant

activity in the setting of human thrombus formation. This may be because of the limited availability of appropriate PAR4 antagonists required for such studies. The most commonly used PAR4 antagonists are the small molecule YD-3 [34], the peptidomimetic tc-YPGKF-NH<sub>2</sub> [35], and the pepducins P4pal-10 and P4pal-i1 [36–38]. However, these agents are either not widely available (YD-3) or have been reported to lack specificity (pepducins) and/or efficacy (tc-YPGKF-NH<sub>2</sub>) in studies using human platelets [39–41].

In order to determine the impact of selectively inhibiting PAR4, we developed a function-blocking anti-PAR4 antibody, and used it to probe for platelet activation events that are reliant on PAR4. Selective inhibition of PAR4 with the anti-PAR4 antibody, but not selective inhibition of PAR1 with atropaxar, impaired platelet procoagulant activity in human isolated platelets and in a whole blood thrombosis assay. Defining the relative functions of the two platelet PARs is of patent interest for any rationalization of PAR4 as an antiplatelet drug target. Our findings demonstrate that thrombin-dependent platelet procoagulant activity – and consequent thrombin generation and fibrin formation – is mediated predominantly by PAR4 and is largely independent of PAR1 in the setting of thrombus formation. These findings suggest that PAR1 and PAR4 have divergent functions on platelets that may be important for thrombus formation and/or stabilization, and that PAR4 antagonists may have additional and/or distinct utility in the prevention of arterial thrombosis.

## Materials and methods

### Materials

A rabbit polyclonal anti-PAR4 antibody was raised against a 21-residue peptide with the sequence GGDDSTPSILPAPR<sup>↓</sup>GYPGQVC [42], matching residues 34–54 of human PAR4, which spans the thrombin cleavage site (indicated by <sup>↓</sup> in the peptide sequence), as previously described [43]. All antibody production protocols were approved by the Alfred Medical Research and Education Precinct Animal Ethics Committee. Antibodies in pooled sera from immunized rabbits were affinity-purified with an AffiGel 10–15 column (BioRad, Hercules, CA, USA) conjugated with the antigen peptide. Preimmune antibody was isolated from sera with protein G–Sephadex (BioRad). The following reagents were stored according to the manufacturers’ recommendations: E5555 (atropaxar, gift from P. Little, University of Queensland Brisbane, QLD, Australia), fluorescein isothiocyanate (FITC)-conjugated anti-rabbit IgG (Millipore, Lake Placid, NY, USA), FITC-conjugated PAC-1 antibody (BD Biosciences, San Jose, CA, USA), AK4 anti-P-selectin antibody (Santa Cruz Biotech, Santa Cruz, CA, USA), FITC-conjugated anti-mouse antibody (Millipore),

Alexa Fluor 488-conjugated annexin-V (Sigma, St Louis, MO, USA), anti-CD9-phycoerythrin (PE) (BD Biosciences), anti-CD41a antibody (BD Biosciences), Alexa Fluor 488-conjugated anti-fibrin antibody (clone 59D8 [44,45]; generous gift from V. Chen, University of New South Wales, Sydney, NSW, Australia), Oregon Green BAPTA-1 AM, Fura Red AM, DM-BAPTA AM (all from ThermoFisher Scientific, Waltham, MA, USA), human  $\alpha$ -thrombin (Sigma; activity expressed in NIH Units [U] obtained by direct comparison with NIH thrombin reference standard), PAR4-activating peptide (PAR4-AP; AYPGKF), PAR1-activating peptide (PAR1-AP; TFLLR) (both from Auspep, Melbourne, Victoria, Australia), ADP (Sigma), cross-linked collagen-related peptide (CRP) (from R. Farndale, University of Cambridge, Cambridge, UK), calcium ionophore A23187 (Sigma), bovine type 1 collagen (Sigma), and hirudin (lepirudin; Celgene, Melbourne, Victoria, Australia). A fluorescence resonance energy transfer (FRET)-based thrombin activity sensor (Thr-SP, a generous gift from S. Diamond, University of Pennsylvania, Philadelphia, PA, USA), was linked to the anti-CD41a antibody via CLICK chemistry as previously described [46], to examine thrombin activity in human whole blood under flow.

#### Human blood samples

All human studies were approved by the Monash University Human Research Ethics Committee. Blood was collected after informed consent had been obtained from healthy adults (aged 21–50 years, of both sexes) who had not taken antiplatelet medications in the past 10 days. Blood was drawn from the antecubital vein with a 19-gauge butterfly needle into syringes containing either one-seventh volume acid citrate dextrose (ACD) (7 : 1 v/v, final concentration) for platelet isolation, or one-tenth volume trisodium citrate (0.32% w/v, final concentration) for whole blood flow experiments, as previously described [47].

#### Mice

All mouse studies were approved by the Alfred Medical Research and Education Precinct Animal Ethics Committee. Mice in these studies were either PAR4-deficient (PAR4<sup>-/-</sup>) [10] or wild-type (PAR4<sup>+/+</sup>). Mouse whole blood was collected into ACD (7 : 1 v/v, final concentration), and platelets were isolated as previously described [47].

#### Detection of PAR4 via flow cytometry

Human or mouse washed platelets ( $5 \times 10^7$  mL<sup>-1</sup>) were incubated with the anti-PAR4 antibody (0.1 mg mL<sup>-1</sup>) for 30 min at 37 °C, and then fixed with paraformaldehyde (1% v/v, final concentration). The suspension was then centrifuged at  $1000 \times g$  for 2 min to obtain the platelet pellet, which was then resuspended in modified

Tyrodé's buffer (12 mM NaHCO<sub>3</sub>, 10 mM HEPES, pH 7.4, 137 mM NaCl, 2.7 mM KCl, 5.5 mM D-glucose, 1 mM CaCl<sub>2</sub>) containing a 1 : 50 dilution of an FITC-conjugated anti-rabbit IgG. After 30 min at room temperature, the samples were centrifuged again, and the platelet pellet was resuspended in modified Tyrodé's buffer and analyzed by use of a flow cytometer (FACSCalibur; BD Biosciences).

#### Platelet aggregation

Human isolated platelets ( $3 \times 10^8$  mL<sup>-1</sup>) were pretreated for 10 min at 37 °C with dimethylsulfoxide (DMSO) (0.1% v/v), the PAR1 antagonist E5555 (0.1  $\mu$ M; minimum concentration required for inhibition of PAR1-AP [48]), the anti-PAR4 antibody (0.1 mg mL<sup>-1</sup>), or a combination of E5555 and the anti-PAR4 antibody. Platelets were treated with one of thrombin (0.1 or 1 U mL<sup>-1</sup>), PAR4-AP (100  $\mu$ M), PAR1-AP (10  $\mu$ M), ADP (10  $\mu$ M), or CRP (10  $\mu$ g mL<sup>-1</sup>). Platelet aggregation was measured with a platelet aggregometer (Helena Laboratory, Beaumont, TX, USA).

#### Microplate-based platelet aggregometry

Human and mouse washed platelets were adjusted to a concentration of  $2 \times 10^8$  platelets mL<sup>-1</sup> in Tyrodé's buffer supplemented with 1.8 mM CaCl<sub>2</sub>, 10% bovine serum albumin, and 0.02 U mL<sup>-1</sup> apyrase. Platelets (100  $\mu$ L) were pipetted into each well of a clear, flat-bottomed 96-well tissue culture plate (BD Falcon (TM); BD Biosciences). Buffer alone was set to the maximum aggregation level, and unstimulated platelets were used set to the baseline light transmission (no aggregation). Agonists were added, and the plate was analyzed in a FLUOstar OPTIMA (BMG Labtech, Ortenberg, Germany) plate reader with a 595-nm excitation filter. After each reading cycle, the plate was subjected to a 5-min shake period (double orbital shake; shaking width of 4 mm), and this was repeated for a total of 10 cycles. Platelets were maintained at 37 °C throughout the experiment with the in-built incubator in the plate reader. Ability to aggregate was calculated as  $\frac{OD(\text{No Agonist}) - OD(\text{Agonist})}{OD(\text{No Agonist}) - OD(\text{blank})} \times 100$  at the time point when aggregation was at a maximum.

#### Platelet calcium signaling dynamics

Calcium flux was measured in human isolated platelets with a dual-dye (ratiometric) microimaging assay, as previously described [49]. Briefly, isolated human platelets ( $3 \times 10^8$  mL<sup>-1</sup>) were loaded with Oregon Green BAPTA-1 AM (0.625  $\mu$ M) and Fura Red AM (0.625  $\mu$ M) at 37 °C for 30 min, as in [49]. Dye-loaded platelets were subsequently treated with PAR antagonists (as above), and stimulated with thrombin (1 U mL<sup>-1</sup>). Cells were monitored in real time, and emission at 500–570 nm and

600–710 nm was recorded every 10 s for 10 min (Nikon A1r). The increase in cytosolic calcium concentration (nm) was determined according to [49].

#### Platelet activation events

The platelet activation events measured were integrin activation (PAC-1 antibody binding), P-selectin expression (AK4 antibody binding), and PS exposure (annexin-V binding). Human isolated platelets ( $5 \times 10^7 \text{ mL}^{-1}$ ) were pretreated with PAR antagonists as described above, and incubated with either FITC-conjugated PAC-1 antibody (1 : 100), AK4 anti-P-selectin antibody (1 : 100) followed by an FITC-conjugated anti-mouse antibody (1 : 10 000), or Alexa Fluor 488-conjugated annexin-V (1 : 100), prior to stimulation with either PAR1-AP, PAR4-AP, thrombin, CRP, or calcium ionophore A23187, as described above. After stimulation, platelets were resuspended in modified Tyrode's buffer for flow cytometry analysis (FACSCalibur; BD Biosciences).

#### Whole blood thrombosis assay

Human whole blood collected in citrate (3.2%) was preincubated for 15 min at 37 °C with both PE-conjugated anti-CD9 antibody ( $4 \mu\text{g mL}^{-1}$ ) and Thr-SP ( $5 \mu\text{M}$ ), and one of hirudin ( $800 \text{ U mL}^{-1}$ ), DMSO (1% v/v), PAR-1 antagonist E5555 ( $1 \mu\text{M}$ ), anti-PAR4 antibody ( $0.2 \text{ mg mL}^{-1}$ ), or the combination of both PAR inhibitors. In parallel experiments, an Alexa Fluor 488-conjugated anti-fibrin antibody ( $5 \mu\text{g mL}^{-1}$ ) was substituted for the thrombin sensor in order to analyze fibrin deposition. Whole blood was recalcified with 5–7.5 mM  $\text{CaCl}_2$  (final concentration) to initiate coagulation, and drawn over glass microslides ( $1 \times 0.1\text{-mm}$  internal diameter; Vitrotubes, Vitrocom, NJ, USA) coated with bovine type 1 collagen ( $250 \mu\text{g mL}^{-1}$ ) by use of a Harvard pump (Instech Laboratories, Plymouth Meeting, PA, USA) at a fixed flow rate of  $0.06 \text{ mL min}^{-1}$ , resulting in a wall shear rate of  $600 \text{ s}^{-1}$ . Dual-color confocal fluorescence images were recorded at excitation wavelengths of 488 nm and 561 nm, and collected through a  $\times 40$  water immersion objective. Confocal Z-stacks were continuously recorded (16-bit images of  $512 \times 512$  pixels;  $317 \times 317 \mu\text{m}$ , Z-step  $0.5 \mu\text{m}$ ; Nikon A1r, with NIS software) for 2 min, and modified, calcium-free Tyrode's buffer was then passed over the thrombi, and Z-stacks encompassing the entire height of the thrombus field were recorded over a period of 10 min. Offline analysis of thrombi parameters was performed with NIS software. Image series were initially thresholded empirically, and the same threshold was then applied to all subsequent experiments, as identical experimental and confocal settings were used throughout. Platelet thrombi were defined by the use of anti-CD9-PE, and thrombin activ-

ity and fibrin volume were quantified according to the average fluorescence of the thrombus field. Data were normalized against the hirudin baseline, and expressed as a percentage of the control.

#### Statistical analyses

Statistical analyses were performed with GRAPHPAD PRISM (version 6.0, La Jolla, CA, USA). Significance was defined at  $P < 0.05$  as determined with either an unpaired, two-tailed Student's *t*-test or one-way ANOVA with Fisher's LSD test for multiple comparisons, as advised and indicated in the relevant figure legends.

## Results

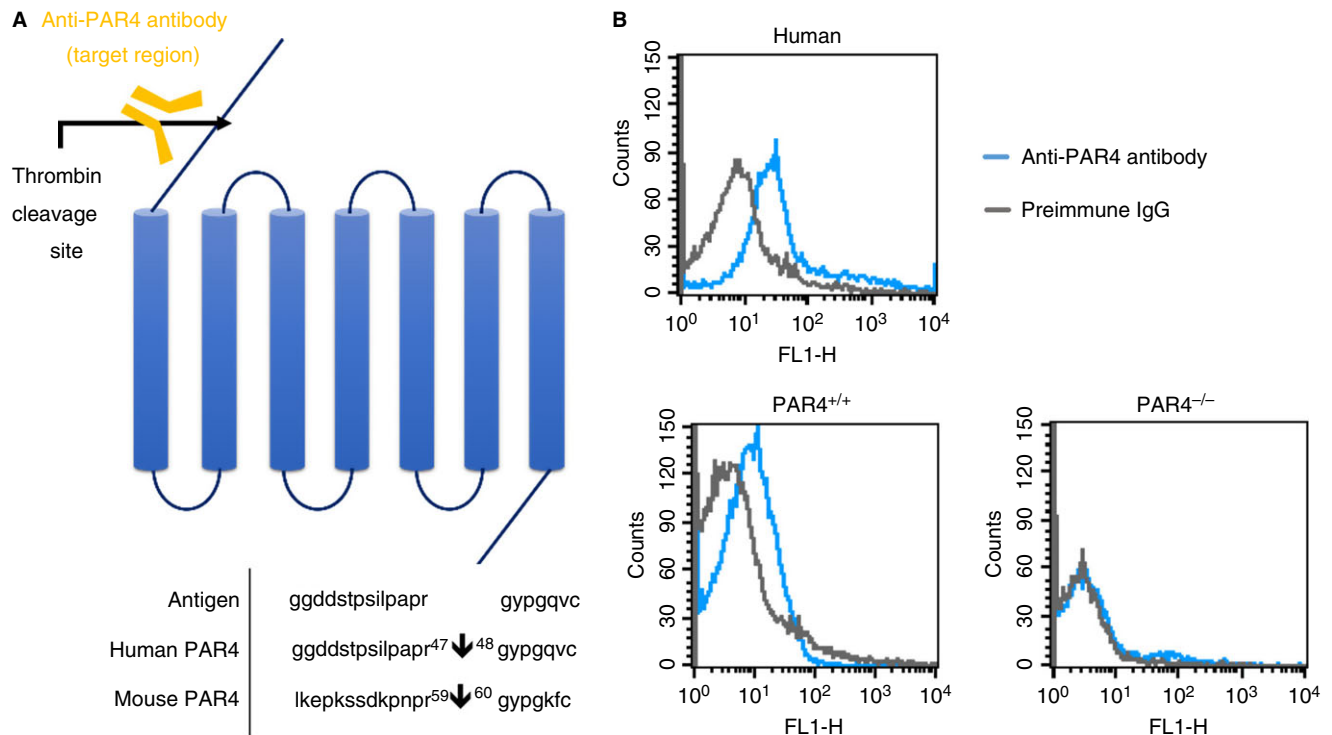
#### Selective inhibition of PAR4-mediated platelet activation by an anti-PAR4 antibody

A rabbit polyclonal anti-PAR4 antibody was raised against a peptide with sequence matching a region spanning the thrombin cleavage site of human PAR4 (Fig. 1A). Affinity-purified anti-PAR4 antibody recognized both human and mouse PAR4 on platelets as assessed by flow cytometry. The anti-PAR4 antibody produced a consistent rightward shift in fluorescence intensity as compared with the same concentration of preimmune rabbit IgG in platelets isolated from either humans or wild-type mice (Fig. 1B). This signal was entirely absent in platelets isolated from PAR4<sup>-/-</sup> mice (Fig. 1B), indicating that the antibody selectively binds to PAR4 on the native platelet surface.

The inhibitory function of the anti-PAR4 antibody was first assessed via aggregation of human washed platelets. Such studies were confounded by the requirement to use thrombin as the agonist (the anti-PAR4 antibody is predicted to inhibit thrombin-induced but not PAR1-AP-induced or PAR4-AP-induced receptor activation). As thrombin activates human platelets via both PAR1 and PAR4, we first established conditions of thrombin-induced platelet aggregation that were largely PAR4-dependent (Fig. S1). To this end,  $0.1 \text{ U mL}^{-1}$  was the minimum thrombin concentration required to reliably induce maximum aggregation of human platelets (Fig. S1C). Responses to  $0.1 \text{ U mL}^{-1}$  thrombin were partially inhibited in a concentration-dependent manner by the PAR1 antagonist E5555, but remained largely intact in the presence of  $0.1 \mu\text{M}$  E5555 (Fig. S1D). This residual response to  $0.1 \text{ U mL}^{-1}$  thrombin in the presence of  $0.1 \mu\text{M}$  E5555 was inhibited in a concentration-dependent manner by the anti-PAR4 antibody, and was almost abolished with an antibody concentration of  $0.1 \text{ mg mL}^{-1}$  (Fig. S1E).

The specificity of this inhibitory effect of the anti-PAR4 antibody was then assessed under these conditions (Fig. 2). Here, aggregation of human platelets stimulated with thrombin ( $0.1 \text{ U mL}^{-1}$ ) was unaffected by pretreatment





**Fig. 1.** The anti-protease-activated receptor (PAR)4 antibody selectively binds to PAR4 on the platelet surface. (A) Schematic of PAR4 showing the region corresponding to the antigen peptide sequence (yellow), which spans the thrombin cleavage site (gray arrow). Note the homology between the mouse and human PAR4 sequences in this region. (B) Representative ( $N = 3-6$ ) flow cytometry traces showing binding of the anti-PAR4 antibody (blue line,  $0.1 \text{ mg mL}^{-1}$ ) versus preimmune rabbit IgG (gray line,  $0.1 \text{ mg mL}^{-1}$ ) to platelets isolated from a human or a wild-type (PAR4<sup>+/+</sup>) or PAR4-deficient (PAR4<sup>-/-</sup>) mouse. Note the complete loss of binding in platelets from the PAR4<sup>-/-</sup> mouse.

with either a PAR1 antagonist (E5555;  $0.1 \mu\text{M}$ ) or the anti-PAR4 antibody ( $0.1 \text{ mg mL}^{-1}$ ) alone, but was abolished in the presence of both inhibitors combined (Fig. 2A). This inhibitory effect was overcome by increased thrombin concentrations (Figs. 2B and S1F), indicating that the anti-PAR4 antibody functions as a competitive inhibitor of thrombin-induced platelet aggregation. The anti-PAR4 antibody had no effect on platelet aggregation induced by PAR4-AP (AYPGKF,  $100 \mu\text{M}$ ; Fig. 2C), demonstrating specificity of the anti-PAR4 antibody against thrombin cleavage of the receptor over other PAR4 activation mechanisms. Furthermore, the anti-PAR4 antibody had no effect on aggregation induced by any of PAR1-AP, ADP, or CRP (Fig. 2D–F). Importantly, PAR1-AP-induced aggregation was abolished by the PAR1 antagonist (E5555;  $0.1 \mu\text{M}$ ), confirming effective PAR1 antagonism under the conditions used here (Fig. 2D).

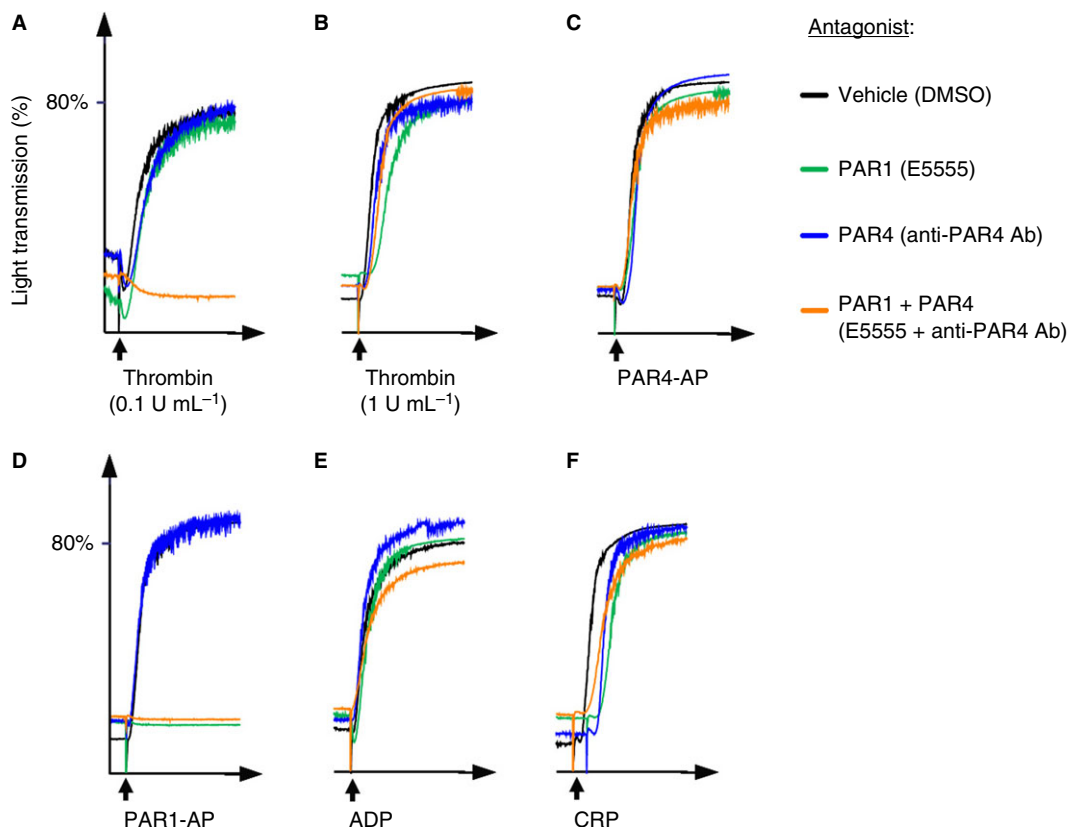
#### Selective inhibition of PAR4 abolishes sustained thrombin-induced calcium signals in human platelets

To confirm the inhibition of thrombin-induced platelet function by the anti-PAR4 antibody, we examined changes in cytosolic calcium flux. In accord with the well-characterized thrombin-induced calcium signaling profile in platelets and other cells [50–53], we observed that thrombin ( $1 \text{ U mL}^{-1}$ ) induced an initial transient increase

in the cytosolic calcium concentration in human platelets that was followed by a sustained plateau (Fig. 3A). Inhibition of PAR1 alone (E5555;  $0.1 \mu\text{M}$ ) significantly attenuated the initial transient increase in the cytosolic calcium concentration (Fig. 3A,B) but had no effect on the sustained increase in the cytosolic calcium concentration (Fig. 3A,C). In contrast, inhibition of PAR4 alone (anti-PAR4 antibody;  $0.1 \text{ mg mL}^{-1}$ ) significantly decreased the initial transient increase (Fig. 3A,B) and abolished the sustained plateau (Fig. 3A,C). Pretreatment of platelets with both PAR inhibitors essentially eliminated all thrombin-induced calcium signaling within the 10-min observation period (Fig. 3A–D). These studies are suggestive of a role for PAR4 in mediating platelet activation events that are reliant on sustained calcium signaling. Therefore, we next examined the effect of PAR4 inhibition on specific platelet activation readouts that are known to be dependent on either acute (integrin  $\alpha_{\text{IIb}}\beta_3$  activation;  $\alpha$ -granule release) or sustained (PS externalization) elevations in intracellular calcium.

#### Selective inhibition of PAR4 is sufficient to impair thrombin-induced PS exposure, but not $\alpha_{\text{IIb}}\beta_3$ activation or P-selectin expression, on human platelets

We examined the effect of PAR inhibition on three separate markers of platelet activation:  $\alpha_{\text{IIb}}\beta_3$  activation



**Fig. 2.** The anti-protease-activated receptor (PAR)4 antibody specifically inhibits thrombin-induced PAR4 activation of human platelets. Shown are representative ( $N \geq 3$ ) aggregation traces of human isolated platelets in the presence of a vehicle control (black; dimethylsulfoxide [DMSO], 0.1% v/v), a PAR1 inhibitor (green; E5555, 0.1  $\mu\text{M}$ ), the anti-PAR4 antibody (blue; 0.1  $\text{mg mL}^{-1}$ ), or both PAR inhibitors (orange). Platelets were stimulated with (A) 0.1  $\text{U mL}^{-1}$  thrombin, (B) 1  $\text{U mL}^{-1}$  thrombin, (C) 100  $\mu\text{M}$  PAR4-activating peptide (PAR4-AP), (D) 10  $\mu\text{M}$  PAR1-activating peptide (PAR1-AP), (E) 10  $\mu\text{M}$  ADP, or (F) 10  $\mu\text{g mL}^{-1}$  collagen-related peptide (CRP). Note that the anti-PAR4 antibody inhibits thrombin-induced (0.1  $\mu\text{M}$ ) platelet aggregation in the presence of concomitant PAR1 antagonism, but that no inhibition is observed at a higher thrombin concentration (1  $\text{U mL}^{-1}$ ) or with other platelet activation mechanisms. Arrows indicate the time point of agonist addition. Ab, antibody.

(PAC-1 antibody binding),  $\alpha$ -granule release (AK4 antibody binding to P-selectin), and PS externalization (annexin-V binding). Inhibition of either PAR1 or PAR4 alone did not significantly affect  $\alpha_{\text{IIb}\beta_3}$  activation (Fig. 4A) or P-selectin expression (Fig. 4B) in response to thrombin (0.1  $\text{U mL}^{-1}$ ), but concomitant inhibition of both receptors produced a non-significant decrease in both  $\alpha_{\text{IIb}\beta_3}$  activation and P-selectin expression. In contrast, PS exposure in response to thrombin (1  $\text{U mL}^{-1}$ ) was significantly reduced in platelets pretreated with the anti-PAR4 antibody alone ( $P > 0.05$ ; Fig. 4C). Pretreatment with the PAR1 inhibitor E5555 (0.1  $\mu\text{M}$ ) also decreased the number of PS-expressing platelets, although combined inhibition of PAR1 and PAR4 caused no further reduction over PAR4 inhibition alone (Fig. 4C). Further evidence of a role for PAR4 in mediating platelet PS exposure was provided by the observation that stimulation of human platelets with PAR4-AP (300  $\mu\text{M}$ ; minimum concentration required to elicit maximal platelet activation; Fig. S1) significantly increased the number of annexin-V-positive platelets ( $P < 0.05$ ) but that stimula-

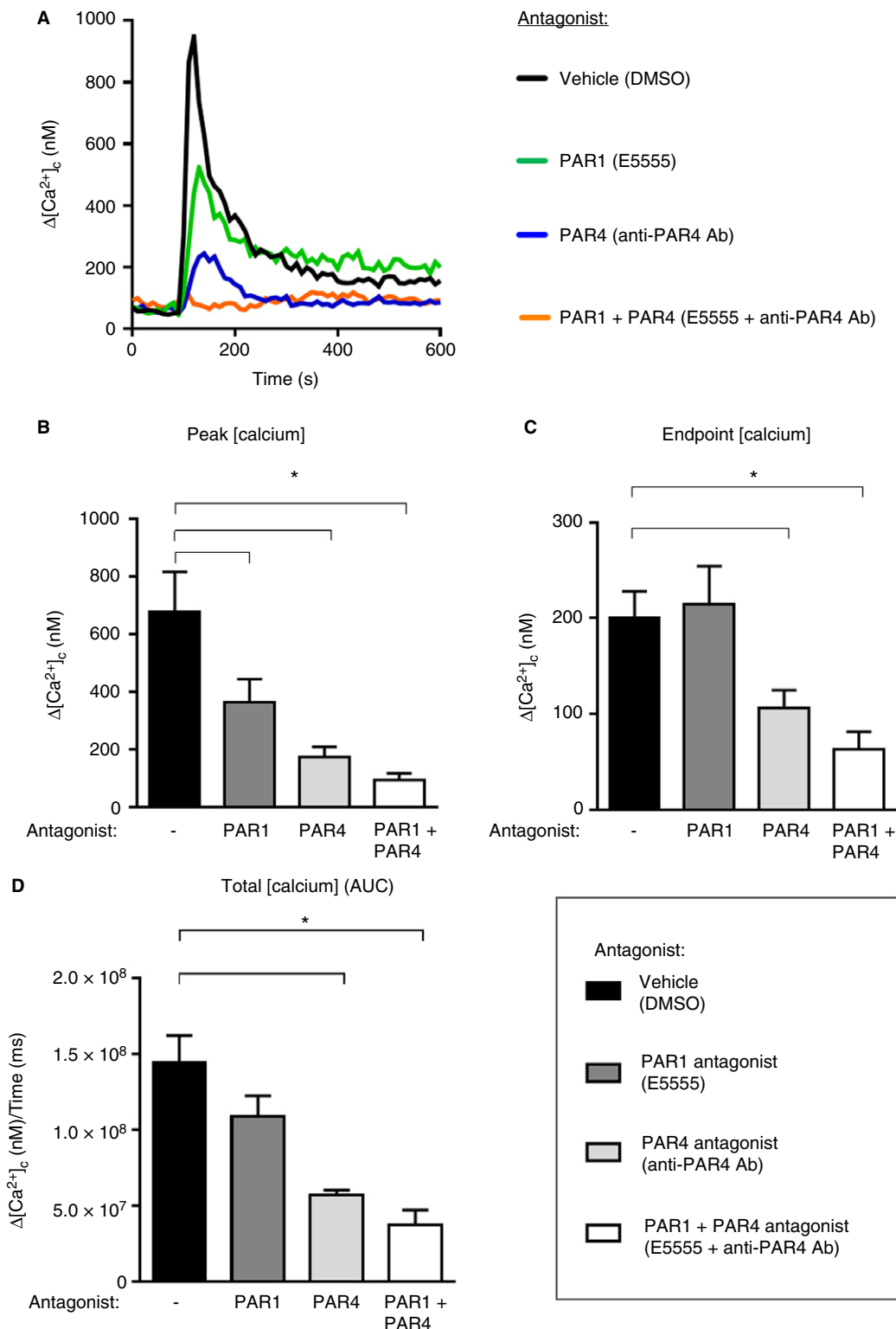
tion with PAR1-AP (30  $\mu\text{M}$ ; 10-fold the concentration required to elicit maximal platelet activation; Fig. S1) did not (Fig. 4D).

As observed for aggregation (Fig. 2), the anti-PAR4 antibody had no effect on any of  $\alpha_{\text{IIb}\beta_3}$  activation,  $\alpha$ -granule release or PS externalization induced by higher concentrations of thrombin (1  $\text{U mL}^{-1}$ ), PAR4-AP, CRP, or, in the case of PS externalization, calcium ionophore (Fig. 4D–F). In addition, only platelets treated with a PAR1 inhibitor (E5555, 0.1  $\mu\text{M}$ ) were unresponsive to platelet activation with PAR1-AP (Fig. 4E–G), further confirming the specificity of PAR1 and PAR4 inhibition in these studies. Together, these findings suggest that selective inhibition of PAR4 is sufficient to impair thrombin-induced PS externalization in human platelets.

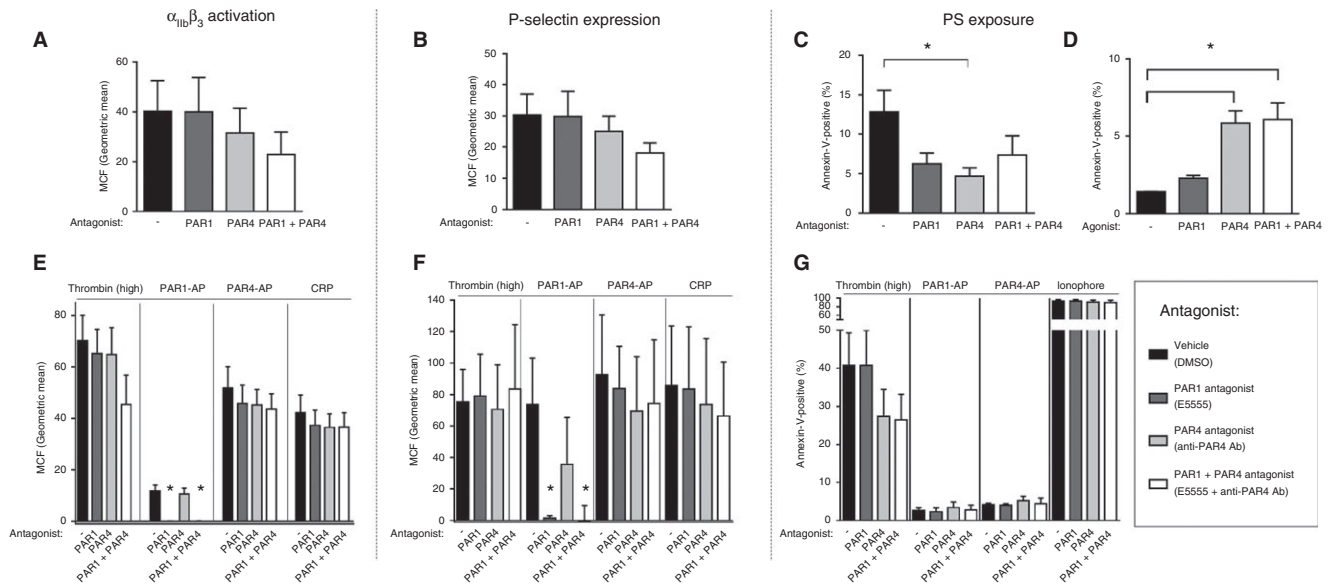
#### *Selective inhibition of PAR4 impairs platelet procoagulant activity during thrombus formation*

A human whole blood thrombosis assay was performed under conditions of coagulation and with a FRET-based





**Fig. 3.** Selective inhibition of protease-activated receptor (PAR)4 abolishes the sustained thrombin-induced calcium signals in human platelets. (A) Representative trace of cytosolic calcium flux in human platelets pretreated with vehicle (black; dimethylsulfoxide [DMSO], 0.1% v/v), a PAR1 inhibitor (green; E5555, 0.1  $\mu\text{M}$ ), the anti-PAR4 antibody (blue; 0.1 mg mL<sup>-1</sup>), or both PAR inhibitors (orange), and stimulated with thrombin (1 U mL<sup>-1</sup> at time = 60 s). (B–D) Real-time responses were analyzed for (B) the initial transient increase in cytosolic calcium concentration ( $\Delta[\text{Ca}^{2+}]_c$ ) (peak), the (C) sustained plateau (endpoint), and (D) total calcium generated (expressed as area under the curve [AUC]). Note that inhibition of PAR4, but not of PAR1, inhibits the sustained plateau, and that inhibition of both receptors essentially abolishes the response. Data are mean  $\pm$  standard error of the mean from three independent experiments (donors); \* $P$  < 0.05 versus control (one-way ANOVA with *post hoc* Fisher's LSD test). Ab, antibody.



**Fig. 4.** Selective inhibition of protease-activated receptor (PAR)4 is sufficient to impair thrombin-induced phosphatidylserine (PS) exposure, but not  $\alpha_{IIb}\beta_3$  activation or P-selectin expression, on human platelets. Thrombin-induced platelet activation events in human platelets were examined by measuring (A, E)  $\alpha_{IIb}\beta_3$  activation (PAC-1), (B, F)  $\alpha$ -granule release (P-selectin via AK4) or (C, D, G) PS exposure (annexin-V) via flow cytometry. Human washed platelets were pretreated with a vehicle control (dimethylsulfoxide [DMSO], 0.1% v/v), a PAR1 inhibitor (E5555, 0.1  $\mu$ M), a PAR4 inhibitor (anti-PAR4 antibody, 0.1 mg mL<sup>-1</sup>), or the combination of PAR inhibitors, before (A–C) thrombin stimulation (0.1 U mL<sup>-1</sup> for 10 min for  $\alpha_{IIb}\beta_3$  activation and  $\alpha$ -granule release; 1 U mL<sup>-1</sup> for 30 min for PS exposure). Note that only PS exposure is inhibited by the anti-PAR4 antibody alone. Note also (D) that PS exposure is increased in response to PAR4-activating peptide (PAR4-AP) (300  $\mu$ M) but not PAR1-activating peptide (PAR1-AP) (30  $\mu$ M). (E, F)  $\alpha_{IIb}\beta_3$  activation and  $\alpha$ -granule release were also examined in response to high thrombin (1 U mL<sup>-1</sup>), PAR1-AP (10  $\mu$ M), PAR4-AP (300  $\mu$ M), and collagen-related peptide (CRP) (10  $\mu$ g mL<sup>-1</sup>). (G) PS exposure in response to thrombin plus CRP (1 U mL<sup>-1</sup> and 10  $\mu$ g mL<sup>-1</sup>, respectively), PAR1-AP (30  $\mu$ M), PAR4-AP (300  $\mu$ M) or calcium ionophore (10  $\mu$ M) was also examined. Data are mean  $\pm$  standard error of the mean of either mean cell fluorescence (MCF, geometric mean) or percentage of positive population, as indicated. \* $P$  < 0.05 versus control (one-way ANOVA with *post hoc* Fisher's LSD test);  $N$  = 6–8. Ab, antibody.

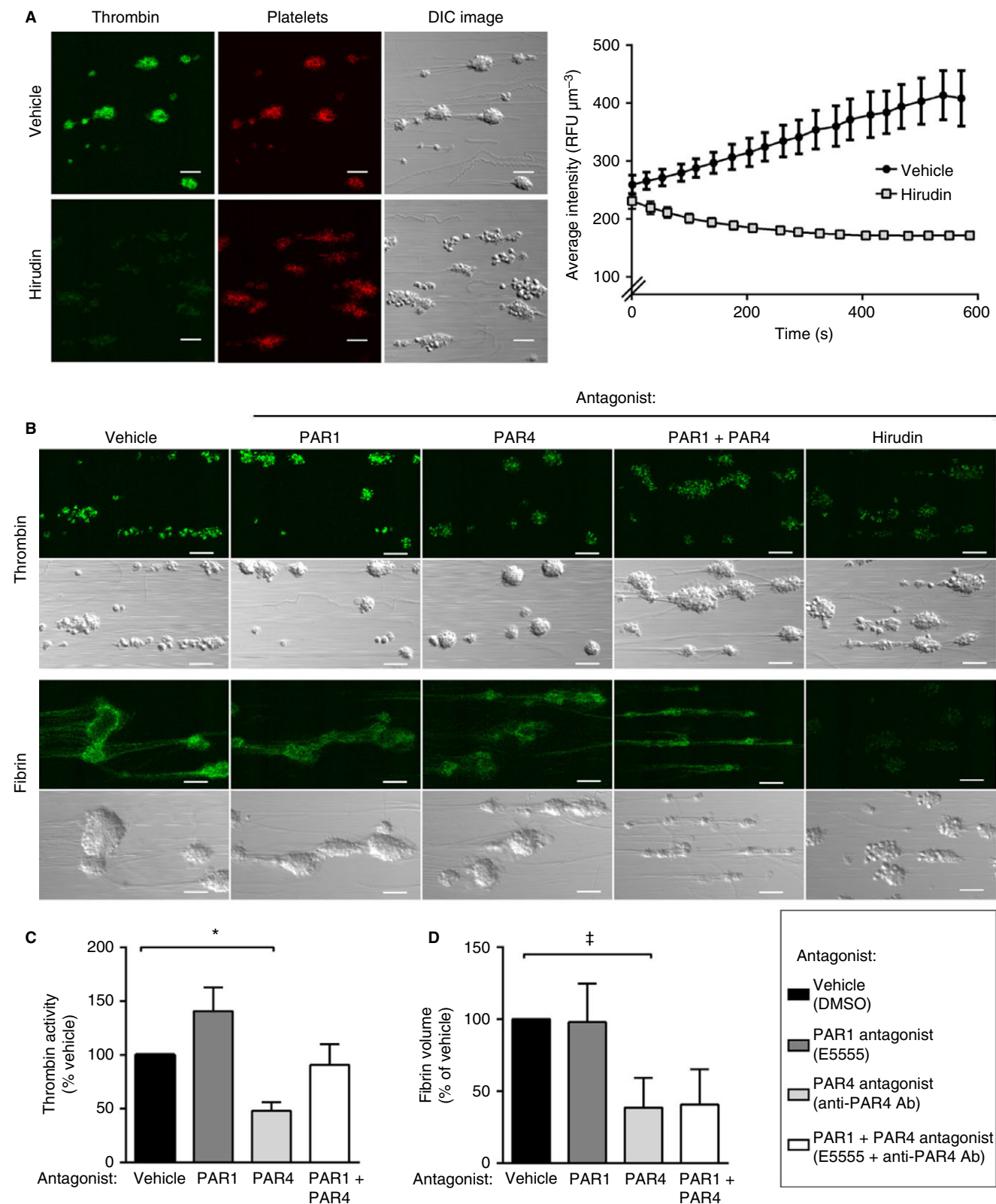
sensor of thrombin activity (ThSP-Ab) [46]. We observed significant thrombin generation on platelet thrombi that was abolished in the presence of the direct thrombin inhibitor hirudin, despite platelet deposition that continued unabated (Fig. 5A). Selective inhibition of PAR4, but not of PAR1, significantly reduced the thrombin activity detected on platelet thrombi (Fig. 5B,C). Curiously, we observed that the PAR1 antagonist E5555 caused an increase in thrombin activity in these experiments (Fig. 5C). The cause of this effect remains unknown, but may relate to the recently reported effects of PAR1 antagonists in promoting platelet activation [48,54]. In similar experiments, selective inhibition of PAR4, but not of PAR1, also reduced the amount of fibrin generation as

compared with the control (Fig. 5B,D). These findings indicate that selective inhibition of PAR4, but not of PAR1, is sufficient to decrease thrombin activity and consequent fibrin formation in the setting of human thrombus formation.

## Discussion

The current study provides the first evidence that PAR4 is required for platelet procoagulant function during thrombus formation in human blood. Here, we used a strategy similar to that previously employed by others [42,55] to develop a function-blocking anti-PAR4 antibody that was an effective and highly specific inhibitor of

**Fig. 5.** Selective inhibition of protease-activated receptor (PAR)4 impairs platelet procoagulant activity during thrombus formation in whole blood. (A) Representative images and quantification of thrombin activity (green; ThSP-Ab) and platelet deposition (red; CD9-phycoerythrin) in a human whole blood thrombosis assay analyzed with confocal microscopy. Note that the direct thrombin inhibitor hirudin (800 U mL<sup>-1</sup>) abolished thrombin activity despite continued platelet deposition. (B–D) Representative images (B) and quantification of (C) thrombin activity ( $N$  = 7) and (D) fibrin deposition ( $N$  = 4) at the 10-min time point of a human whole blood thrombosis assay performed in the presence of a vehicle control (dimethylsulfoxide, 0.1% v/v), a PAR1 inhibitor (E5555, 1  $\mu$ M), a PAR4 inhibitor (anti-PAR4 antibody, 0.2 mg mL<sup>-1</sup>), the combination of PAR inhibitors, or hirudin (800 U mL<sup>-1</sup>). Data are mean  $\pm$  standard error of the mean of four to seven individual donors, and are expressed as average fluorescence intensity of thrombi (RFU  $\mu$ m<sup>-3</sup>) as a percentage of the vehicle control for each donor (C) or total fibrin as a percentage of the vehicle control for each donor (D). \* $P$  < 0.05 and ‡ $P$  = 0.05 versus vehicle control (one-way ANOVA with *post hoc* Fisher's LSD test). Scale bar: 10  $\mu$ m. Ab, antibody; DIC, differential interference contrast microscopy.



thrombin-induced PAR4 activation on human platelets. We used this anti-PAR4 antibody to probe for platelet functions that were primarily driven by PAR4-mediated rather than PAR1-mediated signaling. We observed that

selective PAR4 inhibition attenuated the sustained calcium signal and consequent PS exposure induced by thrombin in human isolated platelets, but not other commonly assessed platelet activation events. These effects

*in vitro* translated to a marked impact on platelet procoagulant activity, thrombin generation and subsequent fibrin formation in a human whole blood thrombosis assay, suggesting an important role for PAR4 during platelet-dependent thrombosis.

Whether PAR1 or PAR4 predominantly drives thrombin-induced platelet procoagulant activity has been the source of some controversy [22,24,56,57]. Here, we used selective antagonists in a series of *in vitro* and *ex vivo* assays to consistently identify PAR4 as the primary mediator of platelet-driven PS exposure. There are several explanations for the discrepancies in the conclusions drawn from our current work and earlier studies implicating PAR1 in this process. First, the initial studies in which selective PAR4 activation failed to induce platelet PS exposure were performed with a weak activator of PAR4 (GYPGKF [56]). We observed here that selective activation of PAR4 with a significantly more potent PAR4 agonist (AYPGFK [58]) was indeed capable of inducing PS exposure. Second, previous studies using inhibitors to implicate PAR1 in platelet procoagulant function have relied on pepducins to selectively inhibit PAR1 versus PAR4 [57]. We and others [39] have observed a lack of specificity of these reagents that was overcome in the current studies with the use of atopaxar and our anti-PAR4 antibody to selectively inhibit PAR1 and PAR4, respectively. Third, technical differences that have been shown to significantly impact on the annexin-V binding assay in isolated platelets [59] may underlie some of the discrepancies. Regardless of these differences, it is important to note that previous studies addressing this issue have been limited to *in vitro* experiments on isolated platelets and measuring surrogate markers of platelet procoagulant activity. On this point, we too observed that selective PAR1 inhibition impairs thrombin-induced PS exposure in human washed platelets. However, when we extended these *in vitro* observations to examine the physiologic consequences of platelet procoagulant activity, the impact of selective PAR4 inhibition was strongly retained, whereas PAR1 inhibition was clearly without effect.

Indeed, the current study is the first to examine the contribution of PAR4 to thrombin generation in a human whole blood thrombosis assay under physiologic flow conditions. Our findings show that PAR4 inhibition was sufficient to significantly impair thrombin generation on platelet thrombi and subsequent fibrin formation, independently of PAR1 inhibition. The extent of impairment in thrombin activity and fibrin production provided by PAR4 antagonism was notably similar, with an approximately 50% reduction in each case. Until very recently, thrombin generation in whole blood has been difficult to examine, with most techniques using closed system approaches in plasma, thus making platelet–thrombin interactions difficult to elucidate [60,61]. The effect of PAR-mediated platelet activation in human plasma has been examined previously, and it was shown that PAR4-

induced activation resulted in more rapid thrombin generation than observed following PAR1-induced activation [22]. Conversely, thrombin generation has been shown to be delayed by PAR4 inhibition, although total amount of thrombin generated was unaffected [24]. Here, we directly measured thrombin generation on platelet thrombi in whole blood under flow conditions known to be optimal for enhanced coagulation [46,62]. Our findings obtained with this approach indicate that PAR4 mediates platelet-dependent thrombin generation and consequent fibrin formation, suggesting that PAR4 inhibition significantly impairs platelet procoagulant activity during thrombosis.

Together, we have used four measures of platelet procoagulant activity to show that inhibition of PAR4 alone is sufficient to impair this important function. It is well known that PAR4 activation generates a more prolonged intracellular calcium signal than PAR1 activation [21,26], and our current study supports previous work [27,63] showing that PAR4 inhibition suppresses this sustained calcium signal. Given that the most notable platelet activation event dependent on a sustained rise in intracellular calcium levels is PS exposure [31,32], our findings build on recently published work suggesting that PAR4 is predominantly responsible for additional procoagulant effects of activated platelets [22,27], and extend this to the setting of human thrombus formation.

Our findings suggest that PAR4 inhibitors might provide a useful strategy for the prevention of arterial thrombosis that is distinct from existing approaches. Whether targeting platelet procoagulant activity for antithrombotic benefit is a viable option remains unknown, but it is worth noting that patients with Scott syndrome – a congenital disorder in which defective platelet PS exposure drastically diminishes the thrombin generation capacity – have a clinically significant bleeding phenotype [64]. There is also emerging evidence that targeting PAR4 might be a useful antithrombotic approach [65–67]. Indeed, recent reports have demonstrated increased expression and function of PAR4 in black Americans that might explain the increased resistance to current antiplatelet drugs and overall poorer cardiovascular outcomes in this population [67,68]. As patients treated with PAR1 antagonists have intact PAR4-mediated platelet responses [69,70], such clinical scenarios provide further impetus for the development of PAR4 antagonists as a distinct antithrombotic therapy.

## Conclusion

The results from the present study demonstrate that PAR4 performs at least one important platelet activation function that is distinct from that of PAR1, providing a further rationale for PAR4 antagonism as a novel target for the prevention of pathologic thrombosis. Given the ongoing clinical demand for improved and varied antiplatelet agents for the safe and effective prevention of



arterial thrombosis, the importance of thrombin-induced platelet activation in this setting, and the clinical setbacks with PAR1 antagonists, further investigation of PAR4 as a suitable target appears to be warranted.

### Addendum

S. L. French, J. F. Arthur, H. Lee, and W. S. Nesbitt performed experiments. S. L. French and J. R. Hamilton wrote the manuscript; J. R. Hamilton designed the research and supervised the study. All authors analyzed data and contributed to editing of the manuscript.

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### Disclosure of Conflict of Interests

The authors state that they have no conflict of interest.

### Supporting Information

Additional Supporting Information may be found in the online version of this article:

**Fig. S1.** Establishment of an assay to examine PAR4-dependent platelet activation by thrombin.

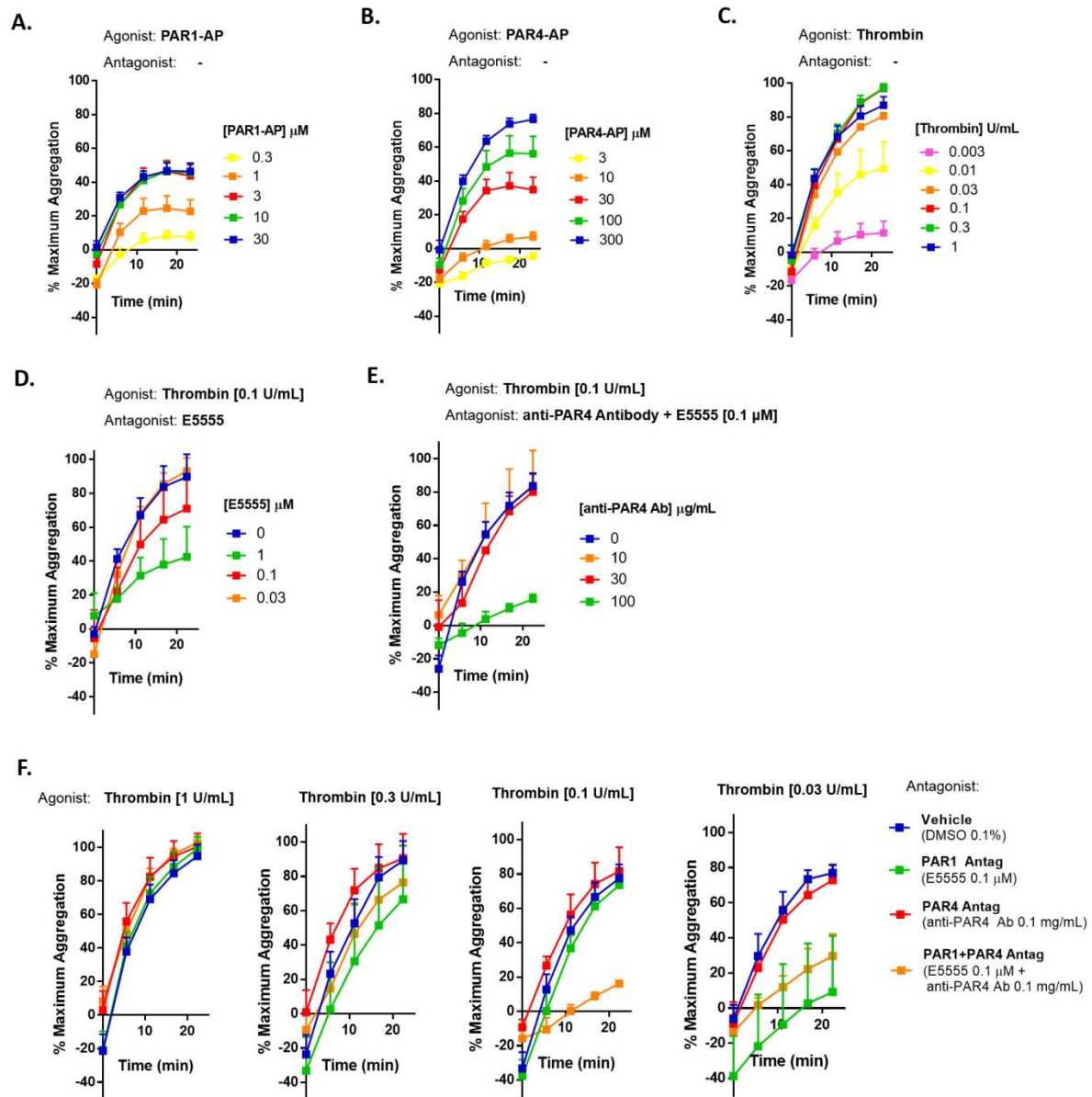
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## 2.3 Supplementary Figures to French et al. “Inhibition of protease-activated receptor 4 impairs platelet procoagulant activity during thrombus formation in human blood” (J Thromb Haemost, 2016)



Supplementary Figure 1.

**Supplementary Figure 1.** Establishment of an assay to examine PAR4-dependent platelet activation by thrombin. Isolated human platelet aggregation concentration response curves to (A) PAR1 activating peptide (PAR1-AP; TFLLR; 0.3-30  $\mu$ M), (B) PAR4 activating peptide (PAR4-AP; AYPGKF; 3-300  $\mu$ M) and (C) thrombin (0.003-1 U/mL) were conducted in a plate reader. (D) A PAR1 antagonist (E5555, 0.1  $\mu$ M (minimum concentration required to inhibit



PAR1-AP-induced platelet activation)) delays thrombin (0.1 U/mL) induced platelet aggregation but does not significantly affect maximal aggregation. (E) Platelet aggregation in response to thrombin (0.1 U/mL) and in the presence of the PAR1 antagonist, E5555 (0.1  $\mu$ M), is inhibited in a concentration-dependent manner by the anti-PAR4 antibody. (F) The effect of PAR inhibition on thrombin-induced platelet activation is overcome at high thrombin concentrations (1 and 0.3 U/mL), whereas at low thrombin concentrations (0.03 U/mL) PAR1 inhibition alone is sufficient to inhibit platelet aggregation. Data are expressed as mean  $\pm$  SEM;  $n \geq 3$ .

## **Chapter 3 - Perinatal lethality of PAR4-deficient mice delivered by primiparous dams reveals spontaneous bleeding in mice without platelet thrombin receptor function**

### **3.1 Introduction**

The studies of Chapter 2 generated a selective and effective experimental PAR4 antagonist and utilised it in a series of *in vitro* experiments to uncover a distinct role for PAR4 on human platelets during thrombosis. These findings suggest PAR4 blockade may be a useful anti-thrombotic strategy. Therefore, this thesis next extended on these *in vitro* studies by investigating the effects of targeting PAR4 *in vivo* by examining the impact of PAR4-deficiency in a mouse model.

Mouse platelets express PAR3 in place of the PAR1 on human platelets, but PAR3 in the mouse does not signal, meaning transduction of thrombin signalling in mouse platelets is entirely reliant on PAR4. Platelets from PAR4<sup>-/-</sup> mice fail to respond to thrombin and PAR4<sup>-/-</sup> mice allow examination of the overall importance of platelet activation by thrombin in (patho)physiology. PAR4<sup>-/-</sup> mice are protected against arterial thrombosis yet exhibit no evidence of spontaneous bleeding. This separation between haemostasis and thrombosis is perhaps surprising. Indeed, the lack of spontaneous bleeding in PAR4<sup>-/-</sup> mice contrasts with that experienced by mice with marked thrombocytopenia, in which high rates of perinatal death due to haemorrhage are observed. Given the rationalisation of PAR4 as an anti-thrombotic drug target in Chapter 2, the studies of Chapter 3 examined in detail the potential for bleeding (perinatal haemorrhage) in PAR4<sup>-/-</sup> mice.

These studies indicate a subtle but significant effect of PAR4-deficiency on spontaneous bleeding in mice, suggesting careful consideration should be given to the safety of any PAR4

antagonists investigated. However, given the discrepancy in platelet PAR expression between mice and humans, these studies also suggest the suitability of any mouse model should be taken into consideration in assessing any *in vivo* studies relating to platelet PAR function.

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## SHORT COMMUNICATION

# Perinatal lethality of *Par4*<sup>−/−</sup> mice delivered by primiparous dams reveals spontaneous bleeding in mice without platelet thrombin receptor function

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**Abstract**

Protease-activated receptor 4 (PAR4) is a cell surface G protein-coupled receptor for serine proteases, such as thrombin. *Par4*<sup>−/−</sup> mice have platelets that are unresponsive to thrombin and thereby allow examination of the importance of thrombin-induced platelet activation in (patho)physiology. *Par4*<sup>−/−</sup> mice are protected against arterial thrombosis but show no evidence of spontaneous bleeding. This contrasts with the bleeding experienced by mice with marked thrombocytopenia, such as those with genetic deficiency of the transcription factor, nuclear factor erythroid 2 (*Nfe2l3*<sup>−/−</sup>), that have high rates of perinatal death due to hemorrhage. Given this discrepancy in spontaneous perinatal bleeding between mice without platelets and those without thrombin-induced platelet activation mechanisms, we examined in detail the immediate postnatal survival of *Par4*<sup>−/−</sup> pups. We observed significant postpartum loss of *Par4*<sup>−/−</sup> pups derived from *Par4*<sup>+/−</sup> intercrosses that was restricted to a dam's first litter; only 9% of surviving pups genotyped as *Par4*<sup>−/−</sup> in first litters and this normalized from the second litter onward (26%). A similar perinatal lethality in pups delivered by primiparous dams occurred in mice lacking platelets (*Nfe2l3*<sup>−/−</sup>; 10%) but not in those lacking fibrinogen (*Fga*<sup>−/−</sup>; 26%). These data, provide the first evidence of spontaneous bleeding in *Par4*<sup>−/−</sup> mice, suggest that a dam's first litter provides a greater hemostatic challenge than subsequent litters, and uncovers an important role for platelets—and more specifically thrombin-induced platelet activation—in hemostasis during these more traumatic births.

**Introduction**

Protease-activated receptors (PARs) are the target of the most recently-developed antiplatelet drugs [1]. PARs are attractive targets for antiplatelet agents because they mediate platelet activation by thrombin, the most potent endogenous activator. Human platelets have two PARs, PAR1, and PAR4. The first PAR1 antagonist was recently approved for the prevention of myocardial infarction and peripheral arterial disease [1] and PAR4 antagonists are in clinical trial for similar purposes [2]. Mouse platelets express non-signaling PAR3 in place of the PAR1 on human platelets, meaning transduction of thrombin signaling in mouse platelets is entirely reliant on PAR4. Platelets from *Par4*<sup>−/−</sup> mice fail to respond to thrombin [3,4] and *Par4*<sup>−/−</sup> mice allow examination of the overall importance of platelet activation by thrombin in (patho)physiology. Predictably, *Par4*<sup>−/−</sup> mice are protected against arterial thrombosis [3,5]. Yet despite this, *Par4*<sup>−/−</sup> mice exhibit no evidence of spontaneous bleeding; they are born at the expected frequency, have no obvious internal or external hemorrhage, and have a normal lifespan [4,6]. This lack of bleeding in *Par4*<sup>−/−</sup> mice contrasts with that observed in mice that have a marked thrombocytopenia, such as those with genetic deficiency of the transcription factor, nuclear factor erythroid 2 (*Nfe2l3*<sup>−/−</sup>). *Nfe2l3*<sup>−/−</sup> mice

**Keywords**

Hemostasis, perinatal hemorrhage, protease-activated receptors, thrombin

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have arrested megakaryocyte development that results in no (or very few) circulating platelets [7]. These mice with (near) absolute thrombocytopenia experience spontaneous bleeding, including significant levels of perinatal hemorrhage and subsequent death due to the hemostatic trauma of birth [7].

Given this discrepancy in the level of perinatal bleeding between mice without platelets and mice without thrombin-induced platelet activation mechanisms, we tracked in detail the genotypes of pups resulting from *Par4*<sup>+/−</sup> intercrosses. We observed significant loss of *Par4*<sup>−/−</sup> pups in the immediate postpartum period, but only in primiparous dams (i.e., the first litter from any given female). A similar perinatal lethality in pups delivered by primiparous dams occurred in mice lacking platelets (*Nfe2l3*<sup>−/−</sup>) but not in those lacking fibrinogen (*Fga*<sup>−/−</sup>), suggesting the effect was most likely primarily due to impaired thrombin-mediated platelet signaling. These data provide the first evidence of a spontaneous bleeding phenotype in *Par4*<sup>−/−</sup> mice.

**Methods**

Generation of mice deficient in PAR4 (*F2lr3* [4]), fibrinogen (*Fga* [8]), or platelets (*Nfe2l3* [7]) has been described previously. *Par4*<sup>−/−</sup> mice were either inbred ≥ 6 generations into C57BL/6 or were 50:50 C57BL/6:129/Sv as indicated. *Nfe2l3*<sup>−/−</sup> and *Fga*<sup>−/−</sup> were all 50:50 C57BL/6:129/Sv. All mice examined were derived from heterozygous intercrosses and all genotypes were confirmed by Southern blot analysis on biopsies taken from 10–14-day old pups. Genotypes of live

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offspring were stratified by an individual female's litter number. Statistical significance was tested by  $\chi^2$  analysis (GraphPad Prism 7).

## Results

In accord with previous observations [4,6], genotyping of all live births across a large number of *Par4*<sup>+/-</sup> intercrosses in an inbred C57BL/6 background revealed approximately the expected rate of production of *Par4*<sup>-/-</sup> mice (21%; 39 of 184; Table I). However, when births were stratified by the order of an individual female's litter, *Par4*<sup>-/-</sup> mice surviving to P10 were markedly reduced in the first litter (9%; 5 of 57; Table I;  $P = 0.01$ ). This significant reduction in the number of *Par4*<sup>-/-</sup> mice surviving their first 10 days was completely corrected from the second litter onward (Table I). Analysis of the number of pups born at P1-3 revealed almost no loss of mice (4 of 184) between P3 and P10. Further, the average size of the surviving first litter from *Par4*<sup>+/-</sup> intercrosses (6.3 pups per litter) was less than in subsequent litters (7.5) (Table I), closely accounting for the absent *Par4*<sup>-/-</sup> pups. Our data, when combined with the observation that *Par4*<sup>-/-</sup> embryos are present at the expected rate throughout the entirety of embryonic development [6], strongly suggest that a significant fraction of *Par4*<sup>-/-</sup> mice are selectively lost from a mother's first litter in the immediate postnatal period.

Since such an observation has not been previously reported, we next directly compared this loss of *Par4*<sup>-/-</sup> pups from primiparous dams to that experienced by mice deficient in platelets (*Nfe2*<sup>-/-</sup>) or fibrinogen (*Fga*<sup>-/-</sup>). Since neither *Nfe2*<sup>-/-</sup> nor *Fga*<sup>-/-</sup> mice survive at high rates in an inbred C57BL/6 background [7,8], we performed this head-to-head comparison in the outbred (50:50) C57BL/6:129Sv

strain, where markedly improved survival has been observed [7,8]. *Par4*<sup>+/-</sup> intercrosses in this mixed strain produced a similar loss of *Par4*<sup>-/-</sup> pups from primiparous dams (14%; 14 from 98;  $P < 0.05$ ; Table II) that again immediately corrected from the same mothers' second litter (Table II). Again, the average surviving litter size in the first litter of *Par4*<sup>+/-</sup> intercrosses was less than in subsequent litters (6.1 vs 7.3; Table II). A similar trend was observed in pups produced from *Nfe2*<sup>+/-</sup> intercrosses; *Nfe2*<sup>-/-</sup> pups represented only 10% of a mother's first litter (5 of 51;  $P < 0.05$ ; Table II) but 20% in subsequent litters (23 of 113; Table II). In contrast, the rate of *Fga*<sup>-/-</sup> pups from *Fga*<sup>+/-</sup> intercrosses was not different across the mother's litter number, with 26% in first litters (14 of 53; Table II) and 21% in subsequent litters (23 of 107; Table II).

## Discussion

This study provides the first evidence of a spontaneous bleeding phenotype in *Par4*<sup>-/-</sup> mice. Here, *Par4*<sup>-/-</sup> pups were selectively lost from litters delivered by primiparous dams. This effect was mirrored in mice deficient in functional platelets (*Nfe2*<sup>-/-</sup>) but not fibrinogen (*Fga*<sup>-/-</sup>). The specificity of this loss of knockout mice to the first litter of *Par4* and *Nfe2* mice supports the hypothesis that the first litter delivered by a given female endures a more significant hemostatic challenge during birth, and that the postnatal lethality observed here is due to impaired platelet function.

*Par4*<sup>-/-</sup> mice have platelets that do not respond to thrombin. Numerous studies have reported protection against thrombosis in *Par4*<sup>-/-</sup> mice accompanied by prolonged bleeding upon hemostatic challenge (e.g., tail transection [3]). Yet, perhaps surprisingly, there is no report of spontaneous bleeding in *Par4*<sup>-/-</sup> mice. When combined with a number of previous observations, our data indicate postnatal loss of *Par4*<sup>-/-</sup> pups rapidly after birth—a time of major hemostatic challenge. First, *Par4*<sup>-/-</sup> embryos are present at the expected rate throughout the entirety of embryonic development [6]. Second, newborn *Nfe2*<sup>-/-</sup> pups, as well as those with a combined deficiency of fibrinogen and PAR4, that do not survive the immediate postnatal period, exhibit marked intraperitoneal hemorrhage [6]. Therefore, while we did not directly measure hemostasis in the present study, it appears most likely that the loss of *Par4*<sup>-/-</sup> pups observed here in the immediate postnatal period is the result of internal hemorrhage. That this death of neonatal

Table I. Live offspring produced from *Par4*<sup>+/-</sup> intercrosses (C57BL/6).

Mother's litter no.	Genotype: observed (expected)			Total pups (litters)
	+/+	+/-	-/-	
1	20 (14.25)	32 (38.5)	5 (14.25) *	57 (9)
2	12 (13.5)	29 (27)	13 (13.5)	54 (7)
3	12 (12.75)	23 (25.5)	16 (12.75)	51 (6)
4	5 (5.5)	12 (11)	5 (5.5)	22 (4)
≥2 combined	29 (31.75)	64 (63.5)	34 (31.75)	127 (17)
Total	49 (46)	96 (92)	39 (46)	184 (26)

\*Overall distribution is significantly different from expected by chi-square analysis ( $P < 0.05$ ).

Table II. Live offspring produced from *Par4*<sup>+/-</sup>, *Nfe2*<sup>+/-</sup>, or *Fga*<sup>+/-</sup> intercrosses (C57BL/6 x 129/Sv).

Mother's litter no.	Genotype: observed (expected)			Total pups (litters)
<i>Par4</i>	+/+	+/-	-/-	
1	29 (24.5)	55 (49)	14 (24.5) *	98 (16)
≥2	45 (48.25)	102 (96.5)	44 (48.25)	191 (26)
Total	74 (72.25)	157 (144.5)	58 (72.25)	289 (42)
<i>Nfe2</i>	+/+	+/-	-/-	
1	13 (12.75)	33 (25.5)	5 (12.75) *	51 (9)
≥2	29 (28.25)	61 (56.5)	23 (28.25)	113 (17)
Total	41 (41)	94 (82)	29 (41)	164 (26)
<i>Fga</i>	+/+	+/-	-/-	
1	15 (13.25)	24 (26.5)	14 (13.25)	53 (9)
≥2	33 (26.75)	51 (53.5)	23 (26.75)	107 (16)
Total	48 (40)	75 (80)	37 (40)	160 (25)

\*Overall distribution is significantly different from expected by chi-square analysis ( $P < 0.05$ ).

*Par4*<sup>-/-</sup> pups is restricted to those in a mother's first litter suggests the impact of PAR4-deficiency is heavily dependent on the strength of the hemostatic challenge, with the hemostatic trauma of birth presumably greater during a primiparous delivery than in subsequent deliveries.

The observed perinatal loss of *Par4*<sup>-/-</sup> pups appears most likely platelet-dependent given that a similar observation was noted with *Nfe2*<sup>-/-</sup> pups. However, as PAR4 is expressed on a number of other cell types, including some that may contribute to hemostasis such as vascular endothelial cells, we cannot formally rule out a contribution from cells other than platelets to the bleeding observed here. On this point, it is worth noting that a previous study used bone marrow transplantation to show that the impaired hemostasis in *Par4*<sup>-/-</sup> mice after injury (prolonged bleeding after tail clip) is fully attributable to PAR4 deficiency in platelets [3]. Therefore, the present study is highly suggestive of an important role for platelets—and more specifically thrombin-induced platelet activation—in hemostasis during traumatic births. Such a phenotype has not been previously described, and whether or not this observation extends to other commonly-used mouse lines with major platelet defects remains unknown. For example, PAR3 is a thrombin co-factor receptor on mouse platelets. Platelets from *Par3*<sup>-/-</sup> mice exhibit reduced sensitivity to thrombin that is sufficient to impair hemostasis—but this impairment is notably less than that experienced by *Par4*<sup>-/-</sup> mice [9]. It will therefore be of interest to determine if the more subtle impairment in platelet function found in, for example, *Par3*<sup>-/-</sup> mice is sufficient to cause a similar perinatal hemorrhage and death in pups from primiparous dams to that observed here in *Par4*<sup>-/-</sup> mice.

Finally, mice without thrombin (prothrombin-deficient mice; *F2*<sup>-/-</sup>) exhibit 100% embryonic lethality—approximately half of which occurs during mid-gestation due to a PAR1-dependent defect in vascular patterning [10] and the remainder due to hemorrhage at birth [11]. This latter effect is presumably due to the combined input of thrombin to platelet activation and fibrin formation, since both *Fga*<sup>-/-</sup> and *Par4*<sup>-/-</sup> mice are unaffected to term but combined genetic deficiency recapitulates the hemostatic effect of prothrombin-deficiency with almost all *Par4*<sup>-/-</sup>:*Fga*<sup>-/-</sup> pups experiencing fatal postnatal internal hemorrhage [6]. The more detailed analysis of hemostasis at birth performed here indicates thrombin-induced platelet activation may be more relevant to birth trauma hemostasis than thrombin-induced fibrin formation. With this in mind, our study suggests that any impact of PAR inhibition on spontaneous bleeding may be heavily dependent on the strength of the hemostatic challenge.

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## Declaration of interest

The authors report no declarations of interest.

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## **Chapter 4 - Humanising the protease-activated receptor (PAR) expression profile in mouse platelets by knocking PAR1 into the Par3 locus reveals PAR1 expression is not tolerated in mouse platelets**

### **4.1 Introduction**

The studies of Chapter 2 provide rationale for the preclinical evaluation of PAR4 antagonism as an antithrombotic approach. However the studies in Chapter 3 highlight the need for an animal model mimicking human platelet PAR expression, so that appropriate safety and efficacy studies can be performed *in vivo*. Since only primate platelets express the PAR1/PAR4 profile, there are significant limits on preclinical *in vivo* research models allowing the requisite investigation into the utility of PAR4 antagonists. The studies of Chapter 4 attempted to address this. Specifically, these studies sought to overcome the limitation of current mouse models by ‘humanising’ the platelet PAR expression profile of the mouse, essentially by replacing PAR3 with PAR1. Here, it was theorised that ‘knock-in’ of PAR1 into the endogenous mouse *Par3* locus in a manner allowing platelet-specific expression would create a mouse with PAR1 and PAR4 on platelets but without ectopic PAR1 expression. However, for reasons that are not yet clear, this approach failed. Taken together with earlier studies using distinct approaches, this study indicates forced expression of PAR1 in mouse platelets is difficult to achieve and suggests other options are required for reliable preclinical screening of PAR antagonists.

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RESEARCH ARTICLE

# Humanizing the Protease-Activated Receptor (PAR) Expression Profile in Mouse Platelets by Knocking *PAR1* into the *Par3* Locus Reveals PAR1 Expression Is Not Tolerated in Mouse Platelets

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## Abstract

Anti-platelet drugs are the mainstay of pharmacotherapy for heart attack and stroke prevention, yet improvements are continually sought. Thrombin is the most potent activator of platelets and targeting platelet thrombin receptors (protease-activated receptors; PARs) is an emerging anti-thrombotic approach. Humans express two PARs on their platelets—PAR1 and PAR4. The first PAR1 antagonist was recently approved for clinical use and PAR4 antagonists are in early clinical development. However, pre-clinical studies examining platelet PAR function are challenging because the platelets of non-primates do not accurately reflect the PAR expression profile of human platelets. Mice, for example, express *Par3* and *Par4*. To address this limitation, we aimed to develop a genetically modified mouse that would express the same repertoire of platelet PARs as humans. Here, human *PAR1* preceded by a lox-stop-lox was knocked into the mouse *Par3* locus, and then expressed in a platelet-specific manner (hPAR1-KI mice). Despite correct targeting and the predicted loss of *Par3* expression and function in platelets from hPAR1-KI mice, no PAR1 expression or function was detected. Specifically, PAR1 was not detected on the platelet surface nor internally by flow cytometry nor in whole cell lysates by Western blot, while a PAR1-activating peptide failed to induce platelet activation assessed by either aggregation or surface P-selectin expression. Platelets from hPAR1-KI mice did display significantly diminished responsiveness to thrombin stimulation in both assays, consistent with a *Par3*<sup>-/-</sup> phenotype. In contrast to the observations in hPAR1-KI mouse platelets, the PAR1 construct used here was successfully expressed in HEK293T cells. Together, these data suggest ectopic PAR1 expression is not tolerated in mouse platelets and indicate a different approach is required to develop a small animal model for the purpose of any future preclinical testing of PAR antagonists as anti-platelet drugs.



**Competing Interests:** The authors have declared that no competing interests exist.

## Introduction

Anti-platelet drugs are the primary therapy for heart attack and stroke prevention, yet improvements are continually sought. Thrombin is by far the most potent endogenous platelet activator, which it achieves via cell surface protease-activated receptors (PARs). Due to the importance of thrombin-induced platelet activation during thrombosis, targeting platelet thrombin receptors has received significant clinical attention, and PAR antagonists are one of the most promising of the emerging anti-thrombotic approaches [1–3]. Humans express two PARs on their platelets, PAR1 and PAR4, both of which are cleaved and activated by thrombin, and both of which are capable of inducing robust platelet activation [4, 5]. PAR1 has a higher affinity for thrombin and was therefore the target of initial drug development. The first PAR1 antagonist, vorapaxar, was recently approved by the FDA for clinical use in the USA [2]. So far, however, vorapaxar has limited clinical utility due to an increase in major bleeding events associated with its use [6, 7]. These limitations on the first PAR1 inhibitor have ignited interest in evaluating the clinical potential of PAR4 antagonists, and these agents are now in early clinical development (NCT02208882) [8].

However, a major limitation in examining platelet PAR function in detail is the absence of a small animal model that accurately reflects the PAR expression profile in human platelets: only primate platelets express the PAR1/PAR4 profile [9]. This places significant limits on preclinical research into the utility of PAR antagonists *in vivo*. Of the commonly-used small animals, for example, platelets from mice, rats and rabbits express PAR3 and PAR4 [10–13], while guinea-pig platelets express PAR1, PAR3 and PAR4 [14]. *Par4*<sup>-/-</sup> mice have been crucial for early proof-of-concept studies into the overall role of PARs in thrombosis [15–17], but are unsuitable for elucidating, for example, the relative anti-thrombotic effects of PAR1 vs PAR4 inhibition.

One way to overcome this limitation is to ‘humanize’ the platelet PAR expression profile of the mouse, essentially replacing mouse PAR3 (Par3) with human PAR1 (PAR1). Initial attempts used platelet-specific expression of a PAR1 transgene in *Par3*<sup>-/-</sup> mice, but were unsuccessful [18]. Possible explanations for the failure of this transgene-based approach include insufficient expression levels or unpredicted gene silencing consequences as a result of nonspecific transgene insertion [18]. We theorized that ‘knock-in’ of PAR1 into the endogenous mouse *Par3* locus in a manner that would allow platelet-specific expression may overcome these earlier issues and would create a mouse that expresses PAR1 and Par4 on platelets, but without further ectopic PAR1 expression. Here, we generate and characterise such a mouse. Despite correct targeting, this genetic approach failed to yield detectable expression or function of PAR1 on mouse platelets. When taken together with earlier studies using distinct approaches, this study indicates that forced expression of PAR1 in mouse platelets will be difficult to achieve and suggests that other options will be required for reliable preclinical screening of any future PAR antagonists.

## Methods and Materials

### Materials

Human  $\alpha$ -thrombin and adenosine 5'-diphosphate sodium salt (ADP) were purchased from Sigma-Aldrich (St Louis, MO, USA). PAR1-activating peptide (PAR1-AP; TFLLR-NH<sub>2</sub>) and PAR4-activating peptide (PAR4-AP; AYPGKF-NH<sub>2</sub>) were synthesised at Monash Institute of Pharmaceutical Sciences (Melbourne, Australia) by Assoc Professor Philip Thompson. The mouse anti-P-selectin (FITC anti-mouse CD62P) and human anti-P-selectin (PE anti-human CD62P) antibodies were purchased from BD Biosciences (San Jose, CA, USA). PE anti-CD45.2,

PE anti-CD41a (human) antibody and PE anti-CD41 antibody (mouse) were purchased from Abcam (Melbourne, Australia). The PE-anti-GPIIb-tail and non-immune rabbit IgG isotype were a generous gift from Assoc Professor Robert Andrews (Monash University, Melbourne, Australia). The anti-PAR1 antibody (ATAP2), anti-PAR3 antibody (8E8), HRP-anti-actin (I-19) and HRP- anti-mouse IgG were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

### Generation of hPAR1-KI (*Pf4-Cre; Par3<sup>LSL-PAR1</sup>*) mice

All mouse studies were approved by the Alfred Medical Research and Education Precinct Animal Ethics Committee (approval number E/1465/2014/M). *Pf4-Cre; Par3<sup>LSL-PAR1</sup>* mice (hereafter referred to as hPAR1-KI mice) were generated by Ozgene (Perth, Australia). The targeting vector involved exon 2 of *Par3* (*F2rl2*) flanked with a lox-stop-lox and followed by exon 2 of *PAR1* (*F2r*) (Fig 1A). Since exon 1 encodes the signal peptide and exon 2 encodes the entire mature receptor for both PAR1 and Par3 [19], Cre-mediated deletion is predicted to result in the replacement of Par3 with PAR1 and leaves the endogenous *Par3* promoter intact (Fig 1A). The BAC vector comprised unaltered exon 1 of *Par3*, then exon 2 of *Par3* followed by a stop codon and flanked by loxP sites, and followed by *PAR1*. The vector was injected into C57BL/6J blastocysts, chimeras were bred for germline transmission, and the targeted allele bred to homozygosity (*Par3<sup>LSL-PAR1</sup>*). These mice were then bred with *Pf4-Cre* mice to induce deletion of *Par3* and permit the expression of *PAR1* specifically in platelets (*Pf4-Cre; Par3<sup>LSL-PAR1</sup>*, hereafter referred to as hPAR1-KI mice). PCR genotyping of the targeted allele was performed using the following primers: common reverse (AGCTGAAAAATGGAGCGCTTG) with WT forward (TGGGTTCTCTCATCCCGTTTG, predicted size 588bp) or mutant forward (TCCTTCACTTGTCTGGCCATC, predicted size 914bp) (Fig 1A). Deletion of *Par3* was confirmed by PCR using the following primers: forward (CAGTGTGTG GTTTTGTTTCACCT) and reverse (GCCAATCACTGCCGGAAG, predicted size 963bp) (Fig 1A).

### Isolation of mouse and human platelets

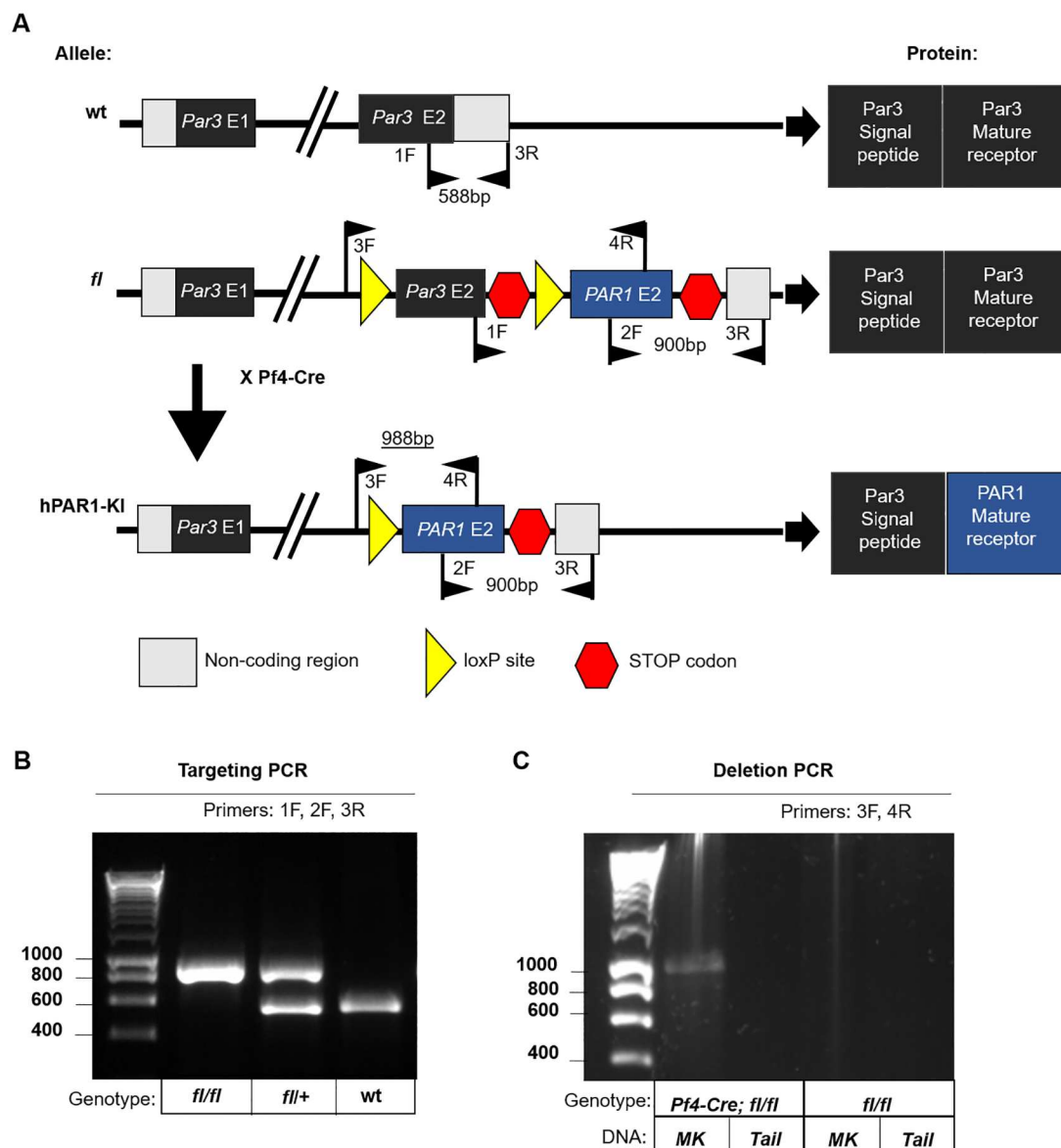
All human studies were approved by the Monash University Human Research Ethics Committee (CF07/0141-2007/0025). Blood was collected after written informed consent was obtained from healthy adults (21–50 years old, of both sexes) who had not taken anti-platelet medications in the past 10 d. Mouse and human blood was drawn into syringes containing acid citrate dextrose (ACD; 1:7 v/v). Platelets were isolated by centrifugation as previously described [20].

### Megakaryocyte DNA extraction

Bone marrow cells were aseptically flushed from the femora and tibiae of mice using a syringe and a 23-gauge needle containing Dulbecco's Modified Eagle Medium (DMEM, Life Technologies) supplemented with 10% foetal bovine serum (FBS), penicillin, streptomycin and glutamine. Post-harvest, thrombopoietin (0.1 µg/mL) was added to the cells, which were allowed to culture for 4 d at 37°C, 5% CO<sub>2</sub>. Cells were then collected and centrifuged at 50 g for 5 min and the pellet was re-suspended in 4 mL of PBS. The cells were layered over a two-step BSA density gradient and allowed to sediment for 45 min at room temperature. The megakaryocyte-rich population was collected and centrifuged at 50 g for 5 min, and the pellet was treated with DNA extraction buffer (Bioline) and digested at 75°C for 10 min.

### Platelet aggregation

Platelet aggregation was measured by light transmission aggregometry in a 96-well plate format. Human and mouse washed platelets (2x10<sup>8</sup> platelets/mL) were added to wells of a clear, flat



**Fig 1. Generation of mice expressing PAR1 in place of Par3 in a platelet-specific manner.** (A) Schematic of the strategy for platelet-specific expression of human PAR1 in place of mouse Par3. Top: wild-type *Par3* (wt) has two exons: exon 1 encodes the signal peptide and exon 2 encodes the complete mature receptor. Middle: In the targeted allele (*Par3*<sup>LSL-PAR1</sup>; fl) an insert of exon 2 of human *PAR1* is placed downstream of exon 2 of *Par3*. *Par3* exon 2 is flanked by loxP sites, allowing Cre-mediated excision. Bottom: Cre-mediated deletion of *Par3* exon 2 allows expression of *PAR1* exon 2, resulting in a predicted chimeric protein consisting of the Par3 signal peptide fused to the mature PAR1 receptor (hPAR1-KI). Heterozygous targeted mice, *Par3*<sup>LSL-PAR1/+</sup> (fl/+); homozygous targeted mice, *Par3*<sup>LSL-PAR1/LSL-PAR1</sup> (fl/fl); platelet-specific replacement of mPAR3 with hPAR1, Pf4-Cre; *Par3*<sup>LSL-PAR1</sup> (Pf4-Cre;fl/fl). (B) PCR genotyping of tail DNA using the primers indicated (black flags): wild type forward (1F), mutant forward (2F), common reverse (3R). (C) Cre-mediated deletion of *Par3* was confirmed by PCR genotyping of individual mice using DNA from both tail (not deleted) and megakaryocytes (MK; deleted), using a forward primer spanning the loxP site (3F) and *PAR1* reverse (4R). A PCR product of predicted size (988bp) was only generated using MK DNA from Pf4-Cre;fl/fl mice. Sequencing of this PCR product confirmed excision of floxed sequences (S3 Fig).

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bottom, 96-well tissue culture plate (Falcon). Platelets were stimulated with either thrombin (0.01–1 U/mL), PAR1-AP (10–100  $\mu$ M), PAR4-AP (10–100  $\mu$ M), or ADP (1–10  $\mu$ M). The plate was analysed at 37°C in a FLUOstar OPTIMA plate reader (BMG Labtech) using a 595 nm excitation filter, for a period of 50 min (10 read cycles with 5 min double orbital shake period between each read). Ability to aggregate was calculated as  $\frac{OD(No\ Agonist) - OD(Agonist)}{OD(No\ Agonist) - OD(blank)} \times 100$  at the time point where aggregation was at a maximum. Optical density was normalised against the blank (maximum) and unstimulated platelets (minimum) and expressed as % maximum.

## P-selectin expression

Flow cytometry was used to detect the expression of P-selectin by activated platelets. Human and mouse platelets, prepared as outlined above, were incubated with an anti-P-selectin antibody and stimulated with thrombin (0.01–1 U/mL), PAR1-AP (10–100  $\mu$ M), or PAR4-AP (30–300  $\mu$ M) for 15 min at 37°C. Samples were then analysed using a FACSCalibur (Becton Dickinson) flow cytometer and FlowJo software.

## Analysis of platelet PAR1 expression

Platelets isolated from either humans or mice ( $5 \times 10^7$  platelets/mL in Tyrode's buffer) were incubated with a PE-conjugated antibody against one of PAR1 (2.5  $\mu$ g/ $10^6$  cells; ATAP2), CD45 (platelet negative control antibody of matching isotype, 2.5  $\mu$ g/ $10^6$  cells), or CD41a (platelet positive control antibody of matching isotype, 2.5  $\mu$ g/ $10^6$  cells). In some experiments, isolated platelets were permeabilized with saponin (0.1% v/v) prior to incubation with antibodies. In these experiments, a PE-conjugated antibody against the cytoplasmic tail of GPIIb was included to confirm effective platelet permeabilization [21]. In all cases, samples were analysed using a FACSCalibur flow cytometer and FlowJo Software. Positive expression was measured as a rightward shift in relative fluorescent intensity compared to the isotype control.

## Western blot

Platelet proteins were analyzed in lysates of human or mouse isolated platelets ( $1 \times 10^9$  platelets/mL), prepared in Laemmli's buffer, denatured (95°C for 10 min), and run on 12% SDS-polyacrylamide gels at 170 V for 40 min. Proteins were transferred to PVDF membranes (Millipore, Lake Placid, NY, USA) at 250 mA for 2 h. Membranes were blocked with TBST (0.01 mM Tris, 50 mM NaCl, 0.01% v/v Tween-20) containing 5% skim milk powder for 30 min at room temperature, incubated with either an HRP-conjugated anti-actin antibody (1  $\mu$ g/mL), anti-PAR1 antibody (1  $\mu$ g/mL) or anti-PAR3 antibody (1  $\mu$ g/mL) at 4°C overnight, followed by a HRP-conjugated anti-mouse IgG at room temperature for 2 h. Enhanced chemiluminescence (ECL) substrate (ThermoFisher Scientific, Waltham, MA, USA) was placed onto the membrane for 1 min before exposure (ChemiDoc Touch Imaging System, Bio-Rad, Hercules, CA, USA).

## Results

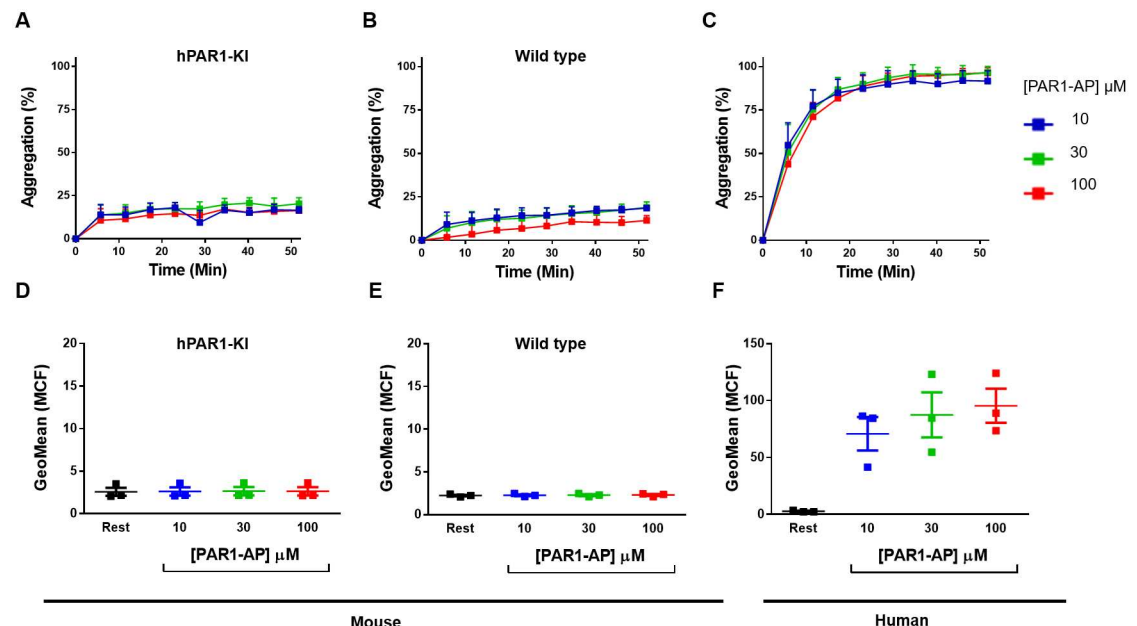
### Generation of hPAR1-KI (*Pf4-Cre; Par3<sup>LSL-PAR1</sup>*) mice

We generated mice in which *PAR1* (*F2R*) was knocked into the *Par3* (*F2lr2*) locus and conditionally expressed in platelets upon *Pf4-Cre*-mediated excision. To this end, exon 2 of *PAR1* (encoding the entire mature receptor) was inserted downstream of exon 2 of *Par3* flanked by a lox-stop-lox (Fig 1A). This approach leaves the promoter and exon 1 of *Par3* (encoding the signal peptide) intact and results in replacement of *Par3* with *PAR1* after *Cre*-mediated excision (Fig 1A). This is predicted to yield the *Par3* signal peptide linked to the complete mature

receptor of PAR1 (Fig 1A). The stability and surface expression of the predicted fusion protein was confirmed in HEK293T cells transiently transfected with vectors expressing the same sequence as that remaining after Cre-mediated excision of the targeted allele, with the predicted Par3/PAR1 fusion protein expressed at similar levels to native PAR1 (S1 Fig). We therefore went on to generate *Par3*<sup>LSL-PAR1/+</sup> mice and confirmed targeting by Southern blot (S2 Fig) and PCR genotyping (Fig 1B). Heterozygous mice were bred to homozygosity (*Par3*<sup>LSL-PAR1/LSL-PAR1</sup>) and crossed with Pf4-Cre mice [22], with experimental mice resulting from crosses of *Pf4-Cre;Par3*<sup>LSL-PAR1/LSL-PAR1</sup> x *Par3*<sup>LSL-PAR1/LSL-PAR1</sup> (Fig 1B). Cre positive offspring (*Pf4-Cre;Par3*<sup>LSL-PAR1/LSL-PAR1</sup>) are hereafter referred to as hPAR1-KI mice and their Cre negative littermates (*Par3*<sup>LSL-PAR1/LSL-PAR1</sup>) as wild-type. Appropriate Pf4-Cre-mediated deletion in the platelet/megakaryocyte lineage was confirmed by PCR of DNA isolated from megakaryocytes versus tail biopsy (Fig 1C). We used a forward primer placed upstream of the loxP site and a reverse primer in each of *Par3* and *PAR1* as indicated (Fig 1A). As predicted, a band of expected size was only generated in megakaryocyte DNA from hPAR1-KI mice. Sequencing of this PCR product confirmed the predicted targeting and Cre-mediated deletion (S3 Fig).

## Platelets from hPAR1-KI mice do not respond to PAR1-selective activation

Mice from these crosses were born at the expected rates, were normal in weight and appearance, and had normal whole blood counts (S1 Table). We assessed PAR1 function in platelets isolated from hPAR1-KI mice using two distinct markers of platelet activation: platelet aggregation and P-selectin expression. PAR1-selective activation by a PAR1-AP failed to aggregate platelets from hPAR1 mice at concentrations up to 100  $\mu$ M (100 times the concentration required to cause platelet activation in human platelets [23]) (Fig 2A). This lack of response



**Fig 2. Platelets from hPAR1-KI mice do not respond to PAR1-selective activation.** Platelets isolated from hPAR1-KI mice, wild type mice, or humans were stimulated with a PAR1-activating peptide (PAR1-AP; TFLLR) and examined for (A–C) platelet aggregation by light transmission aggregometry or (D–F) P-selectin expression by flow cytometry. Note that platelets from either hPAR1-KI or wild type mice failed to respond to PAR1-AP, even at concentrations that were supra-maximal in human platelets. Data are mean  $\pm$  SEM of  $n = 3$  experiments.

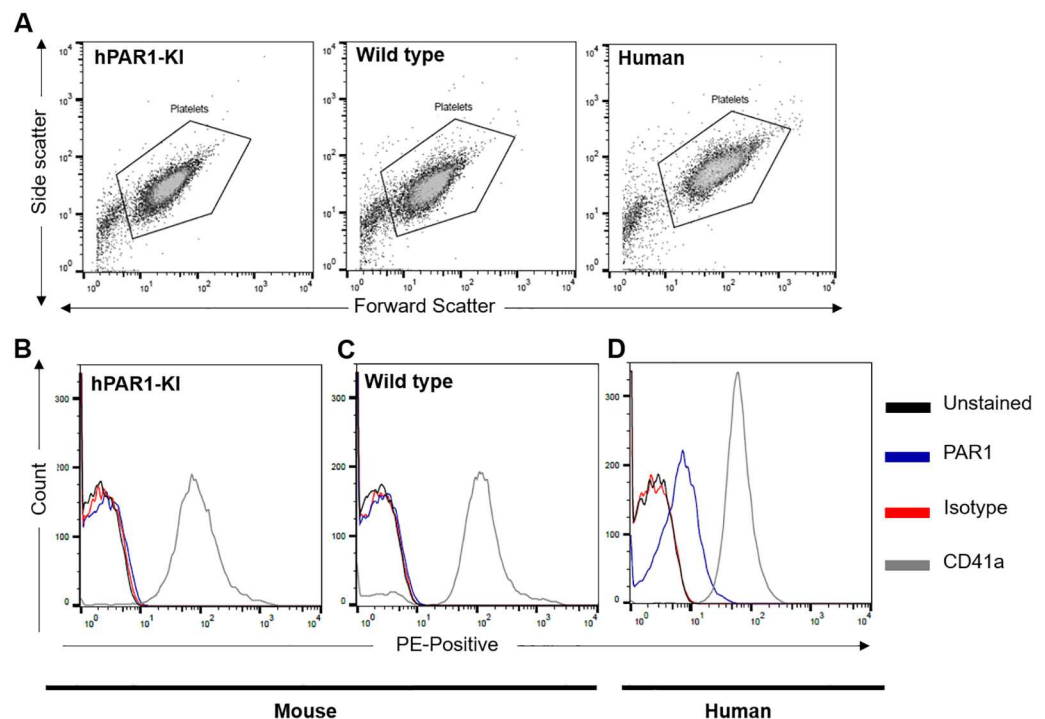
doi:10.1371/journal.pone.0165565.g002

was identical in platelets from wild type mice (Fig 2B). As expected, PAR1-AP induced a robust aggregation response in human platelets (Fig 2C). A similar pattern of responses was observed when P-selectin expression was used as the marker of platelet activation, with PAR1-AP unable to elicit any response in platelets isolated from either hPAR1-KI (Fig 2D) or wild type (Fig 2E) mice, but a robust response in human isolated platelets (Fig 2F). Together, these data indicate lack of functional PAR1 in platelets from hPAR1-KI mice.

### Lack of PAR1 expression on platelets from hPAR1-KI mice

We next examined PAR1 expression in platelets from hPAR1-KI mice. We were unable to detect PAR1 on the surface of platelets from hPAR1-KI mice by flow cytometry (Fig 3A). As with the functional experiments, platelets from wild type mice and humans served as effective negative and positive controls for PAR1 surface expression by this method, respectively (Fig 3B and 3C). In all experiments, the platelet specific marker, CD41, and normal forward/side scatter properties were used to confirm the integrity of the platelets being assessed (Fig 3).

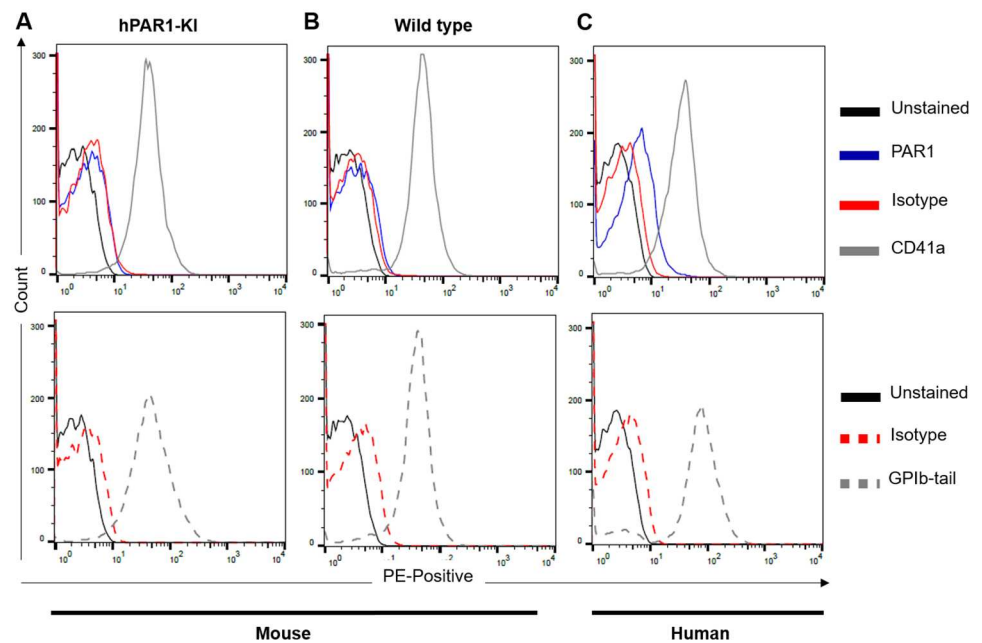
We also tested for PAR1 expression in permeabilized platelets. In these experiments, an antibody against the cytoplasmic tail of GPIIb/IIIa [21] was used as a positive control for successful permeabilization. As predicted, this antibody only recognised an epitope on the internal surface of the platelet membrane, with no staining of intact platelets and strong staining in permeabilized platelets from both mice and humans (S4 Fig). However, we were unable to detect any PAR1 in permeabilized platelets from hPAR1-KI mice (Fig 4A). Again, platelet integrity was confirmed by CD41 expression in all cases, while permeabilization was confirmed by



**Fig 3. PAR1 is undetectable on the surface of platelets from hPAR1-KI mice.** Platelets isolated from hPAR1-KI mice, wild type mice, or humans were either left unstained (black) or were incubated with a PE-conjugated antibody against one of PAR1 (blue) or CD41a (grey; positive platelet control), or an isotype control (red). (A) Platelets were gated by forward and side scatter as shown. (B–D) PAR1 was detected on human platelets but not platelets from either hPAR1-KI or wild type mice. Data shown are representative traces of  $n = 3$  individual experiments.

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**Fig 4. PAR1 is undetectable within platelets from hPAR1-KI mice.** Platelets isolated from (A) hPAR1-KI mice, (B) wild type mice, or (C) humans were fixed and permeabilized prior to incubation with a PE-conjugated antibody against one of PAR1 (blue), CD41a (grey; positive control for platelets), isotype (red), or the cytoplasmic tail of GPIIb (grey dash; positive control for permeabilization). PAR1 was detected in human platelets but not in platelets from either hPAR1-KI or wild type mice. Data shown are representative traces of  $n = 3$  individual experiments.

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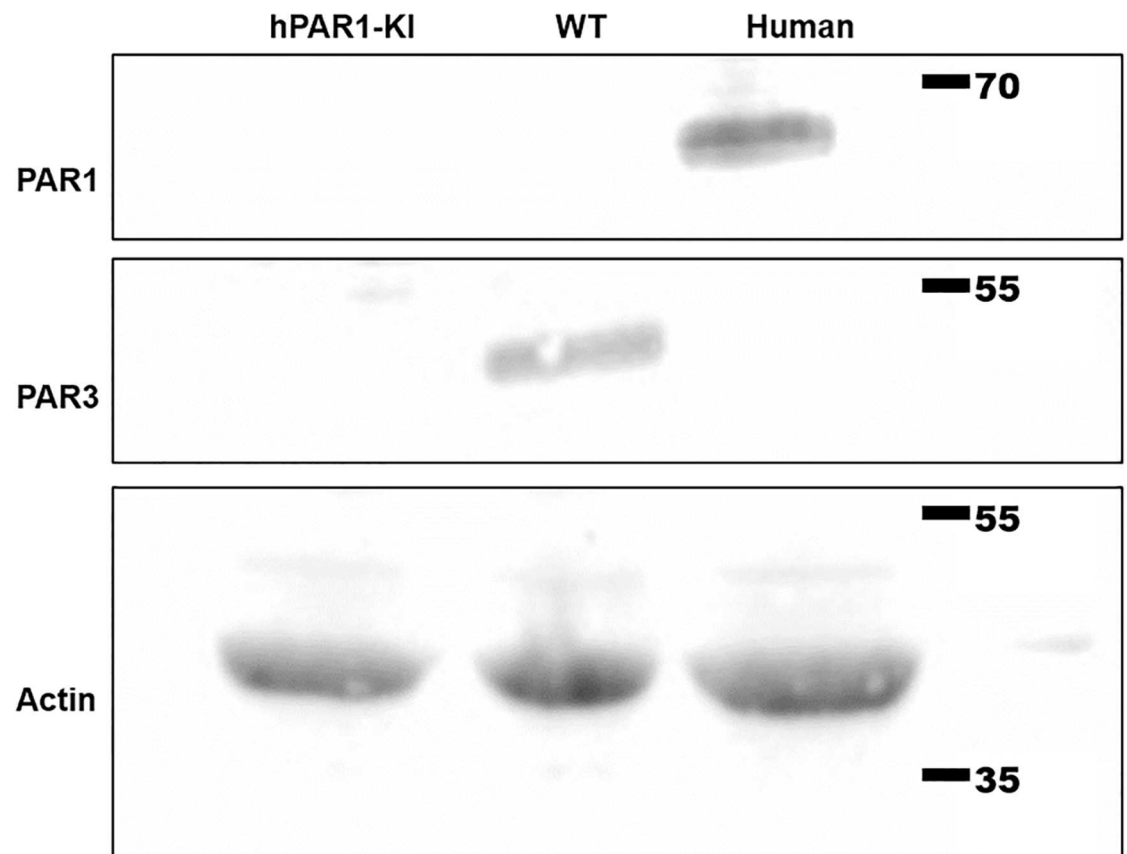
detection with the anti-GPIIb-tail antibody (Fig 4). Platelets from wild type mice and humans again served as controls (Fig 4B and 4C), indicating the anti-PAR1 antibody still binds its epitope in permeabilized cells.

Finally, we also examined PAR1 expression by Western blot. Here, a band at the predicted size of approximately 75 kD was detected in lysates of human platelets, but not of wild-type or hPAR1-KI mouse platelets (Fig 5), mirroring the flow cytometry data. Together, these findings indicate PAR1 is not expressed at detectable levels in platelets from hPAR1-KI mice.

## PAR3 expression and function is disrupted in platelets from hPAR1-KI mice

Our genetic strategy is predicted to disrupt Par3 expression in platelets. This was confirmed genetically by PCR genotyping and sequencing (Fig 1). Regardless, given the lack of PAR1 expression or function in platelets from hPAR1-KI mice, we therefore also probed for Par3 protein in these platelets. We never detected Par3 expression by Western blot in human platelets, but routinely observed a band of the previously reported size (approximately 50kDa) in wild type mouse platelets (Fig 5). This band was absent in platelet lysates from all hPAR1-KI mice examined (Fig 5), further confirming disruption of *Par3* in hPAR1-KI mice. To further examine this, we tested for the predicted functional effects of *Par3*-deficiency.

Platelets from *Par3*<sup>-/-</sup> mice exhibit diminished responsiveness to thrombin [24]. Here, we observed a similar effect in platelets from hPAR1-KI mice. We examined platelet aggregation and P-selectin expression in response to thrombin. Platelets from hPAR1-KI mice exhibited an approximately 3-fold decrease in sensitivity to thrombin in both assays when compared to platelets from wild type mice (Fig 6). Specifically, aggregation of hPAR1-KI platelets was only



**Fig 5. Protein for both PAR1 and Par3 is undetectable in platelets from hPAR1-KI mice.** Whole cell lysates of platelets isolated from hPAR1-KI mice, wild type mice, or humans were probed for either PAR1 or Par3 protein by Western blot. PAR1 was readily detected in human platelets but not in platelets from wild type or hPAR1-KI mice. In the same samples, Par3 was detected in platelets from wild type mice but not in platelets from either hPAR1-KI mice or humans. Actin was used as a protein loading control for all lysates. Blots shown are representative of  $n = 3$  experiments.

doi:10.1371/journal.pone.0165565.g005

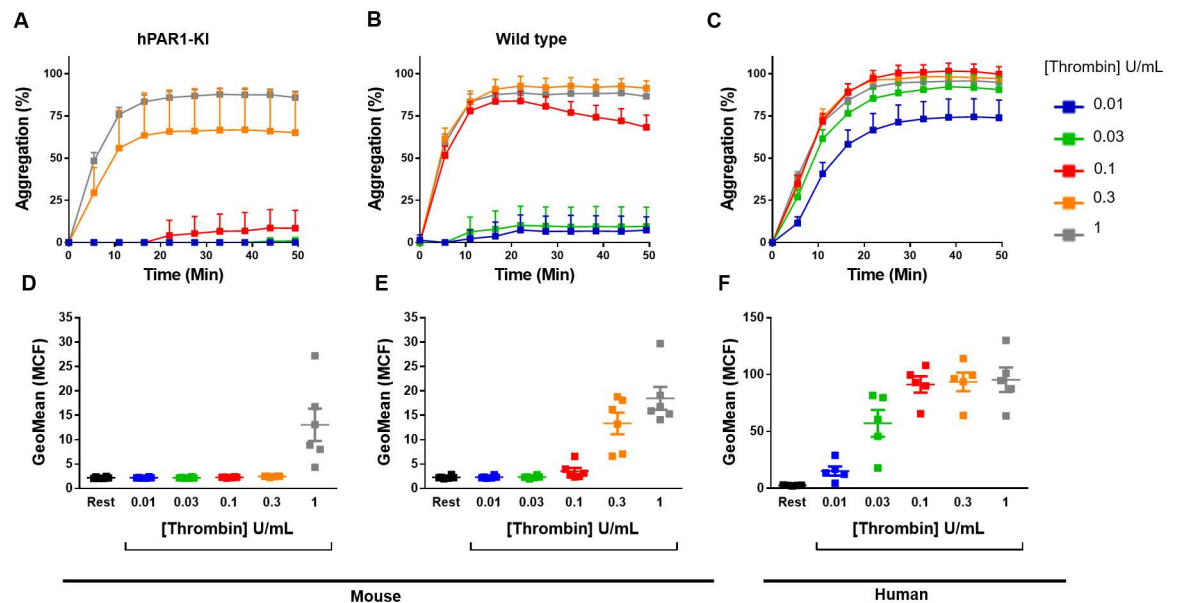
achieved at thrombin concentrations of 0.3 U/mL and above (Fig 6A) compared with 0.1 U/mL and above for wild type platelets (Fig 6B). A similar pattern emerged when examining P-selection expression in response to thrombin: aggregation of hPAR1-KI platelets was only achieved at thrombin concentrations of 1 U/mL (Fig 6D) compared with 0.3 U/mL and above for wild type platelets (Fig 6E).

Importantly, hPAR1-KI platelets responded normally to other agonists, PAR4-AP and ADP (Fig 7). When comparing platelets from hPAR1-KI and wild type mice, near-identical concentration-responses were observed for ADP-induced aggregation (Fig 7A and 7B), PAR4-AP-induced aggregation (Fig 7D and 7E), and PAR4-AP-induced P-selectin expression (Fig 7G and 7H). These functional data are consistent with the phenotype observed in platelets from *Par3*<sup>-/-</sup> mice and, when combined with the failure to detect Par3 expression in hPAR1-KI platelets, suggest correct targeting and *Par3* deletion in hPAR1-KI mice.

## Discussion

PARs are leading targets for new anti-platelet drugs, exemplified by the recent approval of the PAR1 antagonist, vorapaxar, and the early clinical development of the PAR4 antagonist, BMS-986120. Given the increased focus of examining PAR antagonists as anti-platelet agents, there





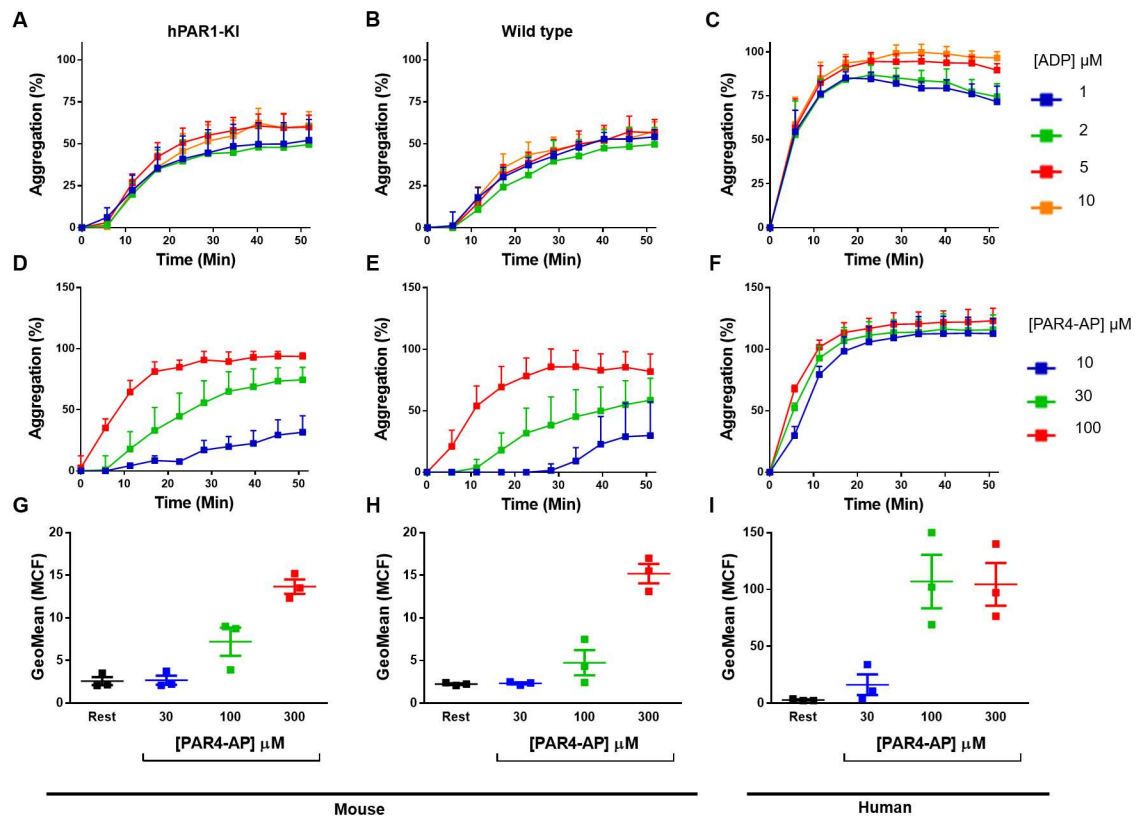
**Fig 6. Platelets from hPAR1-KI mice exhibit diminished responses to thrombin.** Platelets isolated from (A,D) hPAR1-KI mice, (B,E) wild type mice, or (C,F) humans were with thrombin and examined for (A–C) platelet aggregation by light transmission aggregometry or (D–F) P-selectin expression by flow cytometry. Note the approximate 3-fold decrease in sensitivity to thrombin-induced responses in platelets from hPAR1-KI mice versus wild-type mice in both assays. Data are mean  $\pm$  SEM of  $n = 4$ –7 experiments.

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is much interest in developing an appropriate small animal model for *in vivo* testing of such drugs. This has proved to be a major limitation in the field, as the PAR1/PAR4 expression profile of human platelets is only shared amongst primates. Here, we attempted to create a genetically-modified mouse that mimics the PAR1/PAR4 expression profile of human platelets. To do this, we knocked the human *PAR1* gene in to the mouse *Par3* locus in a manner that provided for platelet-specific replacement of *Par3* with PAR1. Correct targeting was confirmed genetically and was apparent functionally. Yet despite this, we were unable to detect expression or function of PAR1 in platelets from these mice, indicating the approach used here is insufficient to ‘humanize’ platelet PAR expression in mice by inducing PAR1 expression in place of *Par3*.

Forced PAR1 expression has been previously achieved using transgene-based approaches in other cell types. For example, mouse PAR1 (*Par1*) was specifically expressed in endothelial cells via the *Tie2* promoter/enhancer [25] and in cardiomyocytes via the  $\alpha$ MHC promoter [26]. In addition, PAR1 has been expressed in mouse mammary gland epithelium [27]. In contrast to these successes, a previous attempt failed to express either *Par1* or PAR1 transgenes in mouse platelets via multiple platelet-specific promoters [18]. The reasons for the inability to express PAR1 in mouse platelets remain unknown. However, given the limitations and lack of control associated with transgene-based approaches, we used a knock-in approach in which *PAR1* was targeted to the *Par3* locus, essentially resulting in replacement of *Par3* with PAR1, and with expression governed by the endogenous mouse *Par3* promoter.

Correct genetic targeting was confirmed by multiple methods, including Southern blot analysis of founders, PCR genotyping across the insert, and PCR analysis of correct Pf4-Cre-mediated excision in the DNA in isolated megakaryocytes. Correct targeting was also apparent by the absence of *Par3* expression in platelets from hPAR1-KI mice. The disruption of *Par3* expression was further supported by functional analyses of these platelets that revealed a



**Fig 7. Platelets from hPAR1-KI mice respond normally to PAR1-independent platelet agonists.** Platelets isolated from (A,D,G) hPAR1-KI mice, (B,E,H) wild type mice, or (C,F,I) humans were stimulated with either (A–C) ADP or (D–I) a PAR4-activating peptide (PAR4-AP) and assessed by (A–F) platelet aggregation or (G–I) P-selectin expression. Note the near-identical concentration-dependent responses in platelets from hPAR1-KI and wild type mice in all cases examined. Data are mean  $\pm$  SEM of  $n = 3$  experiments.

doi:10.1371/journal.pone.0165565.g007

phenotype remarkably similar to that observed in platelets from *Par3*<sup>-/-</sup> mice [24], with an approximately 3-fold decrease in sensitivity to thrombin-induced platelet activation. Together, this detailed analysis indicates correct targeting occurred in these mice and resulted in disruption of Par3 expression. However, no replacement PAR1 expression or function was detected.

The reason behind this lack of expression of PAR1 in mouse platelets is unknown. Our initial functional studies showed that hPAR1-KI platelets did not respond to a PAR1-specific agonist but responded normally to other agonists, suggesting that the receptor was either absent or unable to signal. We probed extensively for protein expression but were unable to detect PAR1 in platelets from hPAR1-KI mice via flow cytometry or Western blot. The lack of detection of PAR1 even in permeabilized platelets (by flow cytometry) or in whole cell platelet lysates (by Western blot) strongly suggest that the PAR1 protein is either not translated or is very rapidly degraded. The inability of mouse platelets to ectopically express PAR1 so far appears unique, but remains unexplained and could arise from a number of critical stages in the development of mature PAR1 protein. While possible that the Par3 signal peptide is insufficient to guide PAR1 to the cell surface [28], the highly conserved nature of signal peptides coupled with our observation of robust surface expression of the fusion protein in HEK293T cells suggest otherwise. Other potential explanations include as yet unidentified mechanisms that appear unique to the mouse platelet, such as poor or defective N-linked glycosylation of the receptor leading to receptor instability and signaling dysfunction [29, 30]. Since the primary aim of this study

was to generate a research tool for *in vivo* thrombosis studies, we did not explore the precise mechanism behind the lack of tolerance for PAR1 expression in mouse platelets. Whether or not replacement of the entire human *PAR1* gene in the mouse *Par3* locus will be sufficient to drive expression in mouse platelets remains unknown. Regardless, it appears that other approaches will need to be examined if a non-primate model is to be used for pharmacological screening of drugs targeting platelet PARs.

## Supporting Information

**S1 Fig. The receptor generated in hPAR1-KI mice can be expressed in HEK293T cells.** The fusion protein predicted to be expressed by hPAR1-KI mice, consisting of the mPAR3 signal peptide attached to the mature hPAR1 receptor, can be expressed on the surface of transfected HEK293T cells. HEK293T cells were transfected with native hPAR1 (purple), the mPAR3/hPAR1 fusion (pink), or empty vector (yellow). Note the similar level of detectable expression of native hPAR1 and the mPAR3/hPAR1 fusion. Also shown is a positive control for surface expression (Thy-1) in cells transfected with the empty vector.  
(TIFF)

**S2 Fig. Southern blot analyses of targeted ES cells and founder mice.** (A) ES cell clones were screened for targeting by Southern blot. Targeted clones are indicated by the expected additional band at 8.3kb (e.g. clones D1 and D4). (B) Founder mice screened by the same Southern blot approach as in (A), showing results for 13 mice from two litters. Seven of the 13 mice genotyped as heterozygous targeted, exhibiting both the wt (11.6 kb) and mutant band (8.3kb).  
(TIFF)

**S3 Fig. Sequence annotation for the hPAR1-KI allele.** (A) Predicted genomic DNA sequence for the hPAR1-KI allele, showing *Par3* exon 1 highlighted in grey, the position of primers 3F and 4R in red, the loxP site in yellow, and *PAR1* exon 2 in blue. Red text indicates the sequence output of the PCR product from primers 3F and 4R, which was identical to the predicted genomic DNA sequence. (B) Annotated sequence of the predicted coding sequence and protein chimera, comprising the Par3 signal peptide (grey text) and PAR1 mature receptor (blue text). Red text indicates sequenced PCR product amplified from genomic DNA of hPAR1-KI mice, aligning to the predicted region of PAR1 coding sequence.  
(TIFF)

**S4 Fig. Confirmation of platelet permeabilization.** Platelets isolated from a mouse (left) or human (right) were either left intact (top) or were permeabilized with saponin (0.1%; bottom) prior to incubation with an antibody against the intracellular C-terminal of GPIb $\alpha$  (grey) or isotype control (red). The GPIb $\alpha$  tail antibody produced a rightward shift over isotype only in permeabilized platelets of both species, confirming successful permeabilization. Data shown are representative traces of n = 3 individual experiments.  
(TIFF)

**S1 Table. Whole blood cell counts from wild type and hPAR1-KI mice.**  
(DOCX)

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**Conceptualization:** SLF RAD JRH.

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**Funding acquisition:** JRH.

**Investigation:** SLF ACP MJM RAD JRH.

**Methodology:** SLF ACP MJM RAD JRH.

**Project administration:** JRH.

**Resources:** RAD JRH.

**Supervision:** RAD JRH.

**Validation:** SLF ACP MJM.

**Visualization:** SLF.

**Writing – original draft:** SLF JRH.

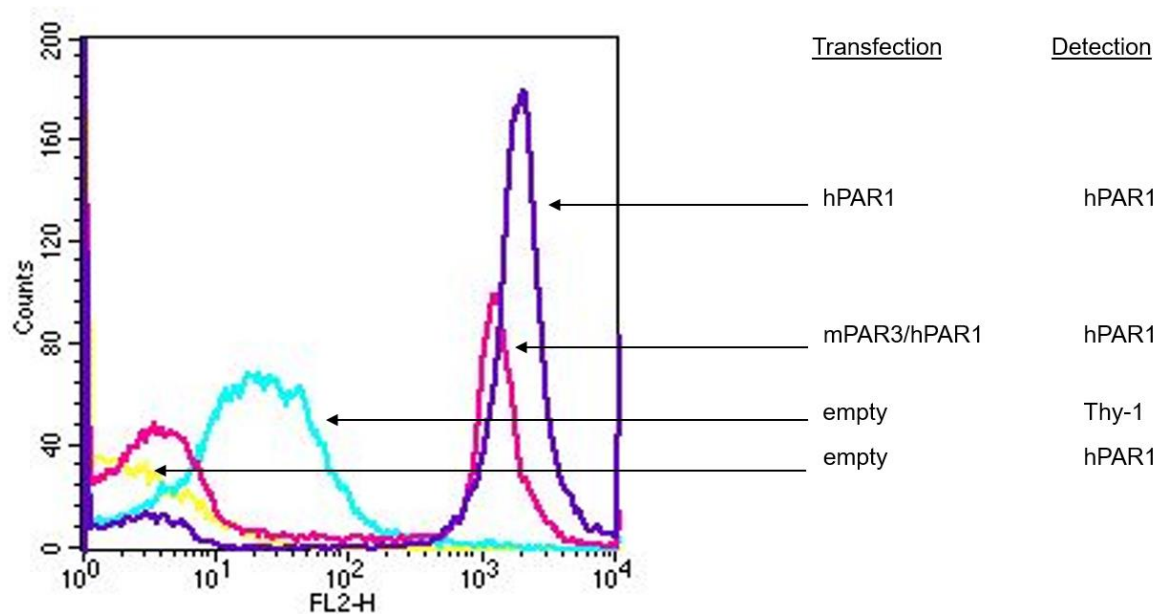
**Writing – review & editing:** SLF ACP MJM RAD JRH.

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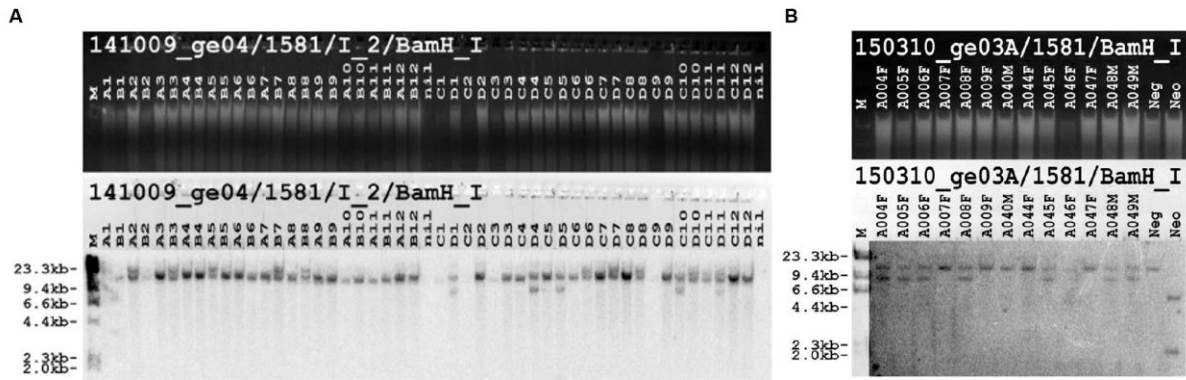
**4.3 Supplementary Figures to French et al. “Humanizing the protease-activated receptor (PAR) expression profile in mouse platelets by knocking *PAR1* into the *Par3* locus reveals PAR1 expression is not tolerated in mouse platelets” (PLoS One, 2016)**



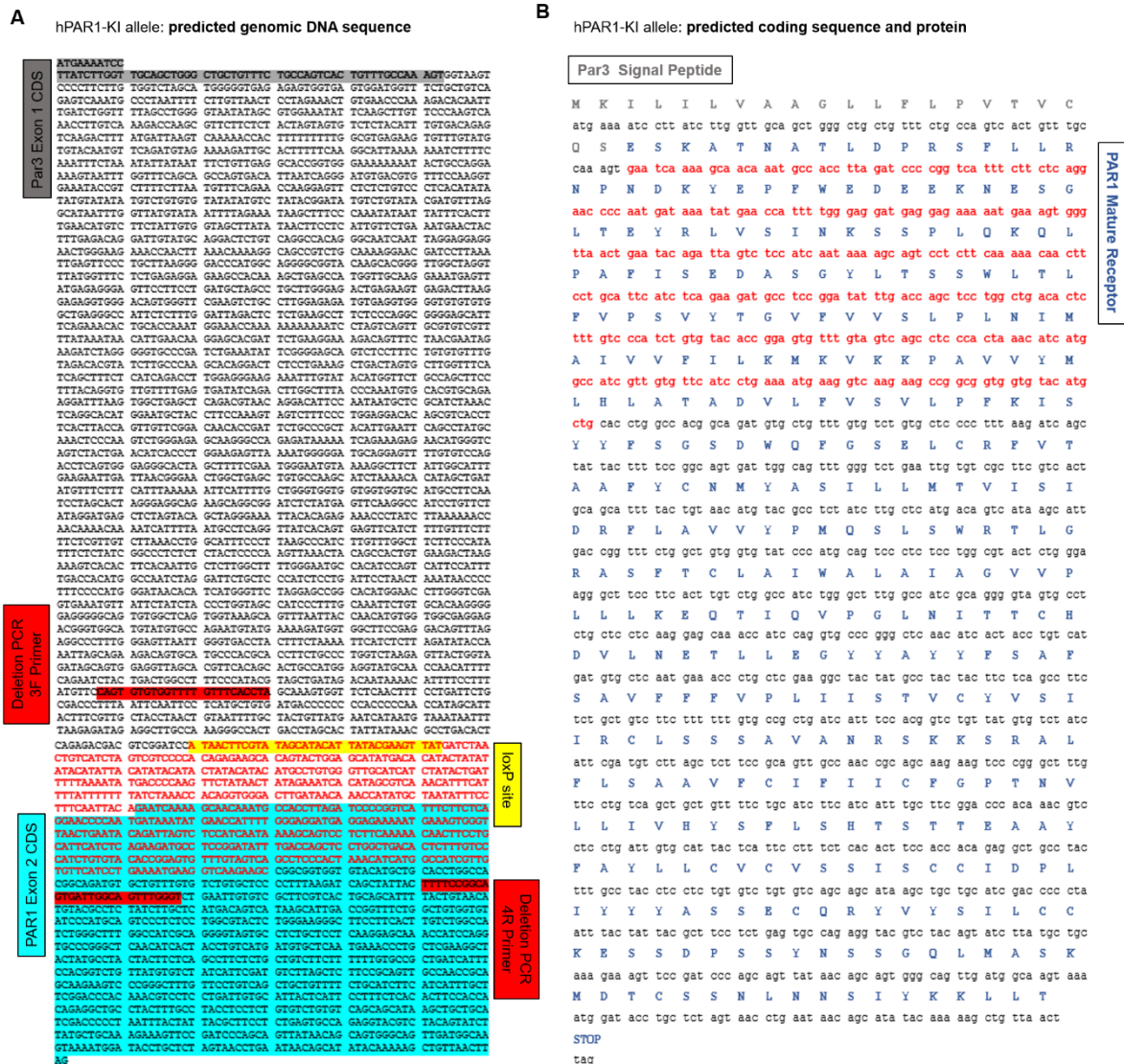
**S1 Fig. The receptor generated in hPAR1-KI mice can be expressed in HEK293T cells.**

The fusion protein predicted to be expressed by hPAR1-KI mice, consisting of the mPAR3 signal peptide attached to the mature hPAR1 receptor, can be expressed on the surface of transfected HEK293T cells. HEK293T cells were transfected with native hPAR1 (purple), the mPAR3/hPAR1 fusion (pink), or empty vector (yellow). Note the similar level of detectable expression of native hPAR1 and the mPAR3/hPAR1 fusion. Also shown is a positive control for surface expression (Thy-1) in cells transfected with the empty vector.



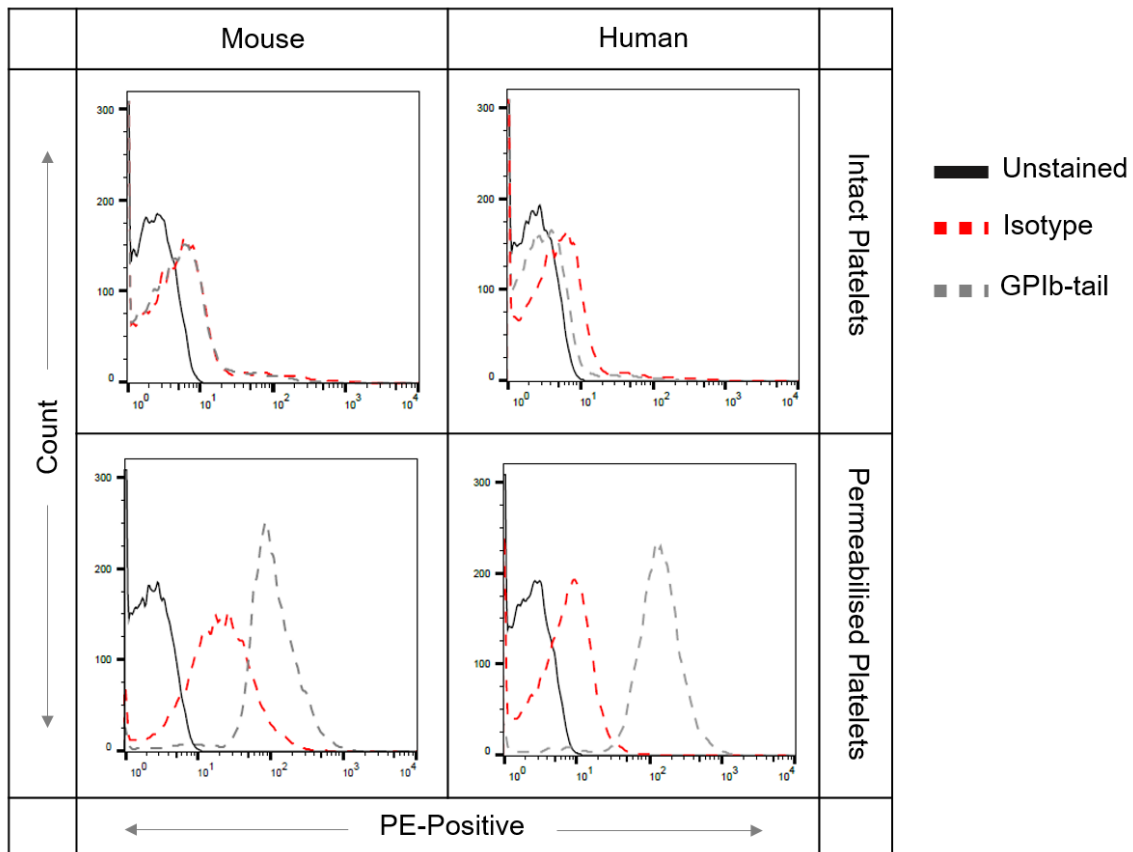


**S2 Fig. Southern blot analyses of targeted ES cells and founder mice. (A)** ES cell clones were screened for targeting by Southern blot. Targeted clones are indicated by the expected additional band at 8.3kb (e.g. clones D1 and D4). **(B)** Founder mice screened by the same Southern blot approach as in (A), showing results for 13 mice from two litters. Seven of the 13 mice genotyped as heterozygous targeted, exhibiting both the wt (11.6 kb) and mutant band (8.3kb).



**S3 Fig. Sequence annotation for the hPAR1-KI allele.** (A) Predicted genomic DNA sequence for the hPAR1-KI allele, showing *Par3* exon 1 highlighted in grey, the position of primers 3F and 4R in red, the loxP site in yellow, and *PAR1* exon 2 in blue. Red text indicates the sequence output of the PCR product from primers 3F and 4R, which was identical to the predicted genomic DNA sequence. (B) Annotated sequence of the predicted coding sequence and protein chimera, comprising the Par3 signal peptide (grey text) and PAR1 mature receptor (blue text). Red text indicates sequenced PCR product amplified from genomic DNA of hPAR1-KI mice, aligning to the predicted region of PAR1 coding sequence.





**S4 Fig. Confirmation of platelet permeabilization.** Platelets isolated from a mouse (left) or human (right) were either left intact (top) or were permeabilized with saponin (0.1%; bottom) prior to incubation with an antibody against the intracellular C-terminal of GPIb (grey) or isotype control (red). The GPIb tail antibody produced a rightward shift over isotype only in permeabilized platelets of both species, confirming successful permeabilization. Data shown are representative traces of n=3 individual experiments.

**S1 Table. Whole blood cell counts from wild type and hPAR1-KI mice.**

	<b>Wild type</b>	<b>hPAR1-KI</b>
<b>WBC (<math>10^3/\mu\text{L}</math>)</b>	7.3 $\pm$ 3.2	7.1 $\pm$ 2.9
<b>RBC (<math>10^6/\mu\text{L}</math>)</b>	9.4 $\pm$ 1.2	9.6 $\pm$ 0.7
<b>PLT (<math>10^3/\mu\text{L}</math>)</b>	887 $\pm$ 50	888 $\pm$ 98

WBC = white blood cells, RBC = red blood cells, PLT = platelets.

## **Chapter 5 - Inhibition of protease-activated receptor 4 (PAR4) with function-blocking antibodies provides equivalent antithrombotic activity in the face of the hyper-reactive Thr120 PAR4 variant**

### **5.1 Introduction**

Despite the lack of an appropriate small animal model to test any novel PAR4 inhibitors *in vivo*, the studies of Chapter 2 coupled with other preclinical and clinical advances with PAR4 antagonists pointed to the overall feasibility of this approach. Specifically, Chapter 2 showed that selective inhibition of PAR4, but not of PAR1, impairs platelet procoagulant function, leading to marked reductions in thrombin generation and fibrin formation during human thrombus formation. This distinct antithrombotic mechanism of action suggests PAR4 inhibition is a viable alternative approach for novel therapy. Toward this goal, a series of small molecule PAR4 inhibitors have been developed, with one of these (BMS-986120) undergoing extensive preclinical testing in non-human primate models and another (BMS-986141) undergoing a phase 2 trial for prevention of transient ischemic attack (NCT02671461). When taken together, there is now substantial rationale for PAR4 antagonism as an antithrombotic approach.

However, as discussed in Chapter 1, a number of PAR4 polymorphisms have recently been described – at least one of which appears to be clinically relevant and may impact on the efficacy of PAR4 antagonists. The recently reported PAR4 variant (at rs773902, encoding a threonine or alanine at amino acid position 120; Thr<sup>120</sup> or Ala<sup>120</sup>), is expressed in 20 – 80% of individuals depending on the population. The Thr<sup>120</sup> variant was shown to render the receptor more sensitive to agonists and less sensitive to antagonists – specifically, near-resistance to the small molecule orthosteric PAR4 antagonist, YD-3. These surprising findings suggest the effectiveness of current strategies to inhibit PAR4 may vary significantly between individuals

and indicate that a different approach to receptor inhibition may be required for indiscriminate PAR4 antagonism across the population.

To address this, the studies of Chapter 5 generated the first human monoclonal inhibitory antibodies against the PAR4 thrombin-cleavage site and examined their activity against the Ala<sup>120</sup> and Thr<sup>120</sup> PAR4 variants. This approach yielded a potent, specific, and highly effective inhibitory antibody that inhibits PAR4 and provides marked antithrombotic effects independently of PAR4 genotype. These findings reveal antibody-mediated inhibition of PAR4 cleavage and activation provides robust antithrombotic activity independently of PAR4 variant expression, and provide rationale for such an approach for improved antithrombotic therapy.

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**Inhibition of protease-activated receptor 4 (PAR4) with function-blocking antibodies provides equivalent antithrombotic activity in the face of the hyper-reactive Thr120 PAR4 variant**

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## Abstract

Thrombin activates human platelets via two protease-activated receptors (PARs), PAR1 and PAR4, and both are antithrombotic drug targets – a PAR1 inhibitor is approved for clinical use and a PAR4 inhibitor is in trial. However, a sequence variant in human PAR4 (rs773902, encoding Thr120 in place of Ala120), expressed in 20 – 80% of individuals depending on the population, renders the receptor more sensitive to agonists and less sensitive to antagonists. Here, we develop the first human monoclonal function-blocking antibody to human PAR4 (hPAR4) and show it provides equivalent efficacy against the Ala120 and Thr120 PAR4 variants. This candidate was generated from a panel of anti-hPAR4 antibodies and was screened for specific inhibitory activity against the Ala120 and Thr120 PAR4 variants in cultured cells, isolated platelets, and an *ex vivo* whole blood thrombosis assay. The antibody binds hPAR4 with high selectivity (16-fold over PAR1, PAR2 and PAR3) and affinity ( $K_D \approx 0.4$  nM) and is capable of near-complete inhibition of thrombin cleavage of either the Ala120 or Thr120 PAR4 variant transfected into HEK293T cells. Platelets from individuals expressing the Thr120 PAR4 variant exhibit increased thrombin-induced aggregation and phosphatidylserine exposure versus those with the Ala120 PAR4 variant, yet the hPAR4 antibody inhibited these responses equivalently ( $IC_{50} \approx 5$   $\mu$ g/ml). In support of these *in vitro* findings, the antibody significantly impairs platelet procoagulant activity in an *ex vivo* thrombosis assay, with equivalent inhibition of fibrin formation and overall thrombus size in blood from individuals expressing the Ala120 or Thr120 PAR4 variant. These findings reveal antibody-mediated inhibition of PAR4 cleavage and activation provides robust antithrombotic activity independently of PAR4 variant and provide rationale such an approach for antithrombotic therapy.

## Introduction

Protease-activated receptors (PARs) are G protein-coupled receptors that are present on the surface of a range of cells and respond to a variety of proteases (for review, see (1)). Human platelets express two PARs, PAR1 and PAR4, and these receptors are primarily responsible for mediating the platelet-activating effects of the key coagulation protease, thrombin (2). Due to this central function in platelet biology, both platelet PARs have been the focus of antithrombotic drug development. The initial strategy was to inhibit PAR1 as it has an approximate 30-fold greater affinity for thrombin than PAR4. This approach led to the recent approval of the first PAR1 inhibitor, vorapaxar, for the prevention of myocardial infarction and peripheral arterial disease (3,4). However, the clinical utility of vorapaxar is limited by increased rates of major bleeding in patients on the drug (5). As a result, there is now emerging interest in targeting PAR4 as a safer antithrombotic approach.

There is substantial rationale for developing PAR4 inhibitors as antithrombotics (for review, see (6,7)). One key point of distinction between PAR1 and PAR4 is the different signaling kinetics of the two receptors and the impact this has on the regulation of platelet function. Specifically, PAR4 contains an anionic sequence downstream of the thrombin cleavage site which serves to prolong the thrombin-receptor interaction (8). One effect of the lower affinity but more prolonged interaction between thrombin and PAR4 versus PAR1 is that activation of PAR4 induces a more sustained, albeit weaker, intracellular signal than the robust and acute signal elicited downstream of PAR1 (9). This has been most obviously observed with the kinetics of PAR-induced calcium signaling. In the setting of platelet function, prolonged calcium signaling drives the procoagulant response. Indeed, selective inhibition of PAR4, but not of PAR1, specifically impairs platelet procoagulant function, leading to marked reductions in thrombin generation and fibrin formation during human thrombus formation (10). This distinct antithrombotic mechanism of action suggests PAR4 inhibition is a viable alternative

approach for novel therapy. Toward this goal, a series of small molecule PAR4 inhibitors have been developed, with one of these (BMS-986141) undergoing a phase 2 trial for prevention of transient ischemic attack (NCT02671461). Proof-of-concept studies were performed using the related compound, BMS-986120, which afforded impressive antithrombotic activity in cynomolgous monkeys with a safety profile that exceeded that of the P2Y<sub>12</sub> antagonist, clopidogrel (11). Together, these studies provide strong rationale for pursuing PAR4 antagonists as novel antithrombotics.

However, a number of PAR4 polymorphisms have recently been described – at least one of which appears to be clinically relevant and may impact on the efficacy of PAR4 antagonists. The recently reported sequence variant in human PAR4 (rs773902, encoding either a threonine or alanine at amino acid position 120; Thr120 or Ala120), is expressed in 20 – 80% of individuals depending on the population, and renders the receptor more sensitive to agonists and less sensitive to antagonists (12,13). Specifically, platelets from individuals expressing the Thr120 variant of PAR4 exhibit increased responsiveness to a PAR4-selective activating peptide and resistance to the small molecule orthosteric PAR4 antagonist, YD-3, when compared with individuals expressing the Ala120 variant (13). In both cases, individuals heterozygous for the SNP displayed an intermediate phenotype. These surprising findings suggest the effectiveness of current strategies to inhibit PAR4 may vary significantly between individuals and indicate that a different approach to receptor inhibition may be required for indiscriminate PAR4 antagonism across the population.

To address this, we utilized VelocImmune® HumAb mice to generate the first human monoclonal inhibitory antibodies against PAR4. We targeted the thrombin cleavage site of the receptor and examined the activity of antibodies against the Ala120 and Thr120 PAR4 variants in cultured cells, isolated platelets, and an *ex vivo* whole blood thrombosis assay. Although there have been a number of previous reports of function-blocking PAR4 antibodies (14-16),



including our own (10), effective inhibition of thrombin-induced receptor activation has been limited. However, our approach yielded a potent, specific, and highly effective inhibitory antibody that inhibits PAR4 and provides marked antithrombotic effects independently of hPAR4 genotype. These findings reveal antibody-mediated inhibition of hPAR4 cleavage and activation provides robust antithrombotic activity independently of hPAR4 variant expression, and provide rationale for such an approach for improved antithrombotic therapy.

## Materials and Methods

### *Materials*

The following reagents were stored according to manufacturer's instructions: Alexa Fluor 488-conjugated goat anti-mouse IgG (BD Biosciences, San Jose, CA, USA), mouse IgG<sub>1</sub> isotype (Santa Cruz Biotechnology, Dallas, TX, USA), anti-CD41a (BD Biosciences), FITC-conjugated anti-FLAG antibody (clone M2, Sigma-Aldrich, St Louis, MO, USA), DyLight650-conjugated anti-fibrin antibody (clone 59D8 (17) generous gift from Dr Vivien Chen, University of New South Wales, Australia), anti-CD9-PE (BD Biosciences), Alexa Fluor 488-conjugated annexin-V (Sigma-Aldrich), hPAR4 immunising peptide (GGDDSTPSILPAPRGYPGQVC-KLH), hPAR4 naked peptide (GGDDSTPSILPAPRGYPGQVC), hPAR4 biotinylated peptide (GDDSTPSILPAPRGYPGQVC-GGGGSKB), hPAR3 biotinylated peptide (AKPTLPIKTFRGAPPNSF-GGGGSKB), hPAR2 biotinylated peptide (SCSGTIQGTNRSSKGRSL-GGGGSKB), hPAR1 biotinylated peptide (SKATNATLDPRS FLLRNP-GGGGSKB) (all from Auspep, Melbourne, Australia), PAR4-activating peptide (PAR4-AP; AYPGKF) and PAR1-activating peptide (PAR1-AP; TFLLR) were synthesized by Prof Philip Thompson (Monash University, Australia), vorapaxar (Axon Medchem, Reston VA, USA), human  $\alpha$ -thrombin (Sigma-Aldrich), calcium ionophore A23187 (Sigma-Aldrich), bovine type 1 collagen (Sigma-Aldrich), and hirudin (lepirudin, Celgene, Summit, NJ, USA). For thrombin measurements in whole blood, a fluorescence resonance energy transfer (FRET)-based thrombin activity sensor (Thr-SP, CPC Scientific, Sunnyvale, CA, USA), was linked to the anti-CD41a antibody via CLICK chemistry as previously described (18).

### *Antibody production and purification*

Immunizations were undertaken in VelocImmune® HumAb mice (generously provided by Regeneron Pharmaceuticals), which have been genetically modified by replacing the variable

regions of the mouse heavy and light chain Ig loci with the corresponding human sequence to produce antibodies that have human variable regions with the mouse constant regions (19,20). All mouse studies were approved by the Monash University Animal Ethics Committee. 14 week old HumAb mice were injected subcutaneously with a KLH-coupled peptide corresponding to a region spanning the thrombin cleavage site of PAR4 (GDDSTPSILPAPR/GYPGQVC-KLH, where / indicates the thrombin cleavage site; 25 µg in Freund's complete adjuvant) with 3 boosts given every 2 weeks for 6 weeks. After the final boost, primary splenocytes were isolated and fused with myeloma Sp2/0 cells, and plated onto 96 well plates to generate antibody-producing hybridomas. Supernatants from the resulting hybridomas were screened for high affinity specific hPAR4 antigen-positive lines using microarray (Arraviet Super Marathon, ArrayJet, UK) and standard ELISA. Cross reactivity to hPAR1, hPAR2 and hPAR3 was also measured. Specific hPAR4 clones were subcloned to monoclonality by limiting dilution. Specific hybridoma lines were expanded, adapted to serum free and suspension culture, and purified anti-hPAR4 antibody was isolated by standard affinity chromatography procedures using Protein A/G sepharose.

#### *Surface plasmon resonance binding studies*

The binding kinetics of purified antibody were determined via surface plasmon resonance (SPR; Bio-Rad ProteOn XPR36). Biotinylated hPAR1, hPAR2, hPAR3 or hPAR4 peptides (25 µg/mL) were captured on the ProteOn NLC biosensor chip consisting of NeutrAvidin surface bound to the alginate polymer. Varying antibody concentrations (6.25 – 100 nM) were flowed across the channels. The data were analyzed using Proteon Manager software with a Langmuir interaction model that assumes a 1:1 binding interaction between the peptide and antibody.

### *Human blood samples*

All human studies were approved by the Monash University Human Research Ethics Committee. Blood was collected after informed consent from healthy adults (21 – 50 years old, of both sexes) who had not taken anti-platelet medications in the past 10 d. Blood was drawn from the antecubital vein using a 19 gauge butterfly needle into syringes containing either one-seventh acid citrate dextrose (ACD; 7:1 v/v, final concentration) for platelet isolation or one-tenth volume trisodium citrate (0.32% w/v, final concentration) for whole blood flow experiments and DNA extraction.

### *PAR4 genotyping*

Genomic DNA was extracted from the buffy coat of blood samples collected in citrate, using the DNEasy Blood and Tissue Kit (Qiagen, Hilden, Germany) as per manufacturer's instructions. Samples were genotyped at rs773902 using the TaqMan SNP Genotyping Assay (Assay C-7493801-10, ThermoFisher Scientific).

### *Flow cytometry*

Antibody binding to native PAR4 on the platelet surface was determined using flow cytometry. Platelets were isolated from human whole blood collected in ACD as previously described (10). Isolated platelets ( $1 \times 10^6$  cells) were then incubated with anti-PAR4 antibody (10  $\mu\text{g/mL}$ ), anti-human CD41a (10  $\mu\text{g/mL}$ ), or isotype control (mouse IgG<sub>1</sub>, 10  $\mu\text{g/mL}$ ) for 30 min at 37°C. Cells were then fixed with paraformaldehyde (2% final concentration) for 30 min at room temperature. The suspension was centrifuged at 1000 x g for 2 min to obtain the platelet pellet, which was then resuspended in modified Tyrode's buffer (12 mM NaHCO<sub>3</sub>, 10 mM HEPES pH 7.4, 137 mM NaCl, 2.7 mM KCl, 5.5 mM D-glucose, 1.8 mM CaCl<sub>2</sub>, 5% bovine serum albumin) containing 10  $\mu\text{g/mL}$  Alexa488-conjugated anti-mouse IgG. After 30 min at 37°C the samples were centrifuged again and the platelet pellet resuspended in modified

Tyrode's buffer and analyzed using a flow cytometer (FACSCalibur; BD Biosciences) and FlowJo software (v10).

#### *PAR4 cleavage assay*

FLAG-tagged Ala120-PAR4 or Thr120-PAR4 expression vectors (pBJ-FLAG-PAR4-120A-296F and pBJ-FLAG-PAR4-120T-296F, a generous gift from Prof. Paul Bray (University of Utah, Salt Lake City, Utah)) were transiently transfected into HEK293T cells using Lipofectamine 2000 (ThermoFisher Scientific), as per the manufacturer's instructions. Cells were harvested 48 h after transfection, washed twice with PBS, and resuspended to  $1 \times 10^6$  cells/mL. Cells were then pre-treated with either anti-PAR4 antibody (1 – 100  $\mu\text{g/mL}$ ), or matched isotype control (mouse IgG<sub>1</sub>; 100  $\mu\text{g/mL}$ ) for 15 min at 37°C, before being stimulated with thrombin (2 U/mL) for 10 min. The reaction was stopped with hirudin (8 U/mL). Samples were then incubated with FITC-conjugated anti-FLAG antibody (10  $\mu\text{g/mL}$ ) for 1 h at room temperature and analyzed by flow cytometry (FACSCalibur) and FlowJo software (v10).

#### *Platelet aggregation*

Platelet aggregation was measured by light transmission aggregometry in a 96-well plate format as previously described (21). Human isolated platelets ( $2 \times 10^8/\text{mL}$ ) were stimulated with varying concentrations of thrombin (0.0178 – 1 U/mL), PAR4-AP (10 – 100  $\mu\text{M}$ ), or PAR1-AP (3 – 30  $\mu\text{M}$ ). The plate was analysed at 37°C in a FLUOstar OPTIMA plate reader (BMG Labtech, Ortenberg, Germany) using a 595 nm excitation filter, for a period of 50 min (10 read cycles with 5 min double orbital shake period between each read). Aggregation was calculated as  $\frac{OD(\text{No Agonist}) - OD(\text{Agonist})}{OD(\text{No Agonist}) - OD(\text{blank})} \times 100$  at the time point where aggregation was at a maximum. Optical density was normalised against the blank (maximum) and unstimulated platelets (minimum) and expressed as % maximum. For antagonist studies, platelets were pre-

treated for 15 min at 37°C with anti-PAR4 antibody (1 – 100 µg/mL), vorapaxar (90 nM), a combination of both, or their vehicle controls (mouse IgG<sub>1</sub> and 1% DMSO, respectively).

#### *Phosphatidylserine exposure*

Phosphatidylserine (PS) exposure on the platelet surface was used as a measure of platelet procoagulant activity. Here, isolated platelets ( $1 \times 10^6$  cells) were stimulated with increasing concentrations of either PAR4-AP (0.1 – 3 mM), thrombin (0.1 – 3 U/mL), or calcium ionophore (20 µM) for 30 min at 37°C. Alexa Fluor 488-conjugated annexin V (1:100 dilution) was added after agonist stimulation for PS detection. After the 30 min stimulation, samples were fixed with paraformaldehyde (2% final concentration) and resuspended in a modified Tyrode's buffer for flow cytometry analysis, as above (FACSCalibur).

#### *Whole blood thrombosis assay*

Human whole blood collected in citrate (3.2%) was pre-incubated for 15 min at 37°C with PE-conjugated anti-CD9 antibody (4 µg/mL), Thr-SP (5 µM), and a Dylight650-conjugated anti-fibrin antibody (5 µg/mL), plus either the anti-PAR4 antibody (0.1 mg/mL) or hirudin (800 U/mL). Whole blood was re-calcified with 10 mM CaCl<sub>2</sub> (final concentration) to initiate coagulation, and drawn over glass microslides (1 x 0.1 mm internal diameter; Vitrotubes, Vitrocom, NJ, USA) coated with bovine type 1 collagen (250 µg/ml) using a Harvard pump (Instech Laboratories, PA, USA) at a fixed flow rate of 0.06 mL/min, resulting in a wall shear rate of 600 s<sup>-1</sup>. Tri-colour confocal fluorescence images were recorded at 488, 561 and 647 nm excitation, collected through a 25x water immersion objective. Thrombi were formed over 3 min of blood flow after which calcium-free Tyrode's buffer was flowed over the thrombi. Confocal z-stacks encompassing the entire height of the thrombus field were continuously recorded (Z step 2 µm; Nikon A1r, with NIS software) over a period of 10 min. Offline analysis of thrombi parameters was performed using NIS software. Image series were initially

thresholded empirically, and then the same threshold applied to all subsequent experiments, since identical experimental and confocal settings were used throughout. Platelet thrombi were defined using anti-CD9-PE, and thrombin activity and fibrin volume were quantified using average fluorescence of the thrombus field. Data were normalised against the hirudin baseline, and expressed as a percentage of the control.

#### *Statistical analyses*

Statistical analyses were performed using GraphPad Prism software (v6.0). Significance was defined at  $p < 0.05$  as determined by either unpaired, two-tailed, Student's *t*-test or one-way ANOVA with Dunnet's post-hoc test for multiple comparisons, as indicated in the relevant figure legends.

## Results

### *Development of a human monoclonal function-blocking antibody against PAR4*

VelocImmune® HumAb mice were immunized with a KLH-linked peptide corresponding to a region of hPAR4 spanning the thrombin cleavage and activation site (GDDSTPSILPAPRGYPGQVC-KLH) (Figure 1A). Serum titres from routine bleeds were monitored and spleens from the highest producing mice were harvested. Splenocytes were then fused with Sp2/0 cells and resulting hybridomas (~ 5000) were screened for binding to the naked version of the immunizing peptide by microarray and confirmed secondarily by ELISA. ELISA-based antigen screening showed that clone mAb-5RC3 (mAb-RC3) bound naked hPAR4 peptide with a 16-fold selectivity over any of hPAR1, hPAR2, or hPAR3 peptides, and did not bind to mouse PAR4 (Figure 1B). SPR confirmed binding of clone mAb-RC3 to the target antigen and estimated it bound the immunizing peptide with a  $K_D$  of ~ 0.4 nM (Figure 1C). We also confirmed mAb-RC3 bound native PAR4 on human platelets. Here, mAb-RC3 (10 µg/mL) bound to a purified human platelet preparation, with a rightward shift in fluorescence intensity over that produced by a mouse IgG<sub>1</sub> isotype (Figure 1D).

### *mAb-RC3 inhibits thrombin cleavage of both Ala120 and Thr120 PAR4 variants*

We next examined the function-blocking activity of the clone mAb-RC3 in a bioassay of thrombin-induced cleavage of PAR4. In HEK293Ts transfected with plasmids containing either Ala120-PAR4 or Thr120-PAR4 with an N-terminal FLAG-tag, thrombin caused a similar concentration-dependent loss of FLAG epitope, with a maximum cleavage of receptor of approximately 60% in both cases (Figure 2A). Under conditions of maximum PAR4 cleavage, mAb-RC3 inhibited receptor cleavage in a concentration-dependent manner (Figure 2B). This effect was not different between the Ala120 and Thr120 PAR4 variants (Figure 2B). Pre-treatment with the highest concentration of mAb-RC3 examined (100 µg/mL) almost



prevented thrombin cleavage of either PAR4 variant, with only ~ 5% residual loss of the FLAG epitope detected (Figure 2B).

*mAb-RC3 inhibits the enhanced platelet aggregation elicited by the Thr120-PAR4 variant*

To date, the heightened PAR4 response in platelets from individuals expressing the Thr120 PAR4 variant has only been demonstrated in response to the PAR4-specific agonist peptide, PAR4-AP, and only in a North American population (13). Therefore, we first confirmed this phenotype in our local (Australian) cohort. In line with previous observations (13), we observed a significant increase in platelet aggregation induced by PAR4-AP in individuals expressing the Thr120 variant versus those expressing the Ala120 variant (Figure 3A). Platelets from heterozygous individuals displayed a similar increase in responsiveness to PAR4-AP (Figure 3A). We next determined whether this hyper-reactivity in platelets from individuals with the Thr120 allele also occurred in response to the physiological PAR4 agonist, thrombin, since this has not been previously reported. Although thrombin-induced aggregation was increased in platelets from Thr120 individuals (homozygous or heterozygous) versus those from Ala120 individuals, this effect was less obvious than that observed in response to PAR4-AP (Figure 3B). To isolate the PAR4-dependant effects of thrombin in this assay, platelets were treated with the PAR1 antagonist, vorapaxar, at a concentration (90 nM) we have previously shown sufficiently blocks PAR1 activation under these conditions (21). Here, we observed a further dampening of the PAR4 genotype-dependent difference in platelet aggregation (Figure 3C). Regardless, mAb-RC3 caused a concentration-dependent inhibition of this thrombin-induced, PAR4-dependent, platelet aggregation. mAb-RC3 almost abolished aggregation at the highest concentrations examined (30 or 100 µg/mL), with indistinguishable effects across donors expressing the various PAR4 variants (Figure 3D) and with an estimated IC<sub>50</sub> of 5.8 µg/mL (Figure 3E).

*mAb-RC3 inhibits the enhanced platelet procoagulant response elicited by the Thr120-PAR4 variant*

The generation of procoagulant platelets is important for thrombus growth and stability (22,23), and thrombin-induced procoagulant platelet production is mediated predominantly by PAR4 (10). Therefore, we next examined the sensitivity of thrombin-induced procoagulant platelet production (assessed by annexin V binding to surface-exposed PS) in donors expressing the Ala120 versus Thr120 PAR4 variant, and the ability of mAb-RC3 to inhibit this response. We observed a strikingly similar trend to the aggregation studies in Figure 3. First, PAR4-AP induced a greater increase in PS on platelets expressing the Thr120 PAR4 variant (heterozygous or homozygous) than on those expressing the Ala120 PAR4 variant (Figure 4A). Second, the same, less obvious, trend was observed in response to thrombin (Figure 4B). Third, mAb-RC3 concentration-dependently inhibited thrombin-induced PS exposure independently of PAR4 genotype, again nearly abolishing the response at the highest concentrations examined (100 µg/mL) in platelets expressing either of the PAR4 variants (Figure 4C).

*mAb-RC3 inhibits human thrombus formation independently of PAR4 variant expression*

Finally, the antithrombotic activity of mAb-RC3 was assessed in a human whole blood thrombosis assay in which coagulation remains intact (10). Specifically, we measured platelet deposition, thrombin generation, and fibrin deposition in real time in blood from individuals expressing the Ala120 or Thr120 PAR4 variant (Figure 5A). Perhaps surprisingly, there was no significant difference in the extent of any of platelet deposition, thrombin generation, or fibrin deposition between donors expressing the various PAR4 variants (Figure 5B,C). Importantly, and in support of the procoagulant platelet response experiments of Figure 4, pretreatment of blood with mAb-RC3 alone was sufficient to significantly reduce thrombin generation (~ 40%) and fibrin formation (~ 60%) in formed thrombi (Figure 5D,E). As in the

*in vitro* experiments, this reduction in thrombin and fibrin occurred independently of PAR4 genotype as it was evident in blood taken from individuals expressing the Ala120 or Thr120 PAR4 variant and in individuals who genotyped as heterozygous at the allele (Figure 5B,C). As predicted from the *in vitro* experiments in Figure 4, the effect of mAb-RC3 on fibrin formation (~ 60% reduction; Figure 6C) was more pronounced than that on platelet deposition (~ 30% reduction; Figure 6B). Together, the overall impact of mAb-RC3 pretreatment was a marked reduction in total thrombus volume (> 50%; Figure 6D). Again, the antithrombotic effect of mAb-RC3 was independent of PAR4 genotype, occurring in 12 of 14 individuals examined (Figure 6E). Indeed, mAb-RC3 only failed to provide an antithrombotic effect in the two individuals with the lowest baseline thrombotic response (Figure 6E). These studies indicate that PAR4 inhibition with the function-blocking monoclonal antibody developed here affords equally effective anti-platelet and antithrombotic effects in individuals regardless of their expression of the sequence variant of PAR4 at rs773902.

## Discussion

We have generated the first human monoclonal function-blocking antibody against the platelet thrombin receptor, PAR4. We utilized VelocImmune® HumAb mice to generate human monoclonal antibodies directed against the thrombin cleavage site of PAR4 – a strategy similar to one we have used previously to generate a specific antagonist of thrombin-induced PAR4 activation (10). One clone (mAb-RC3) demonstrated specific and high-affinity binding to PAR4, and effective inhibition of thrombin-induced PAR4 cleavage and activation that markedly improves on previous antibody-based approaches (14-16). Importantly, mAb-RC3 inhibits the Ala120 PAR4 variant and the hyper-reactive Thr120 PAR4 variant equally well in isolated platelets and provides equivalent antithrombotic effects in blood from individuals regardless of their PAR4 genotype. These studies thereby provide a new agent for PAR4 inhibition that may have broad clinical utility.

There are now a number of PAR4 inhibitors (11,24,25), with one small molecule (BMS-986141) being evaluated in clinical trials (NCT02341638, NCT02671461). However, specificity and efficacy have proven especially challenging. Binding analyses performed here indicate mAb-RC3 binds PAR4 with high affinity ( $K_D \sim 0.5$  nM) and with almost no cross reactivity to other PARs. Furthermore, mAb-RC3 appears significantly more effective than previous antibodies (10,14-16). In addition to the usual challenges of specificity and efficacy, the recent discovery of a clinically significant PAR4 variant (rs773902) has raised additional issues for PAR4 antagonism, with the Thr120-PAR4 variant exhibiting increased sensitivity to agonists and an insensitivity to at least one small molecule antagonist (13). The mAb-RC3 antibody developed here appears to completely overcome this, suggesting antibody-based inhibition of PAR4 may be required for effective and well-controlled antithrombotic activity across the population.

The Thr120-PAR4 variant has previously been associated with up-regulated platelet responses to a PAR4 activating peptide (13). Our findings extend on this to indicate the heightened response also occurs in response to PAR4's physiological agonist, thrombin, with platelets from individuals expressing the Thr120 PAR4 variant exhibiting increased thrombin-induced aggregation. The mechanism by which the amino acid change at position 120 (located in the second transmembrane domain) impacts PAR4 function remains unknown. Our data suggest it is not due to changes in receptor cleavage since the cleavage-site spanning mAb-RC3 was equally effective at inhibiting PAR4 cleavage and activation in Thr120 and Ala120 PAR4 variants. Whether or not interactions at the ligand binding domain within the second extracellular loop are involved remain untested, but are consistent with the findings from us and others (13) showing differences in responses to PAR4 activating peptides and small molecule antagonists that act at this site.

We have previously shown that PAR4 activation via thrombin drives procoagulant platelet activity (10). Here, we extended on these studies to provide the first evidence for a heightened production of procoagulant platelets in individuals expressing the Thr120-PAR4 variant. That this response was also inhibited by mAb-RC3 provides further rationale for its use as an antithrombotic approach. Although a similar trend of increased platelet procoagulant activity in response to PAR4-AP was observed with thrombin, consistent with our experiments examining platelet aggregation, the extent of any increase with thrombin was far less apparent than with PAR4-AP. This observation may explain the lack of correlation between PAR4 genotype and *ex vivo* thrombus formation observed here. We observed no differences in any of platelet deposition, thrombin generation, fibrin formation, or total thrombus volume between donors of different PAR4 genotypes. Whether the increased platelet sensitivity to thrombin is truly insufficient to cause effects in the more involved whole blood system will require more thorough examination – for example in a greater number of individuals and across a range of

thrombogenic conditions. However, regardless of the impact of the sequence variant of PAR4 at rs773902 on thrombus formation, mAb-RC3 alone was sufficient to provide a marked antithrombotic effect in almost all donors examined. As predicted by our in vitro studies, this effect was largely driven by a reduction in platelet procoagulant activity and subsequent fibrin formation, although some contribution by impaired platelet deposition was also apparent. The marked antithrombotic effect provided by mAb-RC3 in blood from donors of all PAR4 genotypes suggests directly targeting the thrombin cleavage site may be required for effective and consistent PAR4 inhibition across the population. Given the efficacy and relative safety of PAR4 inhibition in animal models (11), determining the antithrombotic effects of mAb-RC3 in human blood in combination with other anti-platelet agents that target distinct aspects of thrombus formation (e.g. aspirin, P2Y<sub>12</sub> receptor antagonists) will be of interest.

## **Conclusion**

We have developed a human monoclonal PAR4 antibody that is a highly selective and effective inhibitor of PAR4 cleavage and activation. This function-blocking PAR4 antibody impairs thrombin-induced PAR4 cleavage, platelet activation, and pro-thrombotic activity equivalently in the face of expression of either the Ala120 or Thr120 PAR4 variant. These findings reveal a novel approach to PAR4 inhibition that overcomes the impaired sensitivity of existing approaches against the commonly expressed Thr120-PAR4 variant, and provide rationale for such an approach for improved antithrombotic therapy.

## **Acknowledgements**

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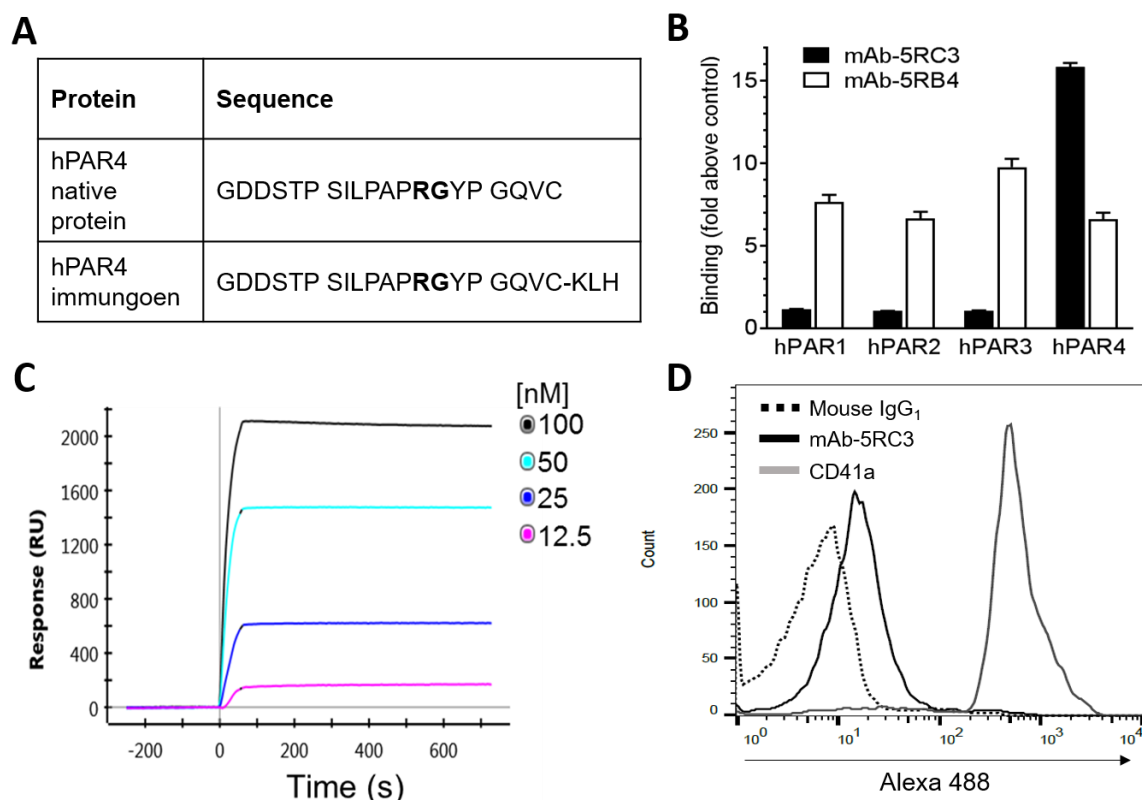
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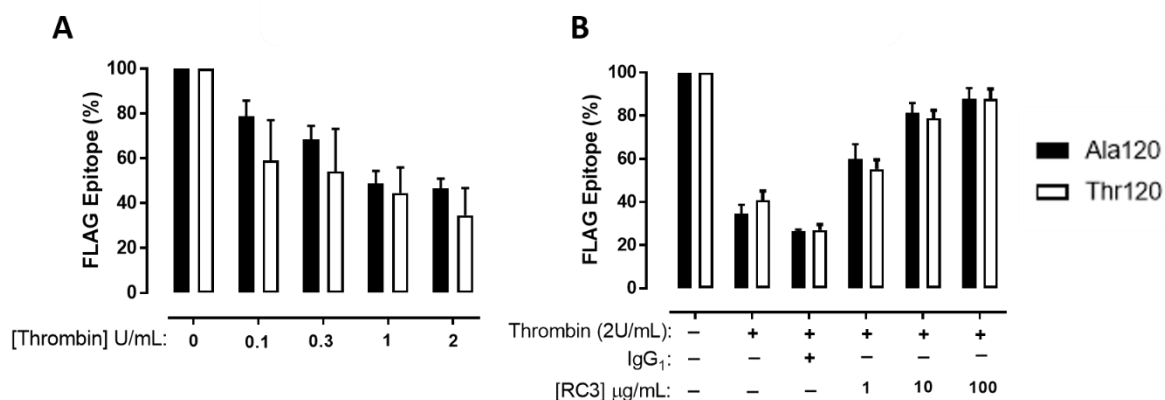
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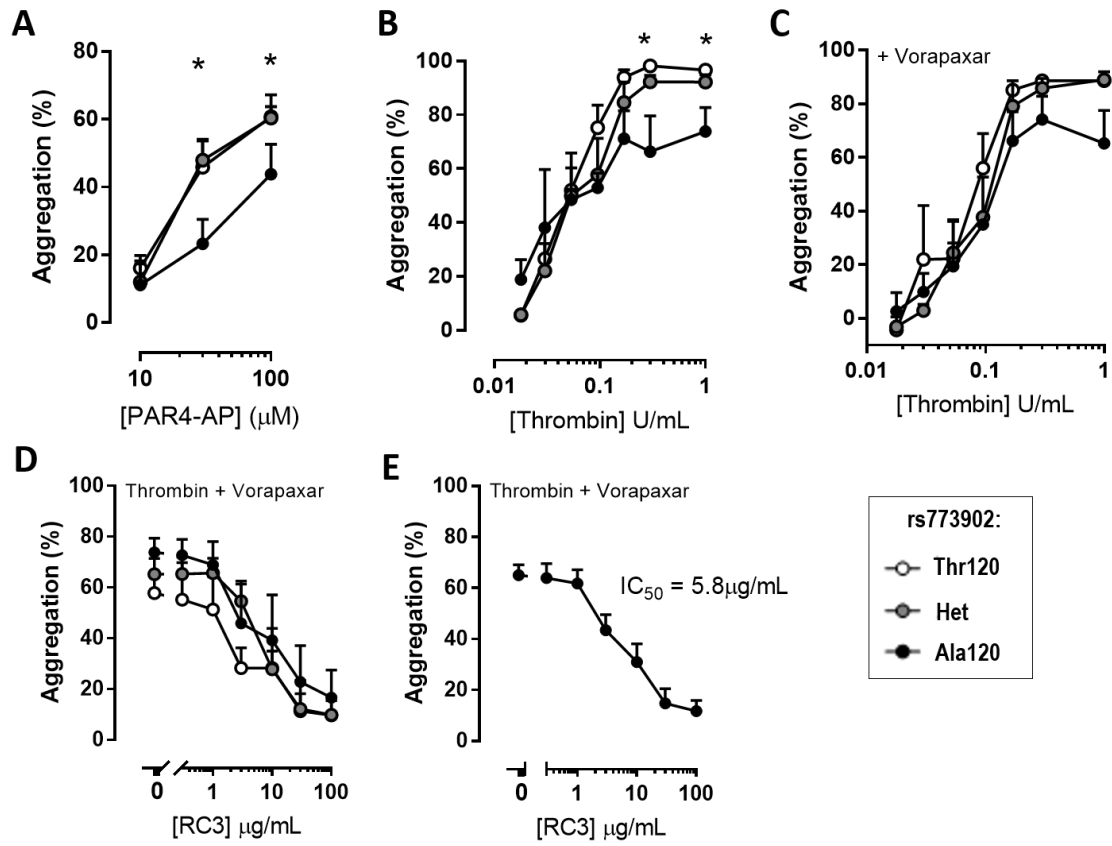
## Figures



**Figure 1. Development of a human monoclonal function-blocking antibody against human PAR4 (mAb-RC3).** (A) HumAb mice were immunized with a KLH-coupled peptide corresponding to a region in the N-terminus of hPAR4 spanning the thrombin cleavage site (GDDSTPSILPAP**RG**YPGQVC-KLH, bold font indicated thrombin cleavage site). (B) Initial ELISA-based antigen screening showed mAb-5RC3 (mAb-RC3) bound native human PAR4 peptide (hPAR4) with a 16-fold selectivity over human PAR1, 2, or 3 peptides, while mAb-5RB4 bound all four peptides similarly. (C) Binding was confirmed by SPR analysis which showed that mAb-RC3 bound native PAR4 peptide with a  $K_D \sim 0.4$  nM. (D) mAb-RC3 (10 $\mu$ g/mL) binding to human PAR4 on the platelet surface was confirmed by flow cytometry. Note a rightward shift in fluorescence intensity compared to the matched Isotype control (mouse IgG<sub>1</sub>, dashed line). The platelet-specific anti-CD41a antibody (grey line) was used as a positive control. (C-D) Data are representative images of n=3 independent experiments.

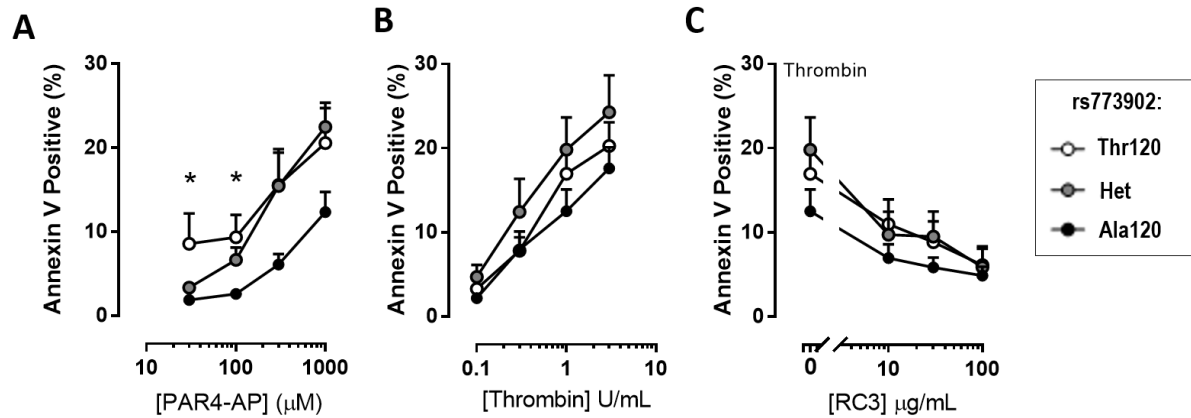


**Figure 2. mAb-RC3 inhibits thrombin cleavage of both Ala120 and Thr120 PAR4 variants.** Thrombin cleavage of PAR4 was assessed in HEK293T cells transiently transfected with either Ala120-PAR4 or Thr120-PAR4 variants with an N-terminal FLAG tag. **(A)** Cells were stimulated with increasing concentrations of thrombin (0.1 – 2 U/mL) and receptor cleavage measured as a loss of FLAG-epitope using a FITC-conjugated anti-FLAG antibody by flow cytometry. **(B)** Pre-incubation of transfected cells with mAb-RC3 (1-100  $\mu$ g/mL) before thrombin stimulation (2 U/mL) provided equivalent and near-complete inhibition of thrombin cleavage of either PAR4 variant. Data is mean  $\pm$  SEM of n=3 experiments.

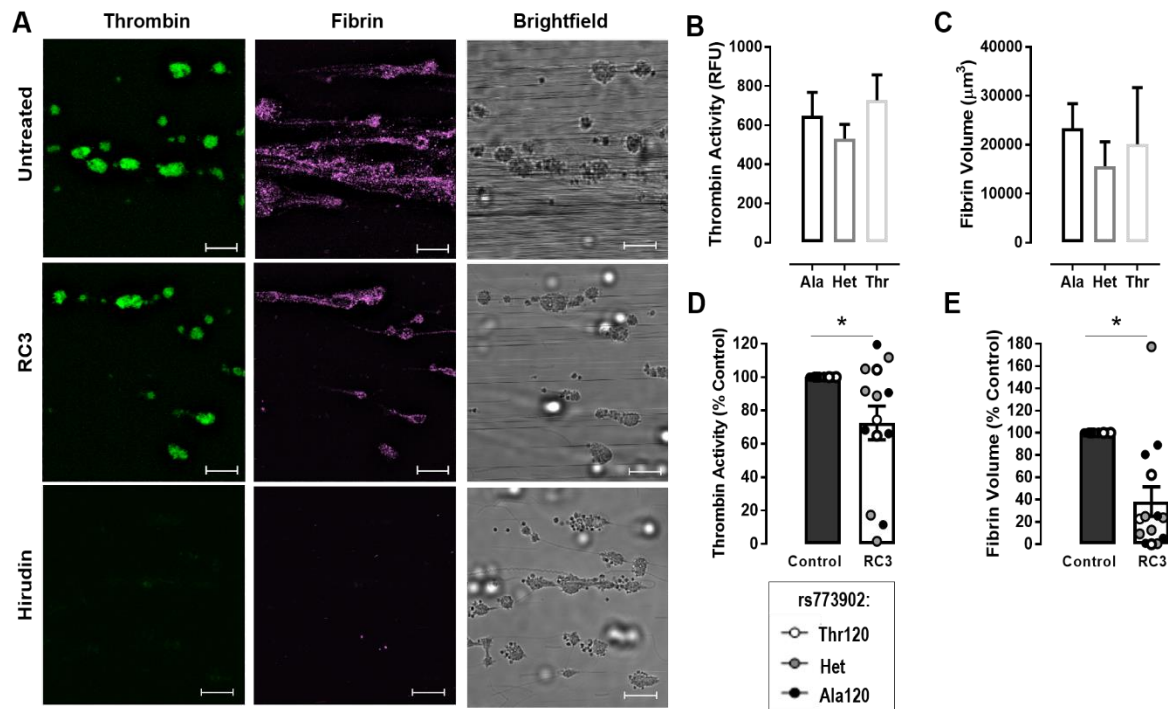


**Figure 3. mAb-RC3 inhibits the enhanced platelet aggregation elicited by the Thr120-PAR4 variant.**

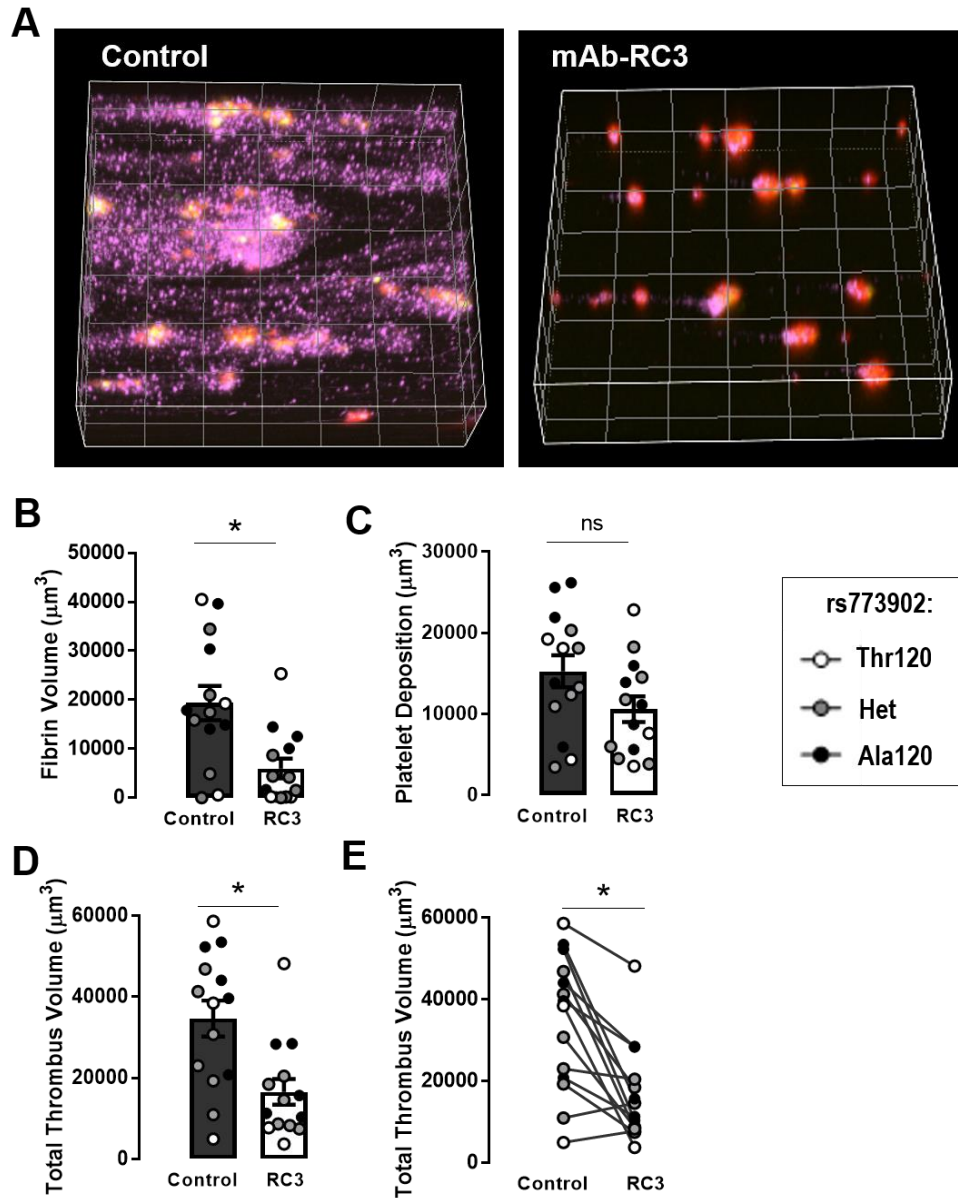
Aggregation of human isolated platelets from donors subsequently genotyped as homozygous at rs773902 for Thr120, Ala120, or as heterozygous. Shown are concentration-response curves to (A) PAR4-AP, (B) thrombin, and (C) thrombin in the presence of the PAR1 inhibitor vorapaxar (90 nM). Also shown are (D) concentration-inhibition curves to mAb-RC3 on responses to thrombin (0.1 U/mL) in the presence of vorapaxar (90 nM), indicating antibody-mediated PAR4 inhibition is equally effective across all genotypes, and (E)  $IC_{50}$  of mAb-RC3 derived from pooled data from (D). All data are mean  $\pm$  SEM from n=3-6 experiments per genotype. \*,  $P < 0.05$  (one-way ANOVA with Dunnet's test for multiple comparisons).



**Figure 4. mAb-RC3 inhibits the enhanced platelet procoagulant response elicited by the Thr120-PAR4 variant.** Production of procoagulant (annexin V-binding) platelets in human isolated platelet preparations from donors subsequently genotyped as homozygous at rs773902 for Thr120, Ala120, or as heterozygous. Shown are concentration-response curves to (A) PAR4-AP and (B) thrombin. Also shown (C) are concentration-inhibition curves to mAb-RC3 on responses to thrombin (1 U/mL), indicating antibody-mediated PAR4 inhibition is equally effective across all genotypes. All data are mean  $\pm$  SEM from n=3-6 experiments per genotype. \*,  $P < 0.05$  (one-way ANOVA with Dunnet's test for multiple comparisons).



**Figure 5. mAb-RC3 inhibits platelet procoagulant activity during *ex vivo* human whole blood flow.** (A) Representative images and (B-E) quantitation of human thrombi formed after 3 minutes of a whole blood thrombosis assay using donors subsequently genotyped as homozygous at rs773902 as Thr120, Ala120, or as heterozygous. (A) Shown are thrombin (green, Thr-SP), fibrin (purple, anti-fibrin antibody) and brightfield image of thrombi. Note that the direct thrombin inhibitor, hirudin (800 U/mL), abolished thrombin activity and fibrin volume despite continued platelet deposition. Scale bar = 20 $\mu\text{m}$ . No differences in (B) thrombin or (C) fibrin volume were observed between thrombi formed in individuals expressing the indicated PAR4 genotypes. Pre-treatment with mAb-RC3 (RC3, 100  $\mu\text{g}/\text{mL}$ ) inhibited both (D) thrombin activity and (E) fibrin volume compared to untreated control. Data is mean  $\pm$  SEM of N = 3-6 per genotype. \*,  $P < 0.05$  (unpaired Student's t test).



**Figure 6. mAb-RC3 inhibits human thrombus formation independently of PAR4 variant expression.** (A) Representative 3D reconstructions (each cubic edge = 20  $\mu\text{m}$ ) and (B-E) quantitation of human thrombi formed after 3 minutes of a whole blood thrombosis assay, showing platelets (red, anti-CD9), thrombin (green, Thr-SP), and fibrin (purple, anti-fibrin antibody). Note pre-treatment of blood with mAb-RC3 (RC3, 100 $\mu\text{g}/\text{mL}$ ) significantly inhibited (B) fibrin volume but not (C) platelet deposition. This translated to (D) a significant reduction in total thrombus volume. Data is mean  $\pm$  SEM of N = 14 (3-6 per genotype). A pairwise analysis is shown in (E).\*,  $P < 0.05$  (unpaired Student's t-test). Ns = not significant.



## Chapter 6 - General Discussion

### 6.1 Key findings of this thesis

The studies described in this thesis examined whether targeting the platelet thrombin receptor, PAR4, is a valid antithrombotic approach. Thrombin-induced platelet activation has been a long coveted antiplatelet strategy. However, the increases in bleeding associated with the clinical PAR1 antagonist, vorapaxar, suggested that safe and effective antithrombotic therapy would require a different approach to manipulating platelet-thrombin signalling. PAR4 is a lower affinity thrombin receptor, and has therefore drawn significant recent attention as a potential antithrombotic drug target that may have less impact on haemostasis. At the beginning of these studies, the function of PAR4 during platelet activation and thrombus formation in humans remained relatively unknown, in large part due to limitations in key reagents. In developing and utilising novel reagents to examine PAR4 function, the studies presented here have brought to light new roles of this receptor and the implications of its inhibition. The major findings of this thesis are:

- 1) PAR4 predominantly mediates thrombin-dependent platelet procoagulant activity – and subsequent thrombin and fibrin formation – largely independently of PAR1 (Chapter 2). This distinct antithrombotic mechanism suggests that PAR4 inhibition is a viable therapeutic strategy.
- 2) To assess PAR4 inhibition *in vivo*, two genetic mouse models were characterised: i) PAR4<sup>-/-</sup> mice, which have platelets that don't respond to thrombin (Chapter 3); and ii) a mouse genetically modified to mimic the human platelet-PAR expression profile (hPAR1-KI mice) (Chapter 4). Due to significant limitations that were revealed with both of these models, these studies suggest that alternative animal models will be required for meaningful preclinical evaluation of PAR4 antagonists.

- 3) A novel human monoclonal anti-PAR4 antibody was developed and characterised. This antibody provided marked antithrombotic effects, including equivalent effects against a common sequence variant in PAR4 that has been shown to be resistant to another PAR4 antagonist (Chapter 5). These studies indicate that blocking the thrombin-cleavage site of the receptor may be required for equivalent PAR4 inhibition across the population.

## **6.2 What is the role of PAR4 during thrombosis?**

This thesis began by endeavouring to determine the specific functions of PAR4 on human platelets during thrombosis, in order to rationalise whether PAR4 inhibition was a valid antiplatelet strategy. At the time of these studies, suitably specific and effective inhibitors of PAR4 were not freely available. Therefore a function-blocking anti-PAR4 antibody was developed and used to probe for platelet activation events reliant on PAR4. The studies of Chapter 2 demonstrate that PAR4 alone, and not PAR1, appears to be more important in the regulation of platelet procoagulant activity – namely, thrombin-induced phosphatidylserine (PS) exposure, consequent platelet-mediated thrombin generation, and fibrin formation.

This role for PAR4 during thrombosis is consistent with what is known regarding receptor structure, activation kinetics, and downstream signalling profiles. It is well known that PAR4 activation generates a more prolonged intracellular  $\text{Ca}^{2+}$  signal than PAR1 activation<sup>69</sup>, due to a longer interaction with thrombin<sup>35</sup> and slower desensitisation of the receptor<sup>80</sup>. It is also well known that a high and sustained rise in intracellular  $\text{Ca}^{2+}$  is required for PS externalisation, and that this surface binds tenase and prothrombinase complexes, leading to the local generation of thrombin and fibrin<sup>102</sup>. The studies in Chapter 2 build on this work by suggesting that PAR4 is predominantly responsible for mediating procoagulant effects of activated platelets, and demonstrating that PAR4 inhibition also directly impacts local thrombin and fibrin generation

under physiological flow conditions. Studies in Chapter 5 illustrate that PAR4 inhibition markedly reduces total thrombus volume, and that this reduction is predominantly due to impaired fibrin formation, rather than platelet deposition on collagen. Given that fibrin formation is an essential component of the stable clot, it follows that PAR4 inhibition could potentially impact overall thrombus stability, however further studies will be required to fully elucidate this. Regardless, these findings demonstrate that PAR4 acts at the interface of platelet aggregation and coagulation, and provides an antithrombotic effect in human blood via a distinct mechanism to other antiplatelet drug targets.

Whether PAR1 or PAR4 drives procoagulant responses has been controversial, with evidence of both receptors playing a role<sup>75, 103-105</sup>. However, this potentially relates to the lack of appropriate inhibitors available at the time of these earlier studies. The studies described in Chapter 2 and Chapter 5 of this thesis developed and characterised two effective and highly specific PAR4 inhibitors. Both function-blocking anti-PAR4 antibodies consistently inhibited thrombin-induced PS exposure, as well as whole blood thrombin and fibrin generation. Indeed, preclinical studies using the newest PAR4 antagonist, BMS-986120, support this notion<sup>98</sup>. BMS-986120 also demonstrated markedly more specificity for PAR4 over PAR1, as well as efficacy against thrombin-induced platelet activation<sup>98</sup>. When used *in vitro* it demonstrated Ca<sup>2+</sup> signalling profiles similar to the ones generated here using an anti-PAR4 antibody<sup>98</sup>. The reagents generated throughout the course of this thesis are therefore of considerable research interest, as they may be used to uncover additional functions of PAR4 in a variety of conditions, including functions in cell types other than platelets.

### **6.3 What is the likely clinical utility of PAR4 antagonism?**

PAR4 inhibition provides an antithrombotic effect via a mechanism distinct from all existing antiplatelet strategies. There are currently no antiplatelet drugs that directly target platelet

procoagulant activity, although the notion of specifically targeting this sub-population of activated platelets is gaining popularity<sup>106</sup>. The rationale for this is that targeting the procoagulant population preserves haemostatic functions of other platelets within the thrombus – namely those displaying the aggregating and contractile phenotype. Studies in Chapters 2 and 5 of this thesis show that specifically inhibiting PAR4 has no major impact on platelet deposition during human whole blood thrombosis, but does reduce fibrin deposition, which translates to a reduction in total thrombus volume. It is well documented that antiplatelet agents targeting platelet adhesion/aggregation pathways, and anticoagulant agents (global inhibition of thrombin) used in combination with the less efficacious antiplatelet drugs (aspirin/clopidogrel), significantly increase clinical bleeding events<sup>107</sup>. Therefore, as PAR4 antagonism acts at the interface of coagulation and platelet activation and leaves primary haemostatic functions intact (i.e. platelet adhesion and aggregation), it may lead to less clinical bleeding that is typically associated with other antiplatelet and anticoagulant drugs. Preclinical studies using the PAR4 antagonist, BMS-986120, strongly support this hypothesis. In non-human primate *in vivo* thrombosis models, PAR4 antagonism prevented thrombotic occlusion at doses that did not impact bleeding times – a therapeutic window not observed with clopidogrel<sup>98</sup>. However, it is of note that thrombin-induced platelet activation is less critical under conditions of high vascular shear<sup>46</sup>, such as those known to occur in stenosed arteries. Therefore, it is unknown whether PAR4 antagonism alone will be sufficiently effective in reducing thrombotic events formed under these conditions, or whether additional antiplatelet interventions will be required. In this regard, studies examining the combined effects of PAR4 antagonism with other agents appear pressing.

Although these preclinical studies show a promising profile with an increased therapeutic window, the overall effect of PAR4 inhibition on haemostasis will require further elucidation. The studies in Chapter 3 of this thesis were the first to reveal a spontaneous bleeding phenotype

in mice genetically deficient in PAR4 – a phenotype which has been overlooked for nearly two decades. Although PAR4<sup>-/-</sup> mice represent a model of complete inhibition of thrombin-platelet signalling, whereas human platelets exposed to a PAR4 antagonist would presumably retain PAR1 signalling, it is not known if the bleeding phenotype is due to the complete absence of thrombin-platelet signalling, or the absence of the physiological function of PAR4 during thrombosis (i.e. procoagulant platelet generation). Chapter 4 of this thesis aimed to generate a mouse model with equivalent platelet-PAR expression to humans, in order to answer this question. However, the failure to achieve PAR1 expression in mouse platelets, in the context of previous failed attempts<sup>108</sup> and other notable differences between murine and human platelet biology, ultimately indicate that other animal models will be more appropriate to conduct meaningful studies. There is, however, some human genetic evidence to guide the appropriateness of PAR4 inhibition as an antithrombotic approach. Patients with Scott syndrome – a rare congenital disorder in which defective platelet PS exposure drastically diminishes their thrombin generation capacity – have a clinically significant bleeding phenotype<sup>109</sup>. Therefore, further studies will be required to determine whether specifically targeting platelet procoagulant activity *via* PAR4 antagonism is a viable option for safe and effective antithrombotic treatment.

The issue of population genetics is another clinical consideration for PAR4 antagonism. Recent studies have revealed a commonly expressed genetic variant of PAR4 (rs773902; encoding either Ala<sup>120</sup> or Thr<sup>120</sup>). The Thr<sup>120</sup> PAR4 variant, expressed in 20 – 80% of people depending on the population, renders the receptor hyper-sensitive to agonists and hypo-sensitive to antagonists<sup>100, 101</sup>. Expression of the Thr<sup>120</sup> variant is racially dimorphic, and present in high proportions in American blacks, sub-Saharan Africa, as well as in Melanesian and Papuan populations<sup>101</sup>. In the American cohort, the demographic in which the Thr<sup>120</sup> PAR4 variant is dominant are known to have higher rates of cardiovascular disease, poorer outcomes,

and therefore more likely to be on antiplatelet drugs. On this point, the hyper-reactivity to PAR4 agonists observed in individuals with the Thr<sup>120</sup> PAR variant persists in the face of current antiplatelet drugs, including vorapaxar<sup>101</sup>. Although the studies in Chapter 5 suggest that the Thr<sup>120</sup> PAR4 variant alone is not sufficient to induce a heightened thrombotic phenotype in at least one *ex vivo* model, further studies with a greater number of donors and different thrombogenic conditions will be required to determine the extent to which PAR4 genotype has a direct impact on whole blood thrombosis. Regardless, the discovery of this commonly expressed PAR4 variant has led to PAR4 antagonism becoming highly desirable strategy for these at-risk populations.

#### **6.4 What does the future hold for PAR4 antagonists?**

The future of PAR4 antagonists is promising – there is a strong mechanistic rationale, preclinical studies have shown a desirable profile and potential to increase the therapeutic window, and the bio-pharmaceutical sector has an interest in continuing to develop improved compounds. The recently developed PAR4 antagonists – BMS-986120 and BMS-986141 – have undergone clinical trials. BMS-986120 was evaluated in a phase 1 dosing study, yet despite efficacy and a lack of adverse events no phase 2 studies of this compound were undertaken. Rather, BMS are investigating the related compound, BMS-986141, which underwent a phase 1 study (NCT02341638) and a subsequent phase 2 trial for the prevention of mini-stroke (NCT02671461). The trial (A Phase 2, Placebo Controlled, Randomized, Double-Blind, Parallel-Arm Study to Evaluate Efficacy and Safety of BMS- 986141 For the Prevention of Recurrent Brain Infarction in Subjects Receiving Acetylsalicylic Acid Following Acute Ischemic Stroke or Transient Ischemic Attack) had a primary efficacy endpoint of a composite of symptomatic ischemic stroke or unrecognized brain infarction, and a primary safety endpoint of a composite of adjudicated major bleeding and adjudicated clinically

relevant non-major bleeding during the treatment period. It was completed in April 2017 but has not yet been reported.

It is far too early to predict the likely clinical success and/or usefulness of PAR4 antagonists, and several key questions remain. Firstly, how well will PAR4 antagonism combine with current standard-of-care agents? This is a central point, since any trial will be conducted in the presence of standard-of-care, which frequently involves dual antiplatelet therapy. With the PAR1 antagonist vorapaxar, for example, the increased bleeding observed is believed to be due to poor compatibility with clopidogrel. Indeed, sub-study analyses show no additional bleeding in patients receiving aspirin plus vorapaxar versus those receiving aspirin alone<sup>48, 50</sup>. Here, it is interesting to note that BMS chose to investigate a patient group being treated with aspirin alone in its first phase 2 trial of its lead PAR4 antagonist. Also unknown are the specific indications most likely to be best served by a PAR4 antagonist. Again, sub-study analyses of the vorapaxar trials may provide pointers. These trials showed the most efficacy in reducing the rate of spontaneous myocardial infarction as well as in prevention of vascular complications associated with peripheral artery disease. This is perhaps unsurprising given the well-known role of thrombin generation in acute myocardial infarction, particularly in patients with a background of unstable angina and/or coronary artery disease<sup>53</sup>. Whether PAR4 antagonism will similarly demonstrate superior efficacy in these clinical situations where thrombin-induced platelet activation are implicated is an obvious place to start for future clinical trials.

Finally, the significant changes in PAR4 pharmacology associated with the Thr<sup>120</sup> PAR4 variant – namely, the resistance to a small molecule PAR4 antagonist – have raised the question as to whether these compounds will also be less effective in some patients. It is also currently unknown how the change in amino acid from Ala to Thr at position 120 alters receptor pharmacology. The crystal structure of PAR4 has not yet been solved, although extrapolations from the crystal structure of PAR1 show critical hydrogen bonds at a similar region<sup>110</sup>, which

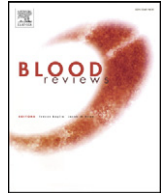
could potentially effect the small molecule binding pocket and/or tethered ligand binding site. One major finding from Chapter 5 of this thesis is that antibody-based inhibition, via blocking the thrombin cleavage site, provides equivalent antithrombotic effects regardless of donor genotype. Further studies will be required to determine how the variant alters receptor biology, what the impact and significance of it will be clinically, and, what method of PAR4 antagonism will provide equivalent inhibition across the population.

## **6.5 Concluding Remarks**

This thesis developed novel reagents to address key limitations in PAR4 research and used these to uncover previously unknown roles of PAR4 during thrombosis. In this way, these studies helped rationalise this receptor as a target for new antiplatelet drugs with a distinct mechanism of action. Although further insights are still to be gained regarding the utility of any PAR4 antagonism in clinical settings, the studies presented here have developed an antibody with therapeutic potential for the safe and effective prevention of arterial thrombosis that warrants investigation in further preclinical studies. These studies suggest targeting PAR4 represents a valid antithrombotic approach.



## **Appendix I**



## REVIEW

# Approval of the first protease-activated receptor antagonist: Rationale, development, significance, and considerations of a novel anti-platelet agent



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## ABSTRACT

Twenty-three years after the discovery of the first thrombin receptor, now known as protease-activated receptor 1 (PAR1), the first drug targeting this receptor is available for human use. The PAR1 inhibitor, vorapaxar (Zontivity, MSD), was recently approved by the FDA for use in the USA for the prevention of thrombotic cardiovascular events in patients with a history of myocardial infarction or peripheral artery disease. In this review, we detail the rationale, development, as well as the clinical significance and considerations of vorapaxar, the original PAR antagonist and the latest anti-platelet agent in the pharmacological armoury against arterial thrombosis.

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## 1. Introduction

## 1.1. There is a significant clinical need for improved anti-thrombotic therapy

Cardiovascular disease, manifesting predominantly as ischaemic heart disease and ischaemic stroke, is by far the most common cause of death and disability in the world, accounting for approximately 30% of all deaths [1]. Strikingly, despite increased awareness and improved management, and in contrast to most other high-impact communicable and non-communicable diseases, the rates of cardiovascular disease rose over the past decade [1]. The increasing burden of conditions such as diabetes, obesity, and depression, as well as an overall ageing population, are likely to ensure that cardiovascular disease rates continue to rise into the foreseeable future, with each of these conditions having increased prevalence of cardiovascular-related morbidity and mortality.

Arterial thrombosis precipitates the most prevalent cardiovascular disease manifestations, most notably acute myocardial infarction (MI), ischaemic stroke, and peripheral artery disease (PAD). Current therapies for the prevention of arterial thrombosis are either anti-platelet agents, which prevent platelet activation, or anticoagulants, which primarily impair fibrin formation at therapeutic doses by inhibiting either the production or activity of thrombin. The efficacy of these agents is tempered by the attendant bleeding risk [2]. Given that activated

platelets are the essential cellular component of arterial thrombi, anti-platelet drugs are the mainstay of current pharmacotherapy for arterial thrombosis, with the current recommendation for prevention of primary or secondary cardiovascular events in patients with acute coronary syndrome (ACS) being anti-platelet therapy alone [3]. Yet current standard-of-care treatments have limited efficacy and improved anti-platelet approaches are sought.

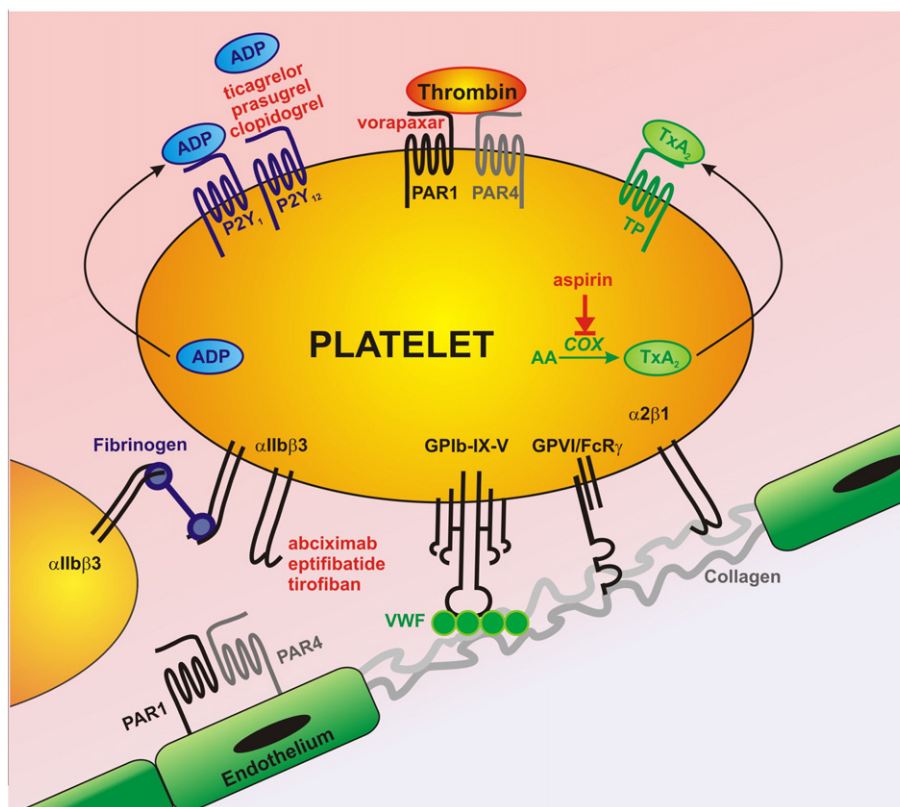
## 1.2. Current anti-platelet agents as anti-thrombotics

The processes controlling platelet function in the setting of arterial thrombosis have been extensively studied for the purpose of rationalising the development of improved anti-platelet therapies. Platelet behaviour during thrombus formation is generally agreed to involve, in order, cell adhesion, activation, and aggregation [4] (see Fig. 1). Understanding these processes has been highly informative in predicting the success of anti-platelet approaches.

- i) **Adhesion:** In response to vascular damage, such as atherosclerotic plaque rupture in the case of ACS, platelets adhere to the damaged vessel wall through a complex between sub-endothelial proteins and cognate receptors on the platelet surface. Rapid initial adhesion is mediated by von Willebrand factor (vWF) binding to the glycoprotein (GP) Ib-IX-V receptor complex [5], although this interaction does not support stable adhesion and a second adhesive step between collagen and GPVI and/or the integrin  $\alpha_2\beta_1$ , as well as fibrinogen and the integrin  $\alpha_{IIb}\beta_3$ , is required for firm platelet adhesion to the vessel wall [4].
- ii) **Activation:** following adhesion, the captured platelets are activated and undergo a series of morphological and functional changes

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**Fig. 1.** Activation and adhesion receptors on the platelet. In response to vascular damage, platelets rapidly adhere to the vessel wall via the interaction of subendothelial von Willebrand factor (VWF, green spheres) with the platelet glycoprotein (GP) Ib-IX-V complex, and collagen (grey strands) with GPVI and integrin  $\alpha_2\beta_1$ . Adherent platelets become activated, and this activation is enhanced by signalling pathways initiated by: thromboxane receptors (TP) activated by thromboxane A<sub>2</sub> (TxA<sub>2</sub>) generated from arachidonic acid (AA) by cyclooxygenase (COX); P2Y<sub>1</sub> and P2Y<sub>12</sub> receptors activated by ADP; and PAR1 and PAR4 activated by thrombin. Platelet activation culminates in the activation of integrin  $\alpha_{IIb}\beta_3$  which binds fibrinogen (blue dumbbell), VWF and fibronectin (not shown) to mediate platelet aggregation. Current anti-platelet agents (in red) include the P2Y<sub>12</sub> inhibitors clopidogrel, prasugrel and ticagrelor; the COX inhibitor, aspirin;  $\alpha_{IIb}\beta_3$  inhibitors, abciximab, eptifibatide and tirofiban; and the most recent addition, vorapaxar, which inhibits PAR1.

including shape change, spreading, release of granular contents, and the local generation of thrombin at the platelet surface as an endpoint of coagulation [6]. This platelet activation is primarily driven by the triumvirate of thrombin, ADP, and thromboxane A<sub>2</sub> (TxA<sub>2</sub>).

- iii) **Aggregation:** regardless of the adhesion and activation mechanism, the final common result is the activation of the integrin  $\alpha_{IIb}\beta_3$ , which engages fibrinogen (also vWF and fibronectin) to mediate platelet aggregation and ultimately thrombus formation [6].

Based on the experiences with current and previously trialled anti-platelet drugs, the most useful therapeutic strategy is to interfere with the activation phase of the platelet response. Specifically, interfering with platelet activation may largely preserve the primary haemostatic functions of the platelet by sparing initial adhesion at sites of vascular injury. By way of example, antagonists of the major platelet integrin,  $\alpha_{IIb}\beta_3$ , such as tirofiban, eptifibatide, and abciximab, are by far the most potent platelet inhibitors as they inhibit platelet aggregation regardless of the activating pathway. However, these drugs all cause significant bleeding complications [7] due to their disruption of the haemostatic function of platelets also dependent on this pathway. Although  $\alpha_{IIb}\beta_3$  inhibitors are currently indicated for acute use during percutaneous coronary intervention (PCI), the high bleeding risk and restrictive intravenous (i.v.) route of administration have limited their use in long-term preventative settings [8].

The current guidelines for treatment of ACS (comprising ST-elevation myocardial infarction (STEMI), non-ST-elevation myocardial infarction (NSTEMI), and unstable angina) and associated diseases are dual therapy with aspirin (an inhibitor of thromboxane A<sub>2</sub> synthesis) and a P2Y<sub>12</sub> receptor antagonist, such as the thienopyridines, clopidogrel

or prasugrel, or the cyclopentyl-triazolo-pyrimidine, ticagrelor [3] (American guidelines tend to favour the use of clopidogrel, whereas European guidelines favour either ticagrelor or prasugrel, with clopidogrel as an alternative.). However, aspirin and clopidogrel prevent just 15 and 17% of lethal cardiovascular events respectively and are only marginally more effective in combination [2]. In addition, an increasing number of patients are being reported as resistant to these agents [9–11], while clopidogrel – a prodrug – is metabolised with significant variability within the population [12], providing yet another source of variable efficacy. Large scale clinical trials assessing the efficacy of aspirin and clopidogrel (together and separately) in reducing thrombotic risk in cardiovascular disease (CAPRIE [13] and CHARISMA [14]) have ultimately determined greater efficacy when used in combination, with an associated minor increase in risk of bleeding complications. Therefore, existing anti-platelet therapies have limitations in one or both of safety and efficacy, with no current agent (or combination of agents) affording sufficiently potent, safe, and orally active prevention of arterial thrombosis. As a result, improved anti-platelet approaches are required to meet the significant clinical need for the safe and effective prevention of arterial thrombosis.

## 2. Targeting protease-activated receptors as an anti-platelet approach

### 2.1. Rationale

#### 2.1.1. Thrombin and thrombin receptors are important for platelet activation during thrombosis

The rationale for targeting thrombin-induced platelet activation as an anti-thrombotic approach has long held appeal. First, thrombin

functions at a time and place that is predicted to provide safe and effective anti-platelet activity. As outlined above, thrombin, in combination with ADP and TxA<sub>2</sub>, mediates the platelet activation required for thrombus growth that follows initial cell adhesion. Second, thrombin is the most potent endogenous activator of platelets. This suggests that targeting thrombin-induced platelet activation may provide greater efficacy over existing mechanisms that block the platelet activating functions of TxA<sub>2</sub> (aspirin) and ADP (clopidogrel and co.). Third, existing drugs that block the production (e.g. rivaroxaban) or activity (e.g. dabigatran) of thrombin are effective as anti-thrombotics. However these agents provide global inhibition of thrombin's actions and, consequently, incur a significant bleeding risk — particularly at the high doses used for the prevention of platelet-rich arterial thrombi. Specifically targeting thrombin-induced platelet activation while leaving the other functions of thrombin intact may mitigate the bleeding risk and provide a more selective effect for arterial thrombosis prevention. In combination, these factors provided the impetus to develop inhibitors of platelet thrombin receptors.

Platelet responses to thrombin are mediated by G protein-coupled protease-activated receptors (PARs) [15,16]. PARs are expressed on the surface of numerous cell types. In the cardiovascular system this includes platelets [17–19], and also leukocytes [20], vascular endothelial and smooth muscle cells [21–23], cardiomyocytes [24,25], and cardiac fibroblasts [26]. Humans express four PARs, with PAR1, PAR3 and PAR4 being activated by thrombin [17–19], and PAR2 by trypsin, tryptase, coagulation factors VIIa and Xa, and membrane-bound serine proteases MTSP1 and TMPRSS2 [27–31]. Human platelets express two thrombin-sensitive PARs, PAR1 and PAR4, and activation of either receptor is capable of inducing platelet activation [17,18,32]. Thrombin-induced platelet activation initiates platelet shape change, promotes platelet aggregation, and provides the procoagulant surface that facilitates secondary coagulation reactions [15,32,33]. PAR1 is the 'high affinity' thrombin receptor on human platelets, responding more sensitively and rapidly to thrombin than PAR4 [32]. However PAR4 can also trigger full platelet activation with higher concentrations (10 to 30-fold) of thrombin [32,34]. Based on this difference in affinity the clinical strategy for PAR inhibition in humans has focussed on blocking PAR1 function.

The cleavage-based PAR activation mechanism is unique. PARs are all activated the same way, whereby the protease agonist cleaves the amino-terminus of the receptor to reveal a cryptic neo-amino terminal sequence known as the "tethered ligand" [17,35]. The newly exposed tethered ligand then activates the receptor by binding intramolecularly to the second extracellular loop [17,35]. This self-activation prompts the conformational change of the receptor that allows interactions with G proteins of the G<sub>q</sub>, G<sub>12/13</sub> and G<sub>i/z</sub> families and consequent intracellular signalling events [36–38] (see Fig. 2). The interactions with G<sub>12/13</sub> drive Rho-dependent cytoskeletal responses involved in platelet shape change [38], whereas G<sub>q</sub>-mediated signalling facilitates the processes important for platelet granule release, activation of cell surface integrins, and platelet aggregation [39]. G<sub>i/z</sub> proteins are involved in inhibition of adenylyl cyclase, thereby removing a brake on platelet activation, while the G<sub>i</sub> protein family is the major source of  $\beta\gamma$  subunits involved in modifying activities of enzymes such as PI3 kinase [37].

Although often described as a dual platelet thrombin receptor system, some distinctions between PAR1 and PAR4 have been shown, including independent intracellular signalling events and distinct activation kinetics. PAR1 is a "high affinity" thrombin receptor, as it contains a hirudin-like sequence (K<sup>51</sup>YEPT<sup>55</sup>) which binds exosite I of the protease and aligns the active site of thrombin with a specific cleavage site in the receptor [40,41]. PAR1 is activated by trace amounts of thrombin (sub-nM range) and is responsible for the initial and rapid rise in intracellular calcium induced in platelets by the coagulation protease [42]. However, the PAR1-induced signal is transient and requires additional support from other platelet agonists such as ADP [43]. In contrast to PAR1, the interaction between PAR4 and thrombin occurs primarily at the active site due to the absence of a hirudin-like thrombin

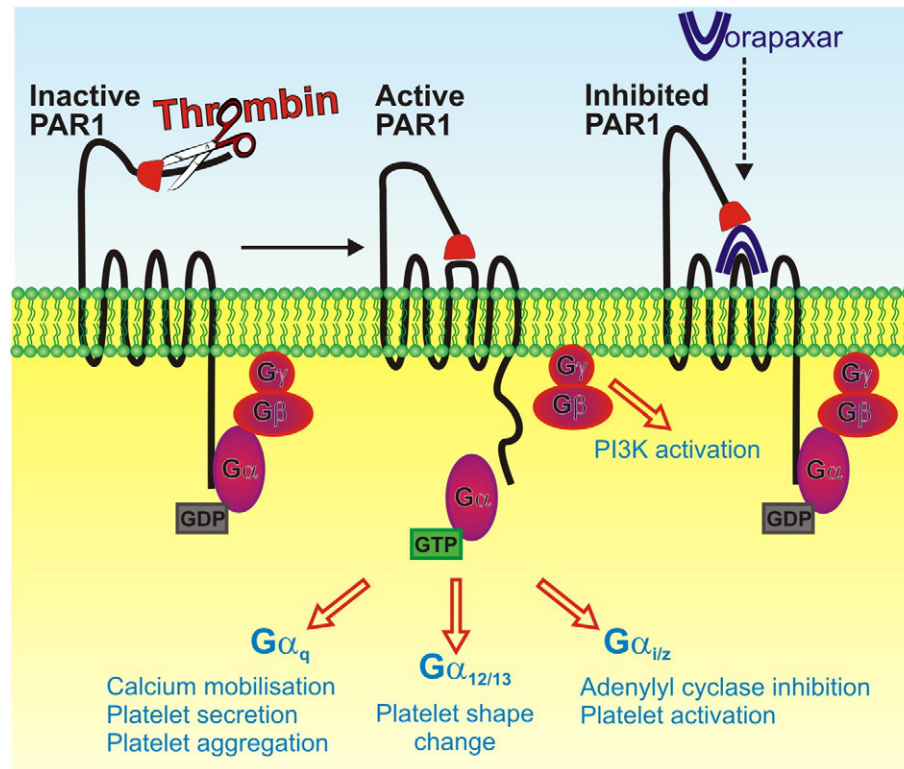
binding domain in the amino-terminal of PAR4. Despite the lack of interaction between PAR4 and exosite I of thrombin, the receptor does bind at the active site with high affinity due to two optimally-positioned proline residues immediately amino-terminal to the thrombin cleavage site [44]. This interaction facilitates slowed dissociation of thrombin from the receptor and results in a slower but more sustained intracellular signalling profile [44]. As a result of these distinct receptor activation and signalling kinetics, PAR4, in contrast to PAR1, is capable of inducing irreversible platelet aggregation in the absence of additional agonist activation [43]. It has therefore been suggested that PAR1 and PAR4 complement each other during thrombin-induced platelet aggregation. However, the major clinical focus has been on inhibiting PAR1 for anti-thrombotic activity due to the significantly greater sensitivity of this receptor to thrombin. Initial pre-clinical studies toward this end were performed in animal studies.

### 2.1.2. Pre-clinical studies

As is the case in many other systems, mouse genetic models and other small animal *in vivo* thrombosis experiments provided pre-clinical proof-of-concept studies. However the interpretation of these studies has always been hampered by the differing platelet PAR profile in these traditional model systems. Of the most commonly used animal models, only non-human primates appear to have an identical platelet PAR profile to humans (PAR1 and PAR4). In contrast, platelets from mice [45], rats [46], and rabbits [47] express PAR3 and PAR4, while platelets from guinea-pigs express PAR1, PAR3 and PAR4 [48]. Despite these limitations, significant insights have been gained from mouse genetic experiments. Mouse platelets do not express PAR1, but instead have PAR3 and PAR4. In a further difference, although mouse PAR4 functions in an analogous manner to human PAR4, mouse PAR3 is incapable of mediating transmembrane signalling by itself, instead functioning as a cofactor that facilitates cleavage and activation of PAR4 at low thrombin concentrations [49,50]. Indeed, PAR3 appears to act as a high affinity receptor for thrombin as it also contains a hirudin-like sequence to facilitate thrombin binding, therefore acting to bind and localise thrombin to the platelet surface and promoting PAR4 cleavage and subsequent signalling. As this model predicts, platelets from PAR4<sup>−/−</sup> mice are unresponsive to thrombin [45,51] and provide a clean genetic model to examine the overall contribution of thrombin-induced platelet activation in (patho)physiology. In this regard, PAR4<sup>−/−</sup> mice are healthy and exhibit no evidence of spontaneous bleeding, but are protected against thrombosis in several distinct *in vivo* models [45,51–53]. Haemostasis is impaired upon challenge as evidenced by an increase in tail bleeding time [45,51]. Perhaps surprisingly, PAR3<sup>−/−</sup> mice are also protected against thrombosis in a series of *in vivo* models, albeit to a lesser extent than observed with PAR4-deficiency [49]. This is of interest given that PAR3<sup>−/−</sup> mice may be viewed as a model of PAR1 inhibition in humans, inasmuch as both sets of platelets rely on PAR4 for thrombin-induced platelet activation. Of note, prolonged tail bleeding times were also observed in PAR3<sup>−/−</sup> mice [49]. Despite the limitations of using mice as a model of human platelet PAR function, these early proof-of-concept studies provided insight into the relative importance of thrombin-induced platelet activation in the setting of *in vivo* thrombosis, and in large part drove the development of PAR1 antagonists as a novel anti-platelet approach.

The development of pharmacological PAR1 inhibitors proved a significant challenge — perhaps largely due to the unique nature of the PAR self-activation mechanism and the consequent issues regarding efficacy for any potential antagonists that must compete with an endogenous agonist with a substantial steric advantage. In addition, receptor cleavage is irreversible and stoichiometrically efficient: multiple receptors can be activated by a single protease molecule. Numerous strategies have been employed to overcome these issues, including peptidomimetics derived from the tethered ligand sequence [54,55], intracellular signalling-inhibiting 'pepducins' [56], as well as receptor function-blocking antibodies [57], all of which have contributed to the





**Fig. 2.** Inhibition of PAR1 by vorapaxar. Thrombin cleaves the extracellular N-terminus of PAR1, exposing a tethered ligand (red cap) that binds to the second extracellular loop of the receptor and induces a conformational change that facilitates G protein interaction with the receptor and consequent intracellular signalling. PAR1 couples to G proteins of the  $G_q$ ,  $G_{12/13}$  and  $G_{i/z}$  families to induce a multitude of intracellular signalling events in platelets. Vorapaxar prevents PAR1 activation by binding at or near the tethered ligand binding site, acting as a competitive antagonist of the endogenous activation mechanism of the receptor.

understanding of PAR biology and led to the eventual development of effective and specific small molecule antagonists.

The earliest PAR inhibitors were peptide-based agents and function-blocking antibodies and provided important tools for experiments investigating platelet PAR function in human cells. The peptide-based inhibitors were synthesised to mimic the endogenous tethered ligand sequence, but with modifications that prevented the receptor being activated once the antagonist had bound. The structure–activity insight gained from the earliest peptidomimetics and further optimisation of this class led to the development of RWJ-58259, which inhibited PAR1-mediated aggregation of human platelets and thrombus formation *in vivo* and *in vitro* in non-human primates [58]. The earliest function-blocking anti-PAR1 antibodies were rabbit polyclonal antibodies targeted against the thrombin cleavage site of the receptor. These agents also inhibited PAR1-dependent aggregation of human platelets [57]. In keeping with the anti-thrombotic activity observed with peptidomimetic antagonists of PAR1, function blocking anti-PAR1 antibodies were also effective anti-thrombotics in non-human primate models [57,58]. Despite this effectiveness, the use of peptide-based agents and therapeutic antibodies in these studies came with the usual pharmacodynamic and pharmacokinetic limitations of such approaches and resulted in investigations into designing selective and potent small molecule antagonists of PAR1. This effort ultimately yielded two agents that were examined in clinical trials: atopaxar (E5555) and vorapaxar (SCH530348).

## 2.2. Development

### 2.2.1. Atopaxar

Atopaxar is a small molecule PAR1 inhibitor. Developed by Eisai and formerly known as E5555, atopaxar is a bicyclic amidine derivative with a molecular weight of 609 Da that competitively binds at or near the

tethered ligand binding site within the second extracellular loop of PAR1 [59]. In initial cell studies, atopaxar was shown to inhibit thrombin-induced aggregation of human platelets [60] and thrombin-induced intracellular calcium signalling in cultured smooth muscle cells [61]. It has an  $IC_{50}$  of 64 nM [60]. *In vivo*, atopaxar has been shown as an effective PAR1 antagonist in small animal models [60,61]. In rabbits, atopaxar reduced thrombin-dependent cerebral vasospasm – one of the major complications of subarachnoid haemorrhage – and was shown to dose-dependently decrease the basilar artery contractile response to thrombin as well as inhibit the upregulation of PAR1 expression in subarachnoid haemorrhage [61]. In a guinea pig model of photochemically-induced thrombosis, orally-administered atopaxar (30 mg/kg) prolonged the time to vessel occlusion by approximately 2-fold [60]. Given the promising efficacy of atopaxar in these small animal models, human trials were undertaken.

The safety of atopaxar was assessed in three phase 2 trials in patients with ACS (LANCLOT-ACS) and with chronic coronary artery disease (LANCLOT-CAD) [62–64]. These trials showed that overall clinically relevant bleeding increased numerically but not significantly, and that major bleeding rates increased in the ACS trial, but not in the CAD trial [62,63]. In addition, higher doses ( $\geq 200$  mg) resulted in an increase in liver enzymes and QT interval prolongation [62]. These effects were dose-dependent with significant liver function abnormalities observed in patients receiving the highest doses of atopaxar [62]. As a result, there are no phase 3 trials for atopaxar currently registered. For this reason, the remainder of this review will focus on the journey of vorapaxar through to FDA approval.

### 2.2.2. Vorapaxar

Vorapaxar is also a small molecule PAR1 inhibitor. Developed by Merck & Co (MSD; originally as Schering-Plough) and formerly known as SCH530348, vorapaxar is a synthetic tricyclic 3-phenylpyridine

analog of the naturally occurring alkaloid himbacine, and has a molecular weight of 591 Da. Similar to atropaxar, vorapaxar is a reversible, competitive antagonist of PAR1 that binds at or near the tethered ligand binding site within the second extracellular loop of PAR1, thus competing with the endogenous activation mechanism of the receptor [65]. Vorapaxar inhibits aggregation of human platelets induced by thrombin and a PAR1 agonist peptide (PAR1-AP) with an  $IC_{50}$  of 47 and 25 nM, respectively, and has a  $K_i$  of 8 nM [65]. Importantly, vorapaxar is administered as a bisulfate salt and has a high oral bioavailability (>90%) and is rapidly absorbed *via* the gastrointestinal tract, with a terminal plasma half-life of 126–269 h [66]. The drug is metabolised primarily by the liver *via* the CYP3A4 pathway and is eliminated almost exclusively as an amine metabolite [67]. As a result, drug interactions that have the potential to effect metabolism of vorapaxar include CYP3A4 inhibitors (e.g. ketoconazole) or inducers (e.g. rifampicin), which may increase or decrease the plasma concentration of vorapaxar respectively. In healthy subjects the drug was well tolerated and long-lasting: administration of a single loading dose (5 to 40 mg) inhibited PAR1-AP-induced platelet aggregation by >90% for more than 72 h [66]. Furthermore, administration of a daily dose of 1, 3, or 5 mg resulted in the same level of inhibition of the platelet aggregation through to day 7 of treatment [66]. To date, and in contrast to the phase 2 findings of atropaxar, no abnormalities in liver function or other adverse events have been associated with long term vorapaxar use.

Early human trials with PAR1 inhibitors showed trends toward clinical benefit in cardiovascular outcomes without an increase in bleeding risk. In a phase 2 trial, vorapaxar was assessed in 773 patients undergoing PCI and with a history of ACS and/or NSTEMI, most of whom were on dual antiplatelet therapy (aspirin plus clopidogrel) [66]. Patients received a loading dose of 10, 20, or 40 mg (or matching placebo) followed by a daily maintenance dose of 0.5, 1.0 or 2.5 mg. The primary endpoint was clinically significant major or minor bleeding, assessed by the Thrombolysis In Myocardial Infarction (TIMI) scale and was found to be not significantly different between the placebo and test groups, even at the highest dosing regimen [66]. Strikingly, although the study was not powered to determine efficacy, a trend toward a reduction in MI events was observed [66]. Given these promising findings, phase 3 trials of vorapaxar were developed using the highest dosing regimen from the phase 2 trials.

### 2.2.3. The Thrombin Receptor Antagonist for Clinical Event Reduction (TRACER) in acute coronary syndrome trial

Vorapaxar was assessed in two large scale phase 3 trials in patients with ACS (TRACER) and stable atherosclerosis (TRA 2°P-TIMI 50). The overall findings of these trials showed that vorapaxar, when combined with standard of care therapy, reduced the risk of cardiovascular events at the cost of increased bleeding [68,69].

TRACER was a randomised, double-blind, placebo-controlled multicentre trial that enrolled 12,944 patients with acute symptoms of coronary ischaemia plus either significant ST changes or elevations in cardiac necrosis markers (troponin or creatine kinase) [69]. Of the study group, 6473 patients received vorapaxar (40 mg single loading and 2.5 mg once daily maintenance) with 6471 receiving placebo. Ninety-six percent of all patients were receiving aspirin and 80% of those were also receiving a thienopyridine [69]. Assessments were performed during initial hospitalisation, with follow-up at 1, 4, 8, and 12 months post-hospitalisation, and every 6 months thereafter, with the aim of a total follow-up of 3 years. The primary end point for TRACER was a composite of cardiovascular death, MI, stroke, recurrent ischaemia, or urgent coronary revascularisation, with the secondary endpoint being a composite of death from cardiovascular causes, MI, or stroke. The key safety end points were a composite of moderate and severe bleeding according to the Global Use of Strategies to Open Occluded Coronary Arteries (GUSTO) classification as well as clinically significant bleeding according to the TIMI classification.

### 2.2.4. TRACER trial: efficacy

Ultimately, the primary endpoint of TRACER was not achieved: a non-significant reduction in the primary end point occurred in patients receiving vorapaxar (18.5%) versus placebo (19.9%) (hazard ratio (HR) 0.92; 95% confidence interval (CI) 0.85–1.01;  $P = 0.07$ ) [69]. The key secondary end point occurred in 14.7% versus 16.4% of patients. Given that superior efficacy of vorapaxar in this setting was not observed, subsequent subgroup analyses were performed but were purely exploratory [69,70]. Analysis of individual end point components revealed that the greatest reduction was observed in the rate of MI (11.1% versus 12.5%; HR 0.88; 95% CI 0.79–0.88;  $P = 0.02$ ). Interestingly, stroke rates between the two treatment groups were similar over 2 years: the vorapaxar group had lower rates of ischaemic stroke (1.1% versus 1.4%; HR 0.79; 95% CI 0.59–1.08;  $P = 0.14$ ) but a higher rate of haemorrhagic stroke (0.3% versus 0.1%; HR 2.73; 95% CI 1.22–6.14;  $P = 0.02$ ). Other efficacy endpoints are summarised in Table 1.

### 2.2.5. TRACER trial: safety

Patients receiving vorapaxar were more likely to experience bleeding events. Moderate or severe bleeding (assessed by GUSTO) occurred in 7.2% of patients receiving vorapaxar compared with 5.2% of those receiving placebo (HR 1.35; 95% CI 1.16–1.58;  $P < 0.001$ ). Clinically significant (TIMI) bleeding occurred in 20.2% of patients in the vorapaxar group versus 14.6% in the placebo group (HR 1.43; 95% CI 1.31–1.57;  $P < 0.001$ ). Most critically, however, intracranial haemorrhage increased more than 5-fold in the vorapaxar group (1.1% versus 0.2%; HR 3.39; 95% CI 1.78–6.45;  $P < 0.001$ ) [69]. This effect was especially noted in patients with a prior history of stroke or transient ischaemic attack (TIA) and increased incrementally over time. Consequently, the planned 3-year follow-up of TRACER was terminated in January 2011 (five months early) in response to recommendations from the data and safety monitoring board.

### 2.2.6. TRACER trial: subgroup analyses

Exploratory subgroup analyses of the TRACER trial have yielded several interesting findings that may warrant further evaluation. Firstly, analysis of MI reduction in TRACER showed that vorapaxar treatment resulted in a 12% hazard reduction (HR 0.88; 95% CI 0.79–0.98;  $P = 0.021$ ) of a first MI of any type [70]. Further, this reduction was largely accounted for by a specific decrease in the rate of type 1 (spontaneous) MI (5.9% vs 7.0%; HR 0.83; 95% CI 0.73–0.95;  $P = 0.007$ ). This is noteworthy, as spontaneous MIs are the most commonly observed: 65% of MIs observed in TRACER were type 1 and were the most frequently observed coronary endpoint. These observations from the original trial data were supported by a recent meta-analysis of all placebo-controlled trials of either atropaxar or vorapaxar in CAD patients [71]. In addition to MI, vorapaxar reduced the primary endpoint in coronary artery bypass graft (CABG) patients with an estimated hazard reduction of 45% (43 patients in the vorapaxar treatment group versus 70 in placebo; 8.2% and 12.9%, respectively, HR 0.55; 95% CI 0.36–0.83;  $P = 0.005$ ) [72]. There was a non-significant increase in both surgical and non-surgical bleeding in this cohort, and the overall results suggest that vorapaxar treatment may be a viable option in this setting. Investigations into the effect of standard-of-care therapy in addition to vorapaxar treatment indicated improved safety and efficacy in the vorapaxar groups who were not receiving high dose (>300 mg) aspirin and/or a thienopyridine [69,70].

### 2.2.7. TRACER trial: overall outcomes

Overall, TRACER concluded that vorapaxar in addition to standard anti-platelet therapy provided no significant net clinical benefit in patients with acute coronary syndromes. Furthermore, vorapaxar significantly increased the risk of major bleeding, most notably intracranial haemorrhage, in a subset of patients. These outcomes had a significant impact on the TRA 2°P-TIMI 50 trial. Firstly, the primary endpoint was revised to a composite of cardiovascular death, MI or stroke, with urgent

**Table 1**

Efficacy and safety endpoints of the TRACER and TRA 2°P-TIMI 50 trials.

	TRACER <sup>a</sup>				TRA 2°P-TIMI 50 <sup>b</sup>			
	Vorapaxar Number of events (percent)	Placebo Number of events (percent)	Hazard ratio (95% CI)	P value	Vorapaxar Number of events (percent)	Placebo Number of events (percent)	Hazard ratio (95% CI)	P value
Primary endpoint	n = 6473 1031 (18.5)	n = 6471 1102 (19.9)	0.92 (0.85–1.01)	0.07	n = 13,225 1028 (9.3)	n = 13,224 1176 (10.5)	0.87 (0.80–0.94)	<0.001 (NNT = 83)
Secondary endpoint	822 (14.7)	910 (16.4)	0.89 (0.81–0.98)	0.02	1259 (11.2)	1417 (12.4)	0.88 (0.82–0.95)	0.001 (NNT = 83)
Cardiovascular death or myocardial infarction	755 (13.5)	843 (14.9)	0.90 (0.81–0.99)	0.03	789 (7.3)	913 (8.2)	0.86 (0.78–0.94)	0.002 (NNT = 111)
Cardiovascular death	208 (3.8)	207 (3.8)	1.00 (0.83–1.22)	0.96	285 (2.7)	319 (3.0)	0.89 (0.76–1.04)	0.15
Myocardial infarction	621 (11.1)	698 (12.5)	0.88 (0.79–0.98)	0.02	564 (5.2)	673 (6.1)	0.83 (0.74–0.93)	0.001 (NNT = 122)
				(NNT = 84)				
Stroke								
Any	96 (1.9)	103 (2.1)	0.93 (0.70–1.23)	0.61	315 (2.8)	324 (2.8)	0.97 (0.83–1.14)	0.73
Ischemic	74 (1.1) <sup>c</sup>	93 (1.4)	0.79 (0.59–1.08)	0.14	250 (2.2)	294 (2.6)	0.85 (0.72–1.01)	0.06
Haemorrhagic	22 (0.3) <sup>c</sup>	8 (0.1)	2.73 (1.22–6.14)	0.02	Data not reported			
Urgent coronary revascularisation	203 (3.8)	189 (3.5)	1.07 (0.88–1.31)	0.49	279 (2.5)	316 (2.6)	0.88 (0.75–1.03)	0.11
Recurrent ischemia with hospitalisation	79 (1.6)	69 (1.5)	1.14 (0.83–1.58)	0.42	Data not reported			
Stent thrombosis definite or probable	61/3549 (1.7) <sup>c</sup>	54/3526 (1.5) <sup>c</sup>	1.12 (0.78–1.62)	0.54	Data not reported			
Death vs any cause	334 (6.5)	318 (6.1)	1.05 (0.90–1.23)	0.52	540 (5.0)	565 (5.3)	0.95 (0.85–1.07)	0.41
Safety Endpoints	n = 6446	n = 6441			n = 13,186	n = 13,166		
GUSTO								
Moderate or severe bleeding	391 (7.2)	290 (5.2)	1.35 (1.61–1.58)	<0.001	438 (4.2)	267 (2.5)	1.66 (1.43–1.93)	<0.001
Severe bleeding	144 (2.9)	87 (1.6)	1.66 (1.27–2.16)	<0.001	Data not reported			
TIMI								
Clinically significant	1065 (20.2)	755 (14.6)	1.43 (1.31–1.57)	<0.001	1759 (15.8)	1241 (11.1)	1.46 (1.36–1.57)	<0.001
Major bleeding	208 (4.0)	136 (2.5)	1.53 (1.24–1.90)	<0.001	Data not reported			
Non-CABG major bleeding	131 (2.7)	71 (1.3)	1.85 (1.39–2.47)	<0.001	287 (2.8)	198 (1.8)	1.46 (1.22–1.75)	<0.001
CABG major bleeding	62/639 (9.7)	49/671 (7.3)	1.34 (0.92–1.95)	0.13	11/175 (7.6)	10/210 (6.1)	1.13 (0.48–2.66)	0.79
Bleeding requiring medical attention	784 (15.2)	564 (11.2)	1.41 (1.26–1.57)	<0.001	Data not reported			
Fatal bleeding	15 (0.4)	8 (0.2)	1.89 (0.80–4.45)	0.15	29 (0.3)	20 (0.2)	1.46 (0.82–2.58)	0.19
Intracranial haemorrhage	40 (1.1)	12 (0.2)	3.39 (1.78–6.45)	<0.001	102 (1.0)	53 (0.5)	1.94 (1.39–2.70)	<0.001

CI = confidence interval; NTT = number to treat; TIMI = Thrombolysis In Myocardial Infarction classification; GUSTO = Global Utilization of Streptokinase and Tissue Plasminogen Activator for Occluded Coronary Arteries classification; CABG = coronary artery bypass graft.

<sup>a</sup> Primary endpoint is a composite of cardiovascular death, myocardial infarction, stroke, or urgent coronary revascularisation. Secondary end point is a composite of cardiovascular death, myocardial infarction or stroke. Percentages are cumulative Kaplan–Meier event rates at 2 years.

<sup>b</sup> Primary endpoint is a composite of death from cardiovascular causes, myocardial infarction or stroke. Secondary endpoint is a composite of death from cardiovascular causes, myocardial infarction, stroke or urgent coronary revascularisation. Percentages are cumulative Kaplan–Meier event rates at 3 years.

<sup>c</sup> Percentages are calculated from raw data, Kaplan–Meier event rates not included in data set.

coronary revascularisation becoming part of the secondary endpoint. Most importantly, patients enrolled in the TRA 2°P-TIMI 50 trial who had a history of TIA and/or stroke were removed from the trial in January 2011 (2 years post enrolment) after the increases in risk of intracranial bleeding were brought to light. These modifications permitted the completion of this trial.

## 2.2.8. Thrombin-Receptor Antagonist for Secondary Prevention of Atherothrombotic Ischaemic Events (TRA 2°P-TIMI 50) Trial

As with TRACER, TRAP 2°P-TIMI 50 was a randomised, double-blind, placebo-controlled, multicentre trial. It was a larger trial in which 26,449 patients were enrolled, 13,224 of whom received vorapaxar (2.5 mg daily, no loading dose) in addition to standard-of-care therapy [68]. Initial recruitment involved patients with stable atherosclerosis and a history of MI or ischaemic stroke (between 2 weeks and 12 months before enrolment) and/or PAD (either after revascularisation or with an ankle-brachial index of <0.85) [68]. As noted above, study treatment in patients who had a prior stroke was discontinued in January 2011, 2 years after the commencement of the trial. In the MI cohort, 98% were receiving aspirin and 78% were also receiving a thienopyridine; in the PAD patients, 88% were receiving aspirin and only 37% a thienopyridine. The primary end point was a composite of cardiovascular death, MI, or stroke, with the key secondary endpoint a composite of cardiovascular death, MI, stroke, or urgent coronary revascularisation. As with TRACER, the key safety endpoints were moderate or severe bleeding (GUSTO) and clinically significant bleeding

(TIMI). The trial covered 3 years, with follow-up visits at 1, 4, 8, and 12 months, and then every 6 months thereafter until trial completion.

### 2.2.9. TRA 2°P-TIMI 50 trial: efficacy

The TRAP 2°P-TIMI 50 trial achieved its primary endpoint. At 3 years, the primary end point had occurred in 9.3% of patients receiving vorapaxar compared with 10.5% in the placebo group (Kaplan–Meier estimates at 3 years; HR 0.87; 95% CI 0.80–0.94;  $P < 0.001$ ) [68]. The major secondary end point was also reduced (11.2% versus 12.4%; HR 0.88; 95% CI 0.83–0.95;  $P = 0.001$ ). Again, of the individual components of these end points, MI was most significantly reduced (5.2% versus 6.1%; HR 0.83; 95% CI 0.74–0.93;  $P = 0.001$ ). Despite these findings there was no significant difference in the rates of death from any cause (5.0% versus 5.3%; HR 0.95; 95% CI 0.85–1.07;  $P = 0.41$ ) [68]. A summary of these endpoints is provided in Table 1.

### 2.2.10. TRA 2°P-TIMI 50 trial: safety

As observed in TRACER, bleeding complications were increased in patients receiving vorapaxar: moderate–severe bleeding (GUSTO) was increased (4.2% versus 2.5%; HR 1.66; 95% CI 1.43–1.99;  $P < 0.001$ ), as was clinically significant bleeding and non-CABG-related major bleeding scores (TIMI) (15.8% versus 11.1% (HR 1.46; 95% CI 1.36–1.57) and 2.8 versus 1.8% (HR 1.46; 95% CI 1.22–1.75), respectively,  $P < 0.001$  for both) [68]. In contrast, there was no significant difference in the rates of CABG-related major bleeding – a lack of effect also observed in the TRACER trial. Intracranial haemorrhage was increased two-fold in the vorapaxar group (1.0% versus 0.5%, HR 1.94; 95% CI 1.39–2.70;



$P < 0.001$ ) but, most critically, fatal bleeding was rare and was not different between the treatment groups (0.3% versus 0.2%, HR 1.46; 95% CI 0.82–2.58;  $P = 0.19$ ) [68]. Bleeding was increased in women, those with a body weight  $< 60$  kg, and patients in the North American trials; however these groups did not test positive for significant interaction [68]. In contrast to TRACER, additional P2Y<sub>12</sub> therapy did not increase the bleeding risk of vorapaxar, yet concomitant aspirin treatment did. The aspirin-naïve patient subgroup did not show any increase in clinically relevant bleeding [68,73].

### 2.2.11. TRA 2°P-TIMI 50 trial: subgroup analyses

TRA 2°P-TIMI 50 was designed to allow analyses of the large, pre-defined subgroups comprising the three distinct atherosclerotic patient cohorts: those with prior MI, ischaemic stroke, or PAD. In support of the TRACER findings, particular benefit was observed in the 17,779 patients with prior MI (8.1% versus 9.7%; HR 0.80; 95% CI 0.72–0.89,  $P < 0.001$ ) [74]. Importantly, patients in this subgroup did not have increased rates of intracranial haemorrhage (0.6% versus 0.4%;  $P = 0.076$ ) [74]. Interestingly, of the 3787 PAD patients, vorapaxar treatment did not reduce the risk of primary or secondary endpoint [75], but did produce some benefit in reducing the risk of peripheral vascular end points, notably limb ischaemia (2.3% versus 3.9%; HR 0.58; 95% CI 0.39–0.86;  $P = 0.006$ ) and peripheral revascularisation (18.4% versus 22.2%; HR 0.84; 95% CI 0.73–0.97;  $P = 0.017$ ) [75]. In concordance with the TRACER trial results, the ischaemic stroke subgroup of TRA 2°P-TIMI recorded no significant efficacy with vorapaxar treatment but an increase in bleeding was observed [76].

As a result of these trials, in May 2014 the FDA approved vorapaxar for the prevention of thrombotic cardiovascular events in patients with a history of MI or PAD. The drug will be sold under the trade name Zontivity. Given the clinical findings from TRACER and TRA 2°P-TIMI, Zontivity will come with a boxed warning contraindicating use in patients with a history of stroke or TIA.

## 2.3. Significance & considerations

### 2.3.1. When will PAR1 antagonists be of most benefit?

Understanding the relative contribution of thrombin-mediated platelet activation to thrombus formation in particular pathologies will likely be of significant use in predicting the clinical success of PAR1 inhibitors. Overall, in patients with no history of stroke or TIA, and with a body weight above 60 kg, the data from TRA 2°P-TIMI 50 translated into 6 fewer cardiovascular deaths at the cost of two intracranial haemorrhages for every 1000 patients treated with vorapaxar [73] (see Table 2). However, detailed analysis of the clinical data with vorapaxar use showed increased benefit of PAR1 inhibition in distinct clinical cohorts. PAR1 inhibition demonstrated a consistent reduction in the rate of type 1 (spontaneous) MI in vorapaxar-treated patients across the phase 2 and phase 3 studies [70]. This is perhaps unsurprising given the well-known role of thrombin generation in acute MI, particularly in patients with a background of unstable angina and/or coronary artery disease [77–79]. That these patients have a noted increase in plasma GPV levels (a marker of thrombin-induced platelet activation) weeks after an acute MI event [78] presumably reflects long-term thrombin-induced platelet activation in this setting. On this point, it is intriguing to note that plasma GPV levels in STEMI patients correlate with the likelihood of occlusion of infarcted arteries and of early recanalisation of these vessels [79].

In addition to acute MI, vorapaxar was shown to reduce the complications associated with PAD, notably limb ischaemia and the requirement for peripheral vascularisation [75]. Such efficacy of PAR1 inhibition in peripheral arteries might also have been predicted from the much previous work investigating the role of thrombin in distinct pathologies. Numerous studies have shown thrombin inhibitors to be more effective at the lower shear rates experienced in peripheral arteries [80–82]. This phenomenon was previously thought due to decreased

**Table 2**

Suggested clinical place of vorapaxar in cardiovascular disease.

Indicated	Contraindicated
Patients with	History of:
1) Coronary artery disease	1) Stroke
2) Peripheral arterial disease	2) TIA
3) In combination with existing antiplatelet drugs	3) Intracranial haemorrhage
	Active pathological bleeding
	Underlying bleeding risk:
	1) Decreased renal and/or hepatic function
	2) Currently taking anticoagulants
	3) Under 60 kg and/or over 75 years
	4) History of bleeding disorders
	Pregnant or breastfeeding women

TIA = transient ischaemic attack;

fibrin production and a consequent impairment of thrombus stability [82], however more recent work has shown that thrombin-dependent platelet activation (via PARs) is also more prevalent at lower blood shear rates [83], presumably due to limited assembly of blood borne coagulation factors on the surface of activated platelets at higher blood flow/shear rates [84].

With the ever-present push toward personalised medicine, identifying the relative role of thrombin-induced platelet activation in patients may be of interest. Clearly, the one-size-fits-all approach to anti-platelet therapy provides a wide spectrum of responses to standard-of-care therapy. On this point, one recent study used thromboelastography and ADP-induced platelet aggregation to predict ischaemic events in aspirin- plus clopidogrel-treated patients 6 months after PCI. After stratification by platelet-fibrin clot strength measured by thromboelastography, this study found that 58% of patients in the highest quartile for ADP-induced aggregation (high residual platelet reactivity) had an ischaemic event by 2 months versus only 2% of patients in the lowest quartile [85,86]. In this regard, patients identified as having high residual platelet reactivity following standard-of-care treatment might receive the greatest benefit from additional administration of the new PAR1 antagonist.

### 2.3.2. How safe is PAR1 antagonism?

In addition to the specific pathologies likely to benefit from PAR1 inhibition (or not), understanding the interaction of vorapaxar with other current anti-platelet drugs is clearly of importance and requires further investigation given the discrepancies in the existing trial data. The TRACER trial showed that bleeding rates (GUSTO) increased in patients receiving vorapaxar in combination with a thienopyridine compared to those not (HR 1.45; 95% CI 1.23–1.71 (no thienopyridine) versus HR 0.95; 95% CI 0.65–1.40 (with thienopyridine);  $P = 0.04$  for interaction) [69]. In contrast, no increase in bleeding was observed in patients co-administered vorapaxar and a thienopyridine in the TRA 2°P-TIMI 50 trial [68]. A further difference between the trial results concerned concomitant aspirin usage, with patients receiving high dose aspirin ( $> 300$  mg daily at both baseline and time of discharge) recording a consistent, albeit non-statistically significant, trend toward higher bleeding and ischaemic outcomes in TRACER [87] but not in TRA 2°P-TIMI 50 [68]. The TRA 2°P-TIMI 50 trial did however report an increase in bleeding in North American patients, and it is worth noting that this patient group was considerably more likely to be receiving high dose aspirin than patients from other countries [68,87]. Regardless of these differences between the trials, bleeding in patients receiving multiple anti-platelet agents is likely to be a major consideration. It remains unknown whether the bleeding complications observed in these trials were due to PAR1 inhibition per se, the concurrent use of three anti-platelet drugs, or to a particular drug combination. On this point, it is worth noting that recent animal studies suggest that strategies that block platelet PAR function



**Table 3**  
Comparison of efficacy and safety of antithrombotic agents in patients with acute coronary syndromes.

Trial details	Vorapaxar	Clopidogrel	Prasugrel	Ticagrelor	Rivaroxaban
	TRA 2°P-TIMI-50 <sup>a</sup> Vorapaxar + aspirin / clopidogrel vs placebo; 2.5 mg daily/3 years; % = Kaplan–Meier at 3 years	CHARMISA <sup>c</sup> Clopidogrel (75 mg daily) + aspirin vs aspirin alone; % = Kaplan–Meier at 27.6 months	TRITON-TIMI-38 <sup>d</sup> Prasugrel + aspirin vs clopidogrel; 60 mg loading dose; 10 mg daily/15 months; % = Kaplan–Meier at 15 months	PLATO <sup>e</sup> Ticagrelor + aspirin vs clopidogrel; 180 mg loading dose; 90 mg twice daily/ 12 months; % = Kaplan–Meier at 12 months	ATLAS-ACS 2-TIMI-51 <sup>f</sup> Rivaroxaban + aspirin / clopidogrel vs placebo; 2.5 mg twice daily/31 months; % = Kaplan–Meier at 24 months
Patient characteristics	Number – 26,449 History of MI, ischemic stroke or PAD	Number – 15,603 Documented coronary disease, cerebrovascular disease, PAD or multiple atherothrombotic risk factors; over 45 years of age	Number – 13,608 Moderate to high-risk ACS with scheduled PCI	Number – 18,624 Hospitalisation for ACS, with or without ST-elevation	Number – 15,526 Recent ACS (hospitalisation within 7 days)
Composite of death from cardiovascular causes, myocardial infarction or stroke	Vorapaxar (%) vs placebo (%); HR (95% CI)	Clopidogrel + aspirin (%) vs aspirin (%); HR (95% CI)	Prasugrel (%) vs clopidogrel (%); HR (95% CI)	Ticagrelor (%) vs clopidogrel (%); HR (95% CI)	Rivaroxaban (%) vs placebo (%); HR (95% CI)
Myocardial infarction	9.3% vs 10.5% HR 0.87 (0.80–0.94) P < 0.001	6.8% vs 7.3% HR 0.93 (0.83–1.05) P = 0.22	9.9% vs 12.1% HR 0.81 (0.73–0.90) P < 0.001	9.8% vs 11.7% <sup>g</sup> HR 0.84 (0.77–0.92) P < 0.001	9.1% vs 10.7% HR 0.84 (0.72–0.97) P = 0.02
Stroke (any)	5.2% vs 6.1% HR 0.83 (0.74–0.93) P = 0.001	1.9% vs 2.0% (nonfatal) HR 0.88 (0.75–1.18) P = 0.59	7.3% vs 9.5% (nonfatal) HR 0.76 (0.67–0.85) P = <0.001	5.8% vs 6.9% HR 0.84 (0.75–0.95) P = 0.005	6.1% vs 6.6% HR 0.90 (0.75–1.09) P = 0.27
Death from any cause	2.8% vs 2.8% HR 0.97 (0.83–1.14) P = 0.73	1.9% vs 2.4% (nonfatal) HR 0.79 (0.64–0.98) P = 0.03	1.0% vs 1.0% (nonfatal) HR 1.02 (0.71–1.45) P = 0.93	1.5% vs 1.3% HR 1.17 (0.91–1.52) P = 0.22	1.4% vs 1.2% HR 1.13 (0.74–1.73) P = 0.56
Key bleeding end points (TIMI except where indicated)	5.0% vs 5.3% HR 0.95 (0.85–1.07) P = 0.41	4.8% vs 4.8% HR 0.99 (0.86–1.14) P = 0.90	3.0% vs 3.2% HR 0.95 (0.78–1.16) P = 0.64	4.5% vs 5.9% HR 0.78 (0.69–0.91) P < 0.001	2.9% vs 4.5% HR 0.68 (0.53–0.87) P = 0.002
Fatal bleeding	Non-CABG related major bleeding 2.8% vs 1.8% HR 1.46 (1.22–1.75) P < 0.001	Non-CABG related major bleeding 2.7% vs 1.3% HR 1.85 (1.39–2.47) P < 0.001	Non-CABG related major bleeding 2.4% vs 1.8% HR 1.32 (1.03–1.68) P = 0.03	Non-CABG related major bleeding 2.8% vs 2.2% HR 1.25 (1.03–1.53) P = 0.03	Non-CABG related bleeding 1.8% vs 0.6% HR 3.46 (2.08–5.77) P < 0.0001
Intracranial bleeding	Clinically significant 15.8% vs 11.1% HR 1.46 (1.36–1.57) P < 0.001	Clinically significant 20.2% vs 14.6% HR 1.43 (1.31–1.57) P < 0.001	Major bleeding 7.9% vs 7.7% HR 1.03 (0.93–1.15) P = 0.57	Major bleeding 7.9% vs 7.7% HR 1.03 (0.93–1.15) P = 0.57	TIMI bleeding requiring medical attention 12.9% vs 7.5% HR 1.79 (1.55–2.07) P < 0.001
	0.3% vs 0.2% HR 1.46 (0.82–2.58) P = 0.19	0.3% vs 0.2% HR 1.53 (0.83–2.82) P = 0.17	0.4% vs 0.1% HR 4.19 (1.58–11.11) P = 0.002	0.3% vs 0.3% HR 0.87 (0.48–1.59) P = 0.66	0.1% vs 0.2% HR 0.67 (0.24–1.89) P = 0.45
	1.0% vs 0.5% HR 1.94 (1.39–2.70) P < 0.001	1.1% vs 0.3% HR 0.96 (0.56–1.65) P = 0.89	0.3% vs 0.3% HR 1.12 (0.58–2.15) P = 0.74	0.3% vs 0.2% HR 1.87 (0.98–3.58) P = 0.06	0.4% vs 0.2% HR 2.83 (1.02–7.86) P = 0.04

HR = hazard ratio; CI = confidence interval; MI = myocardial infarction; PAD = peripheral artery disease; ACS = acute coronary syndromes; PCI = percutaneous coronary infusion; TIMI = Thrombolysis In Myocardial Infarction classification; GUSTO = Global Utilization of Streptokinase and Tissue Plasminogen Activator for Occluded Coronary Arteries classification; CABG = coronary artery bypass graft.

<sup>a</sup> TRA 2°P-TIMI-50 (Thrombin-Receptor Antagonist in the Secondary Prevention of Atherothrombotic Events), Steering Committee, 2012. ClinicalTrials.gov number, [NCT00526474](#).

<sup>b</sup> TRACER (Thrombin Receptor Antagonist Vorapaxar in Acute Coronary Syndromes) Steering Committee, 2012. ClinicalTrials.gov number, [NCT00527943](#).

<sup>c</sup> CHARISMA (Clopidogrel for High Atherothrombotic Risk and Ischemic Stabilization, Management, and Avoidance) Steering Committee, 2006. ClinicalTrials.gov number, [NCT00050817](#).

<sup>d</sup> TRITON TIMI-38 (Trial to Assess Improvement in Therapeutic Outcomes by Optimizing Platelet Inhibition with Prasugrel) Steering Committee, 2007. ClinicalTrials.gov number, [NCT00097591](#).

<sup>e</sup> PLATO (The Study of Platelet Inhibition and Patient Outcomes) Steering Committee, 2009. ClinicalTrials.gov number, [NCT00391872](#).

<sup>f</sup> ATLAS-TIMI-51 (Anti-Xa Therapy to Lower Cardiovascular Events in Addition to Standard Therapy in Subjects with Acute Coronary Syndrome) Steering Committee, 2012. ClinicalTrials.gov number, [NCT00809965](#). Data are presented as modified intention to treat mITT analysis with corresponding p values.

<sup>g</sup> PLATO's primary endpoint was defined as a composite of death from vascular causes, myocardial infarction or stroke. Death from vascular causes was defined as death from cardiovascular causes and any death with or without another known cause.

are more likely to provide safe and effective anti-thrombotic activity when used in combination with aspirin than with clopidogrel [52,88]. Whether or not vorapaxar is likely to be useful in patients classified as “clopidogrel non-responders” remains to be determined but is likely to be of value. Of note, such “clopidogrel non-responders” have been shown to exhibit increased platelet aggregation in response to PAR1 activation [89], further emphasising the potential for vorapaxar in this patient group.

In addition to the moderate increase in bleeding rates across the entire patient cohort, the phase 3 trials of vorapaxar revealed important and significant safety concerns for specific demographics (see Table 2). The FDA approval of vorapaxar came with explicit contraindications for patients with a history of stroke, TIA, intracranial haemorrhage, or active pathological bleeding. Vorapaxar use in geriatric patients or those with severe hepatic impairment was also not recommended due to the inherent increase in bleeding risk in these demographics. Safety and efficacy of vorapaxar in paediatrics or in pregnant or nursing women have not yet been established, but given the controversies surrounding anti-thrombotic management of these populations [90,91] further studies into the safety and efficacy of vorapaxar in these patient groups may be warranted.

Finally, despite a lack of overt off-target effects of vorapaxar throughout these clinical trials, the widespread expression pattern of PAR1 in the cardiovascular system (and beyond) suggests that the long-term effects of systemic PAR1 inhibition will require careful consideration.

### 2.3.3. How does vorapaxar compare with other antithrombotic agents?

Several large-scale clinical trials have been conducted to determine the optimal anti-thrombotic management of ACS patients, including studies on the anti-platelet drugs aspirin, clopidogrel [13,14], prasugrel [92], and ticagrelor [93], as well as anticoagulants such as the direct Xa inhibitor, rivaroxaban [94]. A comparison of these trial results is shown in Table 3. While clearly inappropriate to directly compare the safety and efficacy of these agents across distinct trials with their disparate clinical groups, it is of interest to observe that each drug shows benefit over standard of care therapy in the reduction of at least one clinical end point. With the recent addition of vorapaxar to the anti-thrombotic pharmacological armory, an updated review of the trial data may be warranted.

### 2.3.4. What does the future hold for PAR inhibitors?

The use of vorapaxar beyond the well-defined settings and treatment regimens indicated by TRA 2°P-TIMI 50 awaits further investigation, and approval outside of the United States lies ahead. Given that vorapaxar is orally available, reversible, and sufficiently potent, there are no “second generation” PAR1 antagonists on the horizon as yet. For the time being, it is very much a case of “wait and see” for this first PAR1 inhibitor to be approved. What does the longer term future hold for platelet PAR inhibitors?

With the significant clinical effort behind the development of PAR1 inhibitors as anti-thrombotics, it is surprising that relatively little is known regarding the contribution of PAR4 to human platelet function during arterial thrombosis. Whether or not PAR4 represents a useful anti-platelet target is unknown, although some evidence suggests that PAR4 has distinct functions to PAR1 on human platelets. If true, PAR4 inhibitors may provide a useful adjunct and/or stand-alone therapy for the prevention of arterial thrombosis. For example, previous studies have shown that PAR1 and PAR4 differentially release distinct alpha-granule populations [95] and mediate diverse contributions to the procoagulant function of platelets [96]. Recent work has focussed on developing suitable tools to allow investigations of platelet PAR4 function, in particular PAR4-specific antagonists and mouse models in which the platelet PAR profile is ‘humanised’ to express PAR1 and PAR4 (rather than PAR3 and PAR4). To this end, several PAR4 antagonists have been developed, including pepducins [97], function-blocking antibodies [98]

and the small molecule antagonist, YD-3 [99]. Initial proof-of-concept studies have shown that PAR4 inhibition provides anti-thrombotic activity, albeit in caveated mouse models [53,98,100] and *in vitro* studies [96,101]. Unfortunately, recent efforts to develop a ‘humanised’ platelet PAR mouse have been less productive, with an unsuccessful attempt to express PAR1 as a transgene in PAR3-deficient mice [102]. Further insights into platelet PAR biology utilizing these new tools will improve our understanding of the relative roles of platelet thrombin receptors in (patho)physiology, potentially providing additional rationalisation of these receptors as therapeutic targets.

## 3. Conclusion

The recent approval by the USA's Food and Drug Administration of the first PAR1 antagonist as a novel anti-platelet agent represents a potentially important breakthrough in the treatment of thrombotic cardiovascular events and marks the long path from receptor discovery to clinical drug use. While the journey from the initial discovery of this platelet thrombin receptor in the early 1990s to the approval for use of the first receptor antagonist in 2014 represents a significant achievement and advance in our knowledge of this important platelet activation mechanism, it is not unreasonable to think that significant further insights are yet to be gained.

## Conflicts of interest

The authors declare no conflicts of interest.

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## **Appendix II**

# REVIEW ARTICLE

## Protease-activated receptor 4: from structure to function and back again

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Protease-activated receptors are a family of four GPCRs (PAR1–PAR4) with a number of unique attributes. Nearly two and a half decades after the discovery of the first PAR, an antagonist targeting this receptor has been approved for human use. The first-in-class PAR1 antagonist, vorapaxar, was approved for use in the USA in 2014 for the prevention of thrombotic cardiovascular events in patients with a history of myocardial infarction or with peripheral arterial disease. These recent developments indicate the clinical potential of manipulating PAR function. While much work has been aimed at uncovering the function of PAR1 and, to a lesser extent, PAR2, comparatively little is known regarding the pharmacology and physiology of PAR3 and PAR4. Recent studies have begun to develop the pharmacological and genetic tools required to study PAR4 function in detail, and there is now emerging evidence for the function of PAR4 in disease settings. In this review, we detail the discovery, structure, pharmacology, physiological significance and therapeutic potential of PAR4.

### LINKED ARTICLES

This article is part of a themed section on Molecular Pharmacology of G Protein-Coupled Receptors. To view the other articles in this section visit <http://onlinelibrary.wiley.com/doi/10.1111/bph.v173.20/issuetoc>

### Abbreviations

DRG, dorsal root ganglion; IP<sub>3</sub>, inositol trisphosphate; MLC, myosin light chain; PAR, proteinase-activated receptor; TBD, thrombin binding domain

### Tables of Links

TARGETS	
GPCRs <sup>a</sup>	Enzymes <sup>c</sup>
α <sub>2A</sub> -Adrenoceptor	Calpain
B <sub>2</sub> receptor	Cathepsin G
P2Y <sub>12</sub> receptor	MAPK
PAR1	PKC
PAR2	PLA <sub>2</sub>
PAR3	PLCβ
PAR4	Rho-kinase
Ligand-gated ion channels <sup>b</sup>	
P2X1 receptors	

LIGANDS
ADP
Bradykinin
Thrombin
Trypsin
TXA <sub>2</sub> , thromboxane A <sub>2</sub>
Vorapaxar

These Tables list key protein targets and ligands in this article which are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Pawson *et al.*, 2014) and are permanently archived in the Concise Guide to PHARMACOLOGY 2015/16 (<sup>a,b,c</sup>Alexander *et al.*, 2015a,b,c).

## Introduction

Protease-activated receptors (PARs) have been studied in some detail since the discovery of the prototype receptor, PAR1, in 1991. Humans express four PARs, designated PAR1 to PAR4 (please refer to the Concise Guide to Pharmacology; Alexander *et al.*, 2015a,b,c), that are broadly expressed and have diverse functions across multiple physiological systems. PARs form a small GPCR family characterized by their unique self-activation mechanism following cleavage by specific serine proteases – typically key effector proteases of coagulation and inflammatory pathways. As a result, extensive research has been conducted into the roles of PARs in thrombosis and other cardiovascular diseases, as well as a range of inflammatory conditions. To date, the research spotlight has been firmly focussed on PAR1 and PAR2, with PAR3 and PAR4 largely waiting in the wings.

PAR1 was discovered while attempting to elucidate the mechanism by which thrombin activates platelets and other cells (Vu *et al.*, 1991). PAR1 is a high-affinity thrombin receptor (Vu *et al.*, 1991). The other receptors of this family were subsequently cloned: PAR3 and PAR4 are also activated by thrombin (Ishihara *et al.*, 1997; Xu *et al.*, 1998; Kahn *et al.*, 1998b), with PAR2 thrombin-insensitive and primarily activated by trypsin and trypsin-like proteases (Nystedt *et al.*, 1995). The structure and function of PAR1 and PAR2 are well characterized, and have aided the clinical development of inhibitors of these receptors. The first PAR1 antagonist, vorapaxar, (trade name Zontivity), was approved in late 2014 by the FDA for use in the USA as an anti-platelet agent in patients with cardiovascular disease (Tricoci *et al.*, 2012; Morrow *et al.*, 2012; French *et al.*, 2015). In addition, PAR2 antagonists have undergone pre-clinical studies for the treatment of inflammation in a variety of settings, including arthritis (Kelso *et al.*, 2006; Lohman *et al.*, 2012). Despite these significant advances, there has been comparatively little progress in understanding the functions of PAR4 in health and disease and in targeting this receptor for therapeutic gain. Recent setbacks in the clinical development of PAR1 and PAR2 inhibitors, coupled with an emerging body of evidence suggesting potential utility of targeting PAR4 in disease settings, has re-ignited interest in the physiology and pharmacology of PAR4. This review outlines our current understanding of the physiology, pharmacology, clinical significance and therapeutic potential of PAR4, provided by the recent scientific advances in this area.

## Receptor structure

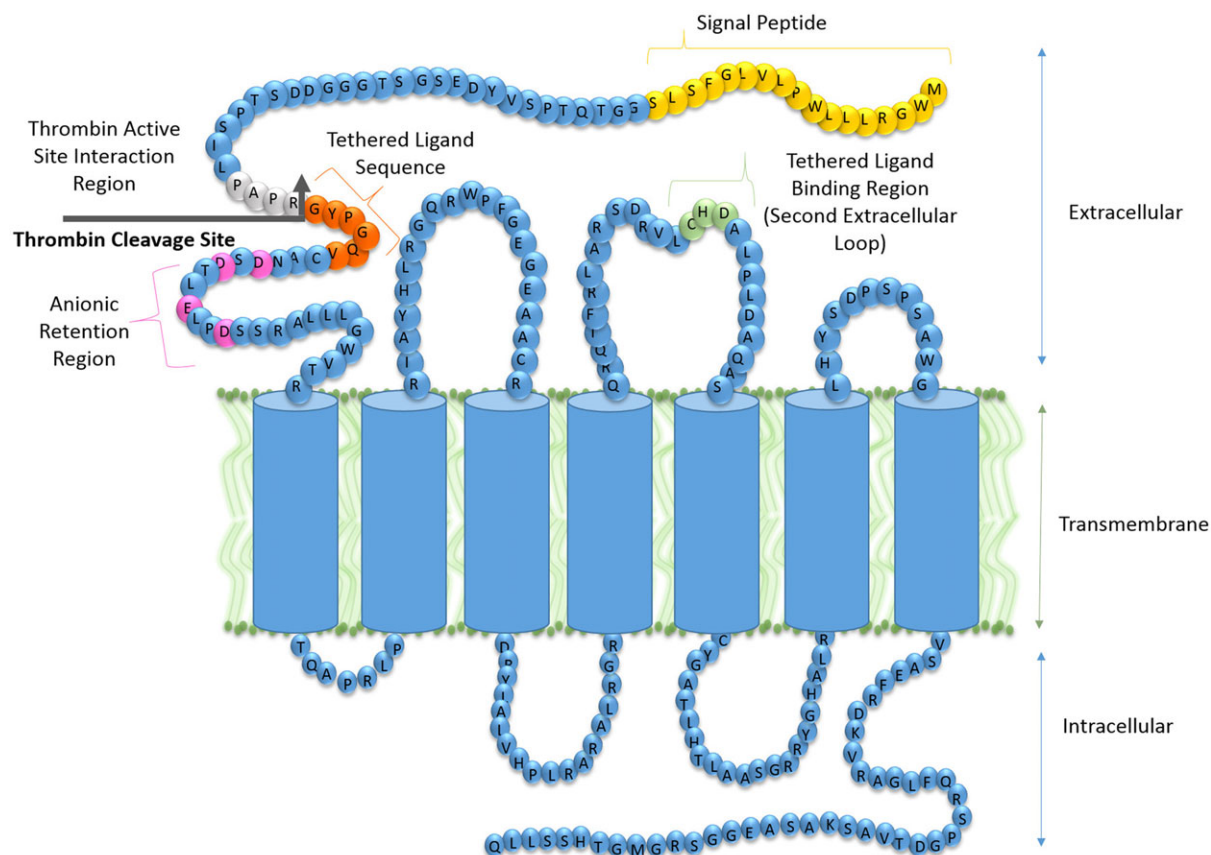
PAR4 was first cloned in 1998 (Xu *et al.*, 1998; Kahn *et al.*, 1998b) and is the most recently cloned member of the PAR family. It is a 300 amino acid seven transmembrane-spanning domain GPCR (Figure 1) that retains most of the core features of the other PARs despite its genetic differences from the other members of the family (Xu *et al.*, 1998; Kahn *et al.*, 1998b). The human PAR4 gene is remarkably smaller than those of the other PARs, and it resides at a distinct location on chromosome 19p12, with the genes for PARs 1–3 located in tandem on chromosome 5q13 (Xu *et al.*, 1998; Kahn *et al.*, 1998a). This genetic divergence of

PAR4 from the other PARs is thought to have arisen from a remote gene duplication and subsequent translocation event which gave rise to ancestral PAR4 and PAR1/PAR2/PAR3 genes (Kahn *et al.*, 1998a). More recently, two gene duplication events have occurred to separate PAR1 and PAR2 from PAR3; and then PAR1 from PAR2 (Kahn *et al.*, 1998a). These genetic differences may underlie some of the key differences in receptor structure and function within the PAR family.

The N-terminus of PAR4 contains a hydrophobic signal peptide sequence, with a signal peptidase cleavage site present at Ser<sup>17</sup>/Gly<sup>18</sup>. The extracellular amino-terminus also contains a serine protease cleavage site at Arg<sup>47</sup>/Gly<sup>48</sup> that is essential for receptor activation. Mutation of the serine protease cleavage site (Arg<sup>47</sup> → Ala) renders the receptor completely unresponsive to proteolytic activation. Site-specific receptor cleavage unmasks a cryptic tethered ligand sequence (GYPGQV) (Xu *et al.*, 1998), which then binds intramolecularly to a defined region in the second extracellular loop of the receptor (Figure 1), resulting in conformational change of the receptor and subsequent coupling to intracellular effectors, in common with other GPCR family members. While this proteolytic activation mechanism is common amongst PARs, there are several notable differences in the structure of PAR4 and the other receptors of the family. First, both the extracellular amino-terminus and intracellular carboxy terminus have little sequence similarity to the corresponding regions of other PARs (Xu *et al.*, 1998). Second, the tethered ligand binding site of PAR4 contains only three core amino acids (CHD; Figure 1) of the consensus sequence conserved in PARs 1–3 (ITTCHDV) (Xu *et al.*, 1998). PAR4 also lacks the high-affinity thrombin binding domain that is present in the other two thrombin receptors, PAR1 and PAR3 (Vu *et al.*, 1991; Mathews *et al.*, 1994; Ishihara *et al.*, 1997; Kahn *et al.*, 1998b).

## Receptor activation and membrane trafficking

The serine proteases capable of cleaving PAR4 are generally key regulators of coagulation (e.g. thrombin and coagulation factor Xa) and/or inflammation (e.g. trypsin and cathepsin G released from neutrophils, and bacterial proteases such as gingipains) (Xu *et al.*, 1998; Kahn *et al.*, 1998b; Sambrano *et al.*, 2001; Cottrell *et al.*, 2004). Additional serine proteases that are specific for arginine/lysine cleavage include coagulation factors VIIa, IXa, Xa, XIa, urokinase and plasmin; however, none of these have significant activity on PAR4, with the exception of factor Xa, which showed small effects at non-physiological concentrations (Xu *et al.*, 1998). Initial studies indicated that PAR4 is activated by thrombin and trypsin at similar concentrations, with an EC<sub>50</sub> of ~5 nM each (Xu *et al.*, 1998) – significantly higher than the EC<sub>50</sub> for thrombin at either PAR1 or PAR3 (~0.2 nM). The comparatively lower affinity of PAR4 for thrombin is a reflection of differences in the macromolecular association between the receptor and enzyme. PAR1 is a high-affinity thrombin receptor due to the presence of a hirudin-like thrombin binding domain (TBD) in its N-terminal exodomain (Mathews *et al.*, 1994). The TBD sequence, K<sup>51</sup>YE<sup>55</sup>PF<sup>55</sup>, interacts with exosite I of thrombin and upon binding causes significant allosteric



### Figure 1

Protease-activated receptor 4: structure. The proposed structure of the 300 amino acid GPCR, human PAR4. Site-specific proteolytic cleavage of the receptor amino-terminus (thrombin cleavage site, Arg<sup>47</sup>/Gly<sup>48</sup>; black arrow) reveals a neo-amino-terminus (tethered ligand sequence; G<sup>48</sup>YPGQV; orange) which binds intramolecularly to the second extracellular loop of the receptor (tethered ligand binding region; green). PAR4 contains two regions for enhanced thrombin interaction – the anionic retention region (pink) which interacts with thrombin's exosite I; and the PAPR sequence (grey) which binds at the active site of thrombin with high affinity.

effects essential for rapid association of thrombin (Jacques *et al.*, 2000). PAR4 does not contain this TBD (Xu *et al.*, 1998). In fact, the evidence indicates that PAR4 has only limited interaction with exosite I of thrombin, as  $\gamma$ -thrombin (which lacks a functional exosite I) activates PAR4 as effectively as  $\alpha$ -thrombin (Xu *et al.*, 1998), and mutations in the exosite I of thrombin have significantly less effect on cleavage of PAR4 than of PAR1 (Ayala *et al.*, 2001). Instead, PAR4 primarily interacts with the active site of thrombin via two optimally positioned proline residues (P<sup>44</sup> and P<sup>46</sup>) in the receptor (Jacques and Kuliopulos, 2003) just upstream of the thrombin cleavage site at Arg<sup>47</sup>/Gly<sup>48</sup>. A possible role for Leu<sup>43</sup> in the thrombin–receptor interaction has also been identified and believed to be in facilitating high-affinity binding of the PAR4 amino-terminal to thrombin's active site (Jacques and Kuliopulos, 2003). PAR4 also contains an anionic cluster, Asp<sup>57</sup>...Asp<sup>59</sup>...Glu<sup>62</sup>...Asp<sup>65</sup>, just C-terminal of the thrombin cleavage site that is thought to interact with cationic residues that border exosite I of thrombin (Ayala *et al.*, 2001) and slow the dissociation rate of thrombin from PAR4 (Jacques and Kuliopulos, 2003). This is reflected in more sustained intracellular signals elicited by PAR4 activation

compared with the more transient signals in response to PAR1 activation (Covic *et al.*, 2002c). These differences in receptor structure and activation mechanisms between PAR1 and PAR4 indicate that these two receptors have the potential to initiate distinct intracellular signalling kinetics and/or events.

Receptor trafficking mechanisms are another key regulator of receptor signalling. This is of particular importance for PARs due to the irreversible, cleavage-based, endogenous activation of these receptors, with continued responsiveness to agonists requiring new receptors to be trafficked to the cell surface. While much is known regarding the receptor trafficking and response regulation of PAR1 and PAR2 (for a detailed review, see Soh *et al.*, 2010), little is known about such mechanisms for PAR4. Defining these pathways is of clear interest. On this point, one apparently important distinction between PAR4 and PARs 1 and 2 relates to receptor desensitization. After receptor cleavage and activation, PARs are rapidly internalized on a phosphorylation signal in regions of the C-terminus of the receptor (Shapiro *et al.*, 1996). However, PAR4 has a shorter C-terminus than either of PAR1 or PAR2 and does not have many of the phosphorylation sites shown to be necessary for desensitization of these receptors (Shapiro



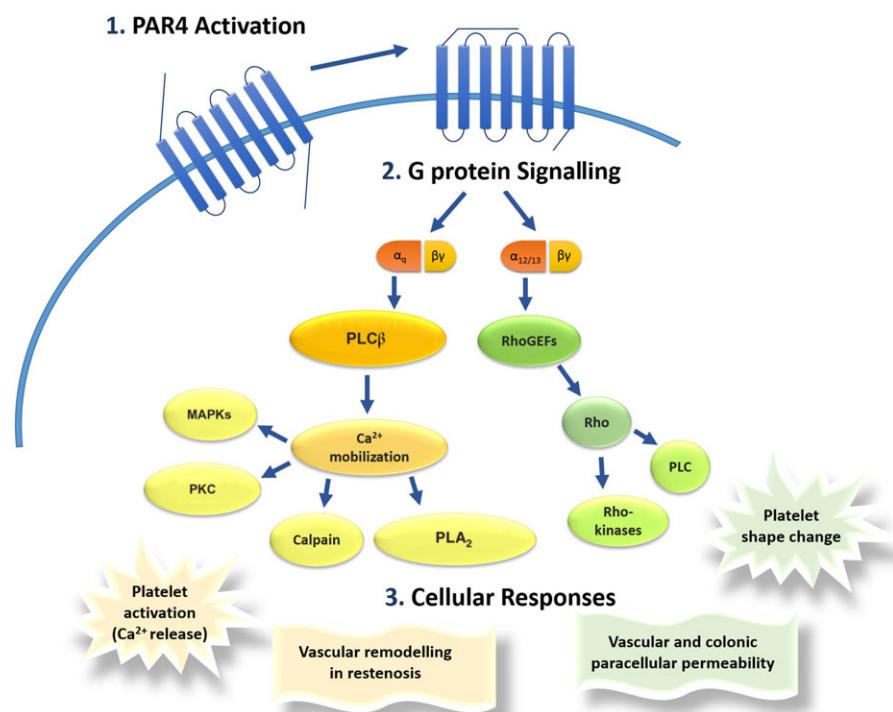
*et al.*, 2000). As a result, agonist-triggered phosphorylation and consequent receptor internalization is significantly slower for PAR4 than for PAR1 or PAR2 (Shapiro *et al.*, 2000), providing a further mechanism for the comparatively prolonged intracellular signalling downstream of PAR4 activation.

Following internalization, PARs are sorted for either degradation (lysosomes) or recycling (endosomes). While activated receptors are predominately packaged into lysosomes for complete signal termination, unactivated receptors can cycle tonically between endosomes and the cell surface to ensure ongoing sensitivity to agonists in the absence of *de novo* receptor synthesis (Hoxie *et al.*, 1993; Shapiro *et al.*, 1996). An additional membrane trafficking pathway for PAR4 is provided by efficient transport from the endoplasmic reticulum (ER), via the Golgi for post-translational modifications such as glycosylation. PAR4 was recently shown to contain an arginine-based ER retention motif within the second intracellular loop that may regulate cell surface expression of the receptor (Cunningham *et al.*, 2012). Intriguingly, co-expression of PAR2 appeared to interfere with this ER retention signal and enhance PAR4 expression on the surface of a human keratinocyte cell line (Cunningham *et al.*, 2012), providing a potential mechanism for cell type and/or context-dependent discrepancies in the regulation of PAR4 surface expression.

## Intracellular signalling

Cleavage and activation of PARs prompts a conformational change in the receptor that allows G protein coupling and initiates multiple intracellular signalling events.  $G_q$ ,  $G_i$  and  $G_{12/13}$  have all been demonstrated to mediate signals in response to activation of PARs, with PAR4 specifically being shown to couple to both  $G_q$  and  $G_{12/13}$  family proteins (Coughlin, 2000; Faruqi *et al.*, 2000), but not  $G_i/G_o$  (Voss *et al.*, 2007) (Figure 2).

PAR4 coupling to  $G_{12/13}$  initiates binding of RhoGEFs (guanine-nucleotide exchange factors that activate Rho) to the  $\alpha$ -subunit. Consequent Rho activation induces a series of Rho-dependent cytoskeletal responses and PLC activation (Figure 2). For example, in platelets, activation of Rho-kinase-dependent cytoskeletal responses via  $G_{12/13}$ -mediated PAR4 signalling triggers platelet shape change (Offermanns *et al.*, 1994; Coughlin, 2000; Dorsam *et al.*, 2002) – one of the key initial events in platelet aggregation during thrombosis. In endothelial cells,  $G_{12/13}$ -mediated PAR4 signalling via Rho is manifested in cytoskeletal responses that drive increases in vascular and colonic paracellular permeability (Dabek *et al.*, 2009; Dabek *et al.*, 2011). Evidence also exists for PAR4 activation eliciting a prolonged signal via  $G_{12/13}$ , which is thought to involve the regulation of MLC phosphorylation and RhoA (Dabek *et al.*, 2011; Duvernay *et al.*, 2013).



**Figure 2**

Protease-activated receptor 4: signalling. Following cleavage and activation of the receptor, PAR4 signals via coupling to  $G_q$  and/or  $G_{12/13}$  family members. The best characterized signalling downstream of  $G_q$  is via  $PLC\beta$ -mediated phosphoinositide hydrolysis and resultant intracellular calcium mobilization. This promotes the activity of several calcium-regulated kinases and phosphatases (MAPKs, PKCs,  $PLA_2$  and calpain) which underlie PAR4-induced cellular responses such as platelet activation and vascular remodelling. The best characterized signalling downstream of PAR4-  $G_{12/13}$  coupling is via RhoGEF and consequent Rho activation, which underlie PAR4-induced cellular responses such as cytoskeletal responses in platelets and vascular smooth muscle and endothelial cells.

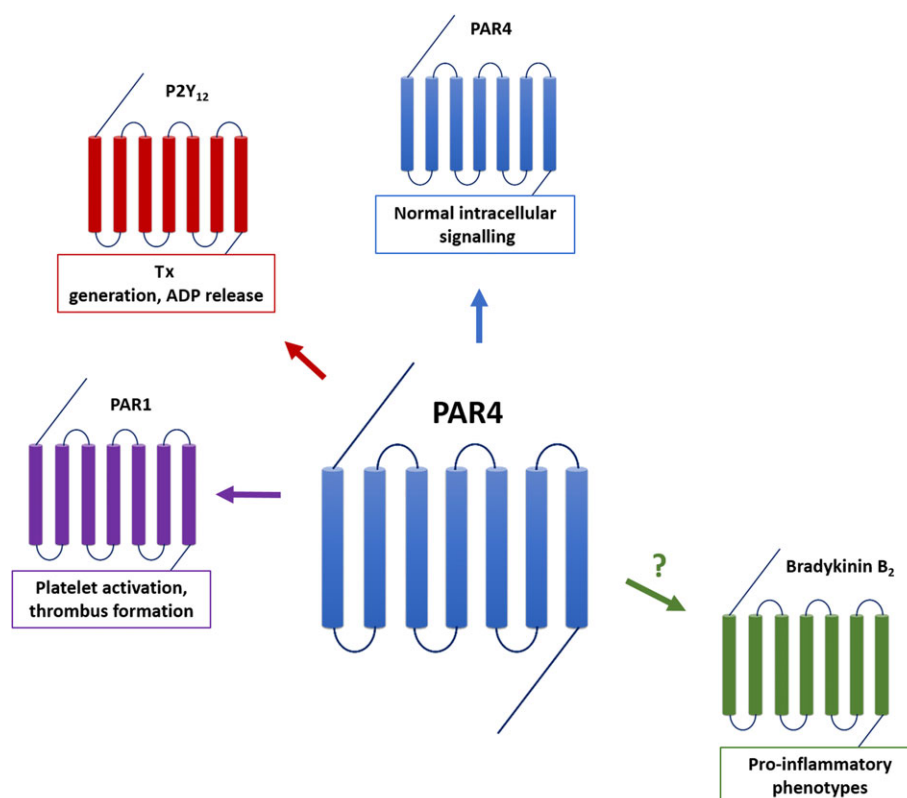
PAR4 coupling to  $G_q$  causes intracellular calcium mobilization via activation of  $PLC\beta$  and consequent phosphoinositide hydrolysis (Figure 2). PAR4-induced calcium mobilization promotes the activity of several calcium-regulated kinases and phosphatases (e.g. MAPKs, PKCs,  $PLA_2$  and calpain) (Coughlin, 2000), with wide-ranging effects. For example, PAR4 activation mediates several key platelet responses that are induced by  $G_q$ -dependent calcium mobilization, including the secretion of platelet storage granules and integrin activation – both of which are critical for effective platelet aggregation (Coughlin, 2000; Coughlin, 2005).

As indicated previously, one key difference with PAR4 is in the kinetics of receptor activation and signalling. Regarding  $G_q$ -mediated signalling, it is known that PAR1 activation stimulates a rapid burst of intracellular calcium mobilization, whereas PAR4 activation elicits a slower rise, which is much more sustained over time (Covic *et al.*, 2002c). This is likely to be a result of the slower cleavage of PAR4 allowing prolonged G protein signalling. Given that mice deficient in  $G_q$  lack thrombin-dependent  $IP_3$  and calcium responses (Gabbeta *et al.*, 1997; Offermanns *et al.*, 1997) and that PAR4 activation stimulates ongoing  $IP_3$  and DAG, differences in calcium mobilization are likely to reflect a result of prolonged  $G_q$  signalling through PAR4 (Lau *et al.*, 1994; Covic *et al.*, 2002c). The potential outcome of this difference in signalling kinetics

is discussed in more detail below, but may be important for ongoing cell signalling under conditions of prolonged agonist exposure. For example, in platelets, PAR1-mediated calcium signalling undergoes rapid desensitization but can be rescued by subsequent PAR4 activation (Falker *et al.*, 2011). In endothelial and smooth muscle cells, long-term  $G_q$ -dependent calcium signalling via PAR4 drives transcriptional and metabolic responses that may be important in vascular remodelling and proliferative processes in restenosis in response to high glucose levels (Pavic *et al.*, 2014).

## Dimerization and crosstalk with other receptors

PAR4 has been shown to form both heterodimers and homodimers (Leger *et al.*, 2006; Arachiche *et al.*, 2013) (Figure 3). Specifically, PAR4 forms stable heterodimers with PAR1 and preventing this association impairs signalling downstream of either receptor (Leger *et al.*, 2006), suggesting reciprocal assistance in the cleavage and activation of PAR1 and PAR4. For example, pharmacological disruption of the PAR1–PAR4 heterodimer effectively prevents carotid artery occlusion in an *in vivo* murine thrombosis model (Leger *et al.*,



**Figure 3**

Protease-activated receptor 4: dimerization partners. There is direct evidence in transfected cell systems for PAR4 homodimers, as well as heterodimers with PAR1 and the P2Y<sub>12</sub> purinergic receptor. Each of these interactions has been shown to be important for normal intracellular signalling and have been implicated in platelet aggregation and thrombus formation. In addition, crosstalk between PAR4 and the B<sub>2</sub> bradykinin receptor may mediate pro-inflammatory effects, either directly or indirectly via transactivation.

2006). In addition, although direct evidence for PAR3–PAR4 heterodimers has not been reported, PAR3, at least in mouse systems, serves as a cofactor for efficient PAR4 activation (Kahn *et al.*, 1998b; Nakanishi-Matsui *et al.*, 2000), suggesting a significant level of interaction between these two PARs. PAR4 has also recently been shown to form homodimers (De La Fuente *et al.*, 2012), with the interacting region mapped to transmembrane helix 4 of the receptor. PAR4 homodimers also appear vital for normal signalling: mutation of a series of hydrophobic residues in transmembrane domain 4 reduced PAR4-mediated calcium mobilization in transfected HEK293 cells (De La Fuente *et al.*, 2012).

Other GPCRs reported to form heterodimers with PAR4 include the P2Y<sub>12</sub> purinergic receptor (Khan *et al.*, 2014), and functional co-operativity – potentially via receptor dimerization – has been reported with the  $\alpha_{2A}$ -adrenoceptor (Grenegard *et al.*, 2008) and B<sub>2</sub> bradykinin receptor (Houle *et al.*, 2005; McDougall *et al.*, 2009) (Figure 3). In human platelets, PAR4 (along with PAR1) is known to work in synergy with the P2Y<sub>12</sub> receptor to contribute to platelet activation and aggregation events, including thromboxane (TXA<sub>2</sub>) generation and ADP release (Holinstat *et al.*, 2006). It has been recently shown that this process is largely mediated by PAR4 and that PAR4 directly interacts with P2Y<sub>12</sub> receptors (Wu *et al.*, 2010; Khan *et al.*, 2014). PAR4 also interacts with the  $\alpha_{2A}$ -adrenoceptor (Grenegard *et al.*, 2008). This interaction has been shown to cause aggregation of aspirin-treated human platelets, via cooperative signalling which eventually triggers the P2X<sub>1</sub> ATP-gated calcium ion channel to overcome the therapeutic effects of aspirin. It has also been suggested that PAR4 communicates with the bradykinin B<sub>2</sub> receptor, responsible for several pro-inflammatory effects. Although the mechanisms and outcomes of this crosstalk are yet to be fully elucidated, it has been shown in models of rat paw oedema and knee joint inflammation that inflammation induced by a PAR4 activating peptide (PAR4-AP) can be inhibited by a B<sub>2</sub> receptor antagonist (Houle *et al.*, 2005; McDougall *et al.*, 2009). These various examples of receptor dimerization and/or crosstalk are indicative of the complexity of PAR4 signalling and how these events are able to affect multiple pathways to alter physiological responses.

## Physiology

### Tissue distribution and expression

PAR4 is broadly expressed, with gene expression by Northern blot most readily detected in the lungs, pancreas, thyroid, testis and small intestine (Xu *et al.*, 1998; Kahn *et al.*, 1998b). Moderate expression has also been detected in placenta, skeletal muscle, lymph nodes, adrenal gland, prostate, uterus and colon (Xu *et al.*, 1998; Kahn *et al.*, 1998b). In the nervous system, PAR4 protein and mRNA have been detected in rat dorsal root ganglion (DRG) non-neuronal cells (conversely, PARs 1–3 were detected in DRG neurons) (Zhu *et al.*, 2005). Despite this extensive expression profile of PAR4, most research has focused on the physiological functions of this receptor in cardiovascular and inflammatory settings. In humans, the key vascular cell types expressing PAR4 include platelets (Xu *et al.*, 1998; Kahn *et al.*, 1998b), leukocytes

(Vergnolle *et al.*, 2002), endothelial cells (Kataoka *et al.*, 2003) and smooth muscle cells (Hollenberg *et al.*, 1999; Hamilton *et al.*, 2001; Vidwan *et al.*, 2010). In addition to the well-known pro-thrombotic and pro-inflammatory actions of PAR4-activating proteases, perhaps one reason for this is that several important animal models, including mice, rats, guinea pigs, rabbits, dogs and monkeys have similar PAR4 expression to humans throughout the vasculature. Platelets are one example of such conserved PAR4 expression – even in the face of variable expression of the other PARs: human platelets express PAR1 and PAR4 (Kahn *et al.*, 1998b); platelets from mice, rats and rabbits express PAR3 and PAR4 (Connolly *et al.*, 1994; Connolly *et al.*, 1996; Nakanishi-Matsui *et al.*, 2000; Khan *et al.*, 2005), while guinea pig platelets express PAR1, PAR3 and PAR4 (Andrade-Gordon *et al.*, 2001). Non-human primates are the only known animal with an identical platelet-PAR expression profile to that of humans (Derian *et al.*, 2003). Not surprisingly, this has been a significant restraint in the pursuit of PAR antagonists as anti-platelet agents. Despite this relatively conserved expression of PAR4 in vascular cells across species, there is some evidence to suggest that PAR4 may function differently between species. Specifically, PAR4 contributes to thrombin responses of mouse endothelial cells (Kataoka *et al.*, 2003) that PAR4 activation causes endothelium-dependent relaxation of rat aorta (Hollenberg *et al.*, 1999). However, in human artery preparations as well as lung fibroblasts, these responses are only elicited when treated with additional inflammatory mediators (Hamilton *et al.*, 2001; Ramachandran *et al.*, 2007). The consequences of this distinct regulation of PAR4 expression by inflammatory cytokines remain unknown but are likely to be important for future studies investigating the roles of PAR4 in inflammation.

### Key physiological functions

PAR4 knockout (PAR4<sup>−/−</sup>) mice have been used to elucidate physiological functions dependent on PAR4 expression (Table 1). Sambrano *et al.* (2001) generated the first PAR4<sup>−/−</sup> mice, in which  $\beta$ -galactosidase was expressed in place of PAR4. Loss of PAR4 function was confirmed by platelet aggregation studies where it was shown that PAR4<sup>−/−</sup> platelets were completely unresponsive to both a PAR4-AP (AYPGKF) and thrombin (Sambrano *et al.*, 2001). PAR4<sup>−/−</sup> mice are normal in appearance, size and fertility, with no platelet, leukocyte or erythrocyte abnormalities (Sambrano *et al.*, 2001). Indeed, no spontaneous phenotype has been reported in PAR4<sup>−/−</sup> mice to date.

**Protection against thrombosis.** The first phenotype reported in PAR4<sup>−/−</sup> mice was protection against thrombosis, with mild associated bleeding (Sambrano *et al.*, 2001). Specifically, it was shown that platelets from PAR4<sup>−/−</sup> mice were completely unresponsive to thrombin, thus creating a clean genetic model in which to test for the overall importance of thrombin-mediated platelet activation for thrombosis *in vivo* (Sambrano *et al.*, 2001). Initially, ferric chloride-induced injury of mouse mesenteric arterioles was utilized. Here, the time to vessel occlusion after injury was prolonged about three-fold in PAR4<sup>−/−</sup> mice (Sambrano *et al.*, 2001). Subsequent studies have shown a similar protection against thromboplastin-

**Table 1**Reported phenotypes in PAR4-deficient (PAR4<sup>-/-</sup>) mice

Phenotype		References
Thrombosis and haemostasis	PAR4 <sup>-/-</sup> mice are protected against arterial/arteriolar thrombosis.	Sambrano <i>et al.</i> , 2001; Weiss <i>et al.</i> , 2002; Lee <i>et al.</i> , 2012a, 2012b
	PAR4 <sup>-/-</sup> mice are protected against pulmonary embolism.	Hamilton <i>et al.</i> , 2004
	PAR4 <sup>-/-</sup> mice display increased tail bleeding times.	Sambrano <i>et al.</i> , 2001; Weiss <i>et al.</i> , 2002; Hamilton <i>et al.</i> , 2004; Lee <i>et al.</i> , 2012b
Inflammation and vascular protection	PAR4 <sup>-/-</sup> mice are protected from cerebral infarct after cerebral artery occlusion.	Mao <i>et al.</i> , 2010
	PAR4 <sup>-/-</sup> mice are protected from tissue oedema.	Busso <i>et al.</i> , 2008
	PAR4 <sup>-/-</sup> mice have diminished vascular remodelling in the setting of diabetes.	Pavic <i>et al.</i> , 2014
Hepatotoxicity	PAR4 <sup>-/-</sup> mice are protected against acetaminophen-induced liver injury.	Miyakawa <i>et al.</i> , 2015

induced pulmonary embolism (Weiss *et al.*, 2002), laser-induced endothelial cell ablation in mesenteric arterioles (Vandendries *et al.*, 2007) and electrolytic-induced (Lee *et al.*, 2012a) and trauma-induced (Lee *et al.*, 2012b) injury of the carotid artery in PAR4<sup>-/-</sup> mice. Bone marrow transplantation studies confirmed that the anti-thrombotic effects observed in PAR4<sup>-/-</sup> mice were due to deficiency of PAR4 in platelets (Hamilton *et al.*, 2004). Intriguingly, it appears that the protection against thrombosis associated with PAR4 deficiency leaves other haemostatic responses intact (Vandendries *et al.*, 2007). PAR4 deficiency in mice is also associated with a mild bleeding phenotype. This haemostatic effect has been most commonly assessed via tail bleeding time, where PAR4<sup>-/-</sup> mice are consistently shown to have prolonged tail bleeding times compared with wild-type mice (Sambrano *et al.*, 2001; Weiss *et al.*, 2002). Despite this commonly reported effect upon active limb trauma, there is no evidence for spontaneous bleeding in PAR4<sup>-/-</sup> mice. PAR4 deficiency also has protective benefits in a mouse model of stroke (transient middle cerebral artery occlusion, in which PAR4<sup>-/-</sup> mice exhibited lower cerebral infarct volume, improved neurological and motor function and reduced blood brain barrier disruption and cerebral oedema, compared with wild-type animals (Mao *et al.*, 2010). Additional models of atherosclerotic plaque formation in ApoE<sup>-/-</sup> have indicated that PAR4 probably does not play a protective role in this setting (Hamilton *et al.*, 2009). However, diabetic mice also deficient in PAR4 exhibit protective vascular remodelling phenotypes (Pavic *et al.*, 2014), indicating that PAR4 may be functionally important in certain diabetic vasculopathies.

**Inflammation.** PAR4 has also been linked to inflammation and neuropathic pain pathways. This is perhaps not surprising given that serine proteases capable of activating PAR4 (i.e. thrombin from coagulation activation, cathepsin G released from neutrophils and trypsin released from mast cells) are well known to drive many pro-inflammatory effects. Most studies investigating the role of PAR4 in inflammatory settings have

examined gain of function phenotypes, most commonly in response to exogenous PAR4 agonists. One of the earliest studies implicating PAR4 in pro-inflammatory reactions found that PAR4 activation mimics the effects of thrombin on leukocyte rolling and adhesion and increases leukocyte migration (Vergnolle *et al.*, 2002). Specifically, superfusion of rat mesenteric venules with either thrombin or a PAR4-AP (but not a PAR1-AP), significantly increased the number of rolling leukocytes, as well as leukocyte recruitment to the peritoneal cavity, suggesting that thrombin-induced effects on leukocytes during inflammation may be mediated largely by PAR4 (Vergnolle *et al.*, 2002). In subsequent studies, PAR4 activation mediated the formation of oedema in rat paws (Hollenberg *et al.*, 2004; Houle *et al.*, 2005; McDougall *et al.*, 2009) via neutrophils and the kallikrein-kinin system (Houle *et al.*, 2005). In support of these observations, PAR4<sup>-/-</sup> mice develop significantly reduced paw swelling in response to tissue-factor-initiated inflammation (Busso *et al.*, 2008).

**Nociception.** PAR4 expression is up-regulated in DRG neurons during nociceptor activation and neurogenic pain (Dattilio *et al.*, 2005; Chen *et al.*, 2013) and in sensory neurons of the colon (Auge *et al.*, 2009), suggesting a potential role for PAR4 in nociception associated with inflammatory pathologies of the viscera, such as inflammatory bowel disease. In addition, PAR4 may be important in transmitting the signals for neurogenic itch (Tsujii *et al.*, 2008; Papoiu *et al.*, 2015; Patricio *et al.*, 2015). PAR4 activation is thought to attenuate nociception via sensory neurons by inhibiting nociceptive signals (Asfaha *et al.*, 2007). These analgesic effects of PAR4 activation have been modelled *in vivo*, where injections of PAR4-agonist peptides in rat paws evoke analgesia in response to both mechanical and thermal stimuli (Asfaha *et al.*, 2007). Further, in the setting of inflammatory bowel disease in mice, colonic delivery of PAR4 agonists reduced basal visceral pain and visceral hypersensitivity caused by PAR2 (Auge *et al.*, 2009). For a detailed review on PAR4 in pain and inflammatory pathologies, see Fu *et al.* (2015). Together, these studies help identify PAR4 as a likely mediator of



multiple pathological processes and provide rationale for the development of pharmacological reagents targeting PAR4.

## Pharmacology

### Agonists

Current agonists of PAR4 consist of amidated peptides based on the native tethered ligand sequence of the receptor (Table 2). A hexapeptide matching the sequence immediately downstream of the thrombin cleavage site of the receptor, GYPGQV-NH<sub>2</sub> (Figure 1) selectively activates PAR4, although prohibitively high concentrations of ~500  $\mu$ M are required for effective activation (Faruqi *et al.*, 2000). The native tethered ligand sequence of mouse PAR4, GYPGKF-NH<sub>2</sub>, activates both human and mouse PAR4 with slightly greater potency (Kahn *et al.*, 1998b). However, an alanine scan of this sequence undertaken to identify residues critical for receptor activation inadvertently uncovered a more potent peptide agonist, AYPGKF-NH<sub>2</sub>, which has become the gold standard PAR4 activator. This peptide has an EC<sub>50</sub> of between 5 and 100  $\mu$ M, depending on the setting (Table 2) and is highly specific for PAR4 (Faruqi *et al.*, 2000; Hollenberg *et al.*, 2004).

### Antagonists

Research into the physiological function of PAR4 relies on the development of specific and effective antagonists. The unique structure and activation mechanism of PARs has long posed a problem for the development of such inhibitors. In particular, activation by proteolytic cleavage is a highly efficient and irreversible system, while the tethered ligand-based activation mechanism requires antagonism of the binding of an agonist intrinsic to the receptor and presumably with considerable steric advantage. These issues have hindered the development of PAR4 antagonists, although recent efforts have begun to overcome these challenges, with several distinct antagonist classes emerging (Table 2).

**Peptidomimetics.** The first approach taken to identify potential inhibitors of PAR4 was the generation of numerous peptide analogues based on the tethered ligand sequences of human, mouse and rat PAR4. The peptidomimetic approach yielded two sequences based on mouse PAR4 and modified by the addition of a *trans*-cinnamoyl (tc) group (Hollenberg and Saifeddine, 2001). The resulting compound, tc-YPGKF-NH<sub>2</sub>, appeared to bind but not activate PAR4 and abolished PAR4-AP-induced aggregation of rat platelets and significantly reduced thrombin-induced platelet aggregation, at least at high concentrations (400  $\mu$ M) (Hollenberg and Saifeddine, 2001). tc-YPGKF-NH<sub>2</sub> also inhibited thrombin-induced platelet aggregation in human platelets (Ma *et al.*, 2005), although there is limited other evidence of this agent inhibiting human PAR4 (Hollenberg *et al.*, 2004).

**Low MW compounds.** Low MW PAR4 antagonists have also been developed via screening of various heterocyclic structures. The first compound identified to selectively inhibit PAR4 was an indazole derivative known as YD-3 [1-benzyl-3

(ethoxycarbonylphenyl)-indazole] (Lee *et al.*, 2001; Wu *et al.*, 2002). YD-3 inhibited thrombin-induced platelet aggregation in rabbits with an IC<sub>50</sub> of 28  $\mu$ M but, in humans, it only partly inhibited platelet aggregation in response to thrombin concentrations lower than 0.5 nM (Wu *et al.*, 2002, 2003). Several studies have aimed to increase the efficacy of YD-3, and have produced several derivatives (Huang *et al.*, 2006; Chen *et al.*, 2008), including the N2-(substituted benzyl)-3-(4-methylphenyl)-2H-indazoles, compounds numbered 19, 25, and 31 (Huang *et al.*, 2006). The IC<sub>50</sub> of these compounds in HL-60 cell lines was reported to be greater than 36  $\mu$ M (Huang *et al.*, 2006). The newest low MW PAR4 inhibitor, ML354, is a substituted indole-based derivative (Young *et al.*, 2013; Wen *et al.*, 2014). This compound inhibited PAR4-AP-induced integrin activation with an IC<sub>50</sub> of 140 nM, but also PAR1-AP-induced integrin activation with an IC<sub>50</sub> of 10  $\mu$ M. Initial studies using these recently developed low MW PAR4 antagonists have helped to generate interest in PAR4 as a novel therapeutic target for anti-angiogenic drugs (Huang *et al.*, 2006). A clinical candidate for PAR4 inhibition, BMS-986120, is currently being evaluated in phase I clinical trials for safety and tolerability for the prevention and/or treatment of thromboembolic disorders (NCT02208882).

**Pepducins.** Pepducins are a distinct class of PAR4 inhibitors that mimic the region of the receptor that binds G proteins and essentially work by blocking the interactions between the receptor and effector G protein (Covic *et al.*, 2002a). Specifically, peptides with a sequence corresponding to the third intracellular loop of a given receptor (the region responsible for interaction with G proteins) are conjugated to an N-terminal palmitate (pal) that anchors the peptide to the lipid membrane of a cell. Once anchored within the membrane, it is proposed that pepducins act by binding G proteins and thus prevent downstream signalling. PARs were used as the target receptors for initial proof-of-concept studies, and pepducins exist against both PAR1 and PAR4 (Covic *et al.*, 2002a). The anti-PAR4 pepducin, P4pal-10 (pal-SGRRYGHALR-NH<sub>2</sub>) inhibited ~85% of thrombin-induced aggregation of both human and mouse platelets at a concentration of 3  $\mu$ M and is effective *in vivo*, as assessed by increased tail bleeding time in mice (Covic *et al.*, 2002a, 2002b). However, the specificity of P4pal-10 remains debatable, as it displays a level of cross-reactivity with PAR1-AP-induced platelet activation (Covic *et al.*, 2002b), as well as collagen and TXA<sub>2</sub> (Stampfuss *et al.*, 2003). A subsequent pepducin (P4pal-i1) targeted against the first intracellular loop (i1) was synthesized to address the issue of specificity and has been shown to inhibit PAR4-AP-induced platelet aggregation without affecting PAR1-AP-induced platelet aggregation (Leger *et al.*, 2006).

**Function-blocking antibodies.** Several function-blocking anti-PAR4 antibodies have also been developed and proven to be useful PAR4 antagonists. The first anti-PAR4 function-blocking antibody was a rabbit polyclonal antibody against the thrombin cleavage site of human PAR4. This antibody inhibited thrombin-induced aggregation of human platelets in the presence of concomitant PAR1 inhibition at a concentration of 1 mg·mL<sup>-1</sup> (Kahn *et al.*, 1999). More recently, a strategy was employed by Mumaw *et al.* (2014) to

**Table 2**  
PAR4 agonists and antagonists

Class	Compound	Target	Agonists		Reference
			Receptor	Cell	
Amidated peptides	GYPGQV-NH <sub>2</sub>	TLS binding site	Human	Transfected mouse fibroblasts	Faruqi <i>et al.</i> , 2000
	GYPGKF-NH <sub>2</sub>	TLS binding site	Human	Transfected mouse fibroblasts	Kahn <i>et al.</i> , 1999
			Rat	Platelets	Hollenberg <i>et al.</i> , 2004
	AYPGKF-NH <sub>2</sub>	TLS binding site	Human	Transfected mouse fibroblasts	Faruqi <i>et al.</i> , 2000
			Rat	Platelets	Hollenberg <i>et al.</i> , 2004
			Rat	Aorta	
Class	Compound	Target	Antagonists		Reference
			Receptor	Cell type	
Peptidomimetics	tc-YPGKF-NH <sub>2</sub>	TLS binding site	Rat, human	Platelets	Hollenberg <i>et al.</i> , 2001; Hollenberg <i>et al.</i> , 2004; Ma <i>et al.</i> , 2005
Low MW compounds	YD-3	TLS binding site	Rabbit, rat, human	Platelets, smooth muscle cells	Wu <i>et al.</i> , 2002; Wu <i>et al.</i> , 2003; Peng <i>et al.</i> , 2004
	19, 25, 31	TLS binding site	Human	Platelets	Huang <i>et al.</i> , 2006
	ML354	TLS binding site	Human	Platelets	Young <i>et al.</i> , 2013; Wen <i>et al.</i> , 2014
Pepducins	P4pal-10	Third intracellular loop	Mouse, human	Platelets	Covic <i>et al.</i> , 2002a, b
	P4pal-11	First intracellular loop	Guinea pigs, human	Platelets	Leger <i>et al.</i> , 2006
Function-blocking antibodies	Rabbit polyclonal	Thrombin cleavage site	Human, rat	Platelets, fibroblasts	Kahn <i>et al.</i> , 1999
	Rabbit polyclonal (Can12)	Anionic region	Mouse, human	Platelets	Mumwaw <i>et al.</i> , 2014
	Mouse monoclonal (14H6, 5 F10)	Thrombin cleavage site	Human	Platelets, HEK-293s	Mumwaw <i>et al.</i> , 2015

<sup>a</sup>IC<sub>50</sub> values correspond to inhibition of thrombin-induced PAR4 activation.

<sup>b</sup>IC<sub>50</sub> values correspond to inhibition of PAR4 agonist peptide-induced activation.

<sup>c</sup>TLS, tethered ligand sequence; Values are concentrations reported to inhibit thrombin-induced PAR4 activation.

raise anti-PAR4 antibodies against the anionic region sequence of PAR4 (C<sup>54</sup>ANDSDTLTPD), just downstream of the thrombin cleavage site. The lead antibody to emerge from these studies, CAN12, not only inhibits thrombin-induced aggregation of human platelets but also inhibits aggregation induced by PAR4-AP, ADP or collagen (Mumaw *et al.*, 2014), suggesting major issues with specificity. CAN12 was shown to inhibit arterial thrombosis in the Rose Bengal carotid artery thrombosis model at 0.5 mg.kg<sup>-1</sup> (Mumaw *et al.*, 2014), indicating cross-reactivity with murine PAR4. Beneficial effects were seen when CAN12 was infused both 10 min pre-injury and 15 min post-injury. A new series of monoclonal antibodies from the same group has been very recently reported and appear to bind either to the anionic region or to the thrombin cleavage site of PAR4 and partially inhibit thrombin-induced cleavage of the receptor in expression systems (Mumaw *et al.*, 2015).

This diverse set of PAR4 antagonists, developed over recent years, have been invaluable for investigations into the physiological roles of PAR4 and have also served to promote targeting of this receptor as a novel therapeutic approach. However, there still remains significant work to be done to develop PAR4 antagonists suitable for potential clinical utility.

## Clinical significance

Cardiovascular disease remains a major clinical problem. Increasingly prevalent conditions such as diabetes, hypertension and coronary artery disease eventuate in diseased and pro-thrombotic vasculature and a common endpoint of these, often co-existing, conditions is the formation of pathological arterial thrombosis leading to myocardial infarction or ischaemic stroke. Together, such conditions account for ~40% of all deaths in Western countries (Mozaffarian *et al.*, 2015). Given that platelets are the key cellular component of arterial thrombi, anti-platelet agents are the primary therapy for the prevention of myocardial infarction and ischaemic stroke. There are several anti-platelet drugs on the market, but a lack of efficacy and safety of these existing drugs has fuelled a need for improved therapies. Given the importance of PARs for platelet activation, PAR antagonists represent one of the leading classes of new anti-platelet drugs.

### Targeting PAR4 as an anti-platelet approach?

Ever since the discovery of PAR1 as the major platelet thrombin receptor, there was strong interest in developing PAR1 antagonists as novel anti-platelet agents. This led to the development of the first PAR1 antagonist, vorapaxar (trade name Zontivity), which was recently granted FDA approval for use in the USA for the prevention of myocardial infarction and peripheral artery disease. Vorapaxar was assessed in two large-scale phase III clinical trials, of which one was terminated early due to a significant increase in intracranial haemorrhage in the vorapaxar-treated cohort (Scirica *et al.*, 2012; Tricoci *et al.*, 2012). Indeed, throughout these trials, vorapaxar was associated with reduction of ischaemic outcomes in some subgroups, whereas in others was associated

with adverse bleeding events (French *et al.*, 2015). Subsequent studies examining blood from patients receiving vorapaxar have confirmed that PAR4-mediated platelet responses remain intact (Judge *et al.*, 2014). As a result, there is growing interest in the clinical potential of PAR4 antagonists as novel anti-platelet agents.

### Clinically significant variants of PAR4

Further support for the targeting of PAR4 as an anti-platelet approach comes from recent studies demonstrating racial differences in the expression and reactivity of platelet PAR4 (Edelstein *et al.*, 2013; Edelstein *et al.*, 2014; Tourdot *et al.*, 2014). Significantly, these differences appear to associate with resistance to current anti-platelet drugs and overall poorer cardiovascular outcomes (Edelstein *et al.*, 2013; Edelstein *et al.*, 2014; Tourdot *et al.*, 2014). Specifically, platelets from healthy Black subjects were hyper-responsive to PAR4-AP when compared with platelets from non-Black subjects. This increased sensitivity persisted in the face of PAR1 antagonism (vorapaxar) or aspirin treatment. This work led to the identification of a genetic variant of PAR4, Ala<sup>120</sup>Thr, which is associated with greater PAR4 reactivity and occurs at a higher frequency in Black subjects than in non-Black subjects. This racial difference may have important clinical implications and potential therapeutic impact on specific population responses, and therefore clinical outcomes, to anti-platelet therapy, and further supports the idea of targeting platelet PAR4 for improved anti-thrombotic therapy.

## Conclusions

Despite the discovery of PAR4 over 15 years ago, comparatively little is known about the function of this receptor. Generation and characterization of PAR4<sup>-/-</sup> mice has yielded phenotypes relating to thrombosis and haemostasis, as well as inflammation and vascular protection – prompting recent interest in the field and leading to the development of improved inhibitors for investigations into PAR4 function. This important ground work now allows several key remaining questions to be addressed. For instance, apart from thrombosis, inflammation, and pain, what physiological functions are there for PAR4? How can PAR4 be best inhibited pharmacologically? What is the importance of PAR4 expression and reactivity differences observed in genetically distinct patient groups and how will this affect treatment options for people with cardiovascular and/or inflammatory diseases? In addition, the limited number of molecular pharmacology studies of PAR4 leaves several other unanswered questions: how important is receptor dimerization for specific cell signalling events? Does the importance of PAR3/PAR4 co-factoring in mouse platelets extend to other receptors in other settings? Does biased signalling occur for PAR4, as it does for both PAR1 and PAR2? The recent increase in interest in PAR4 will, we hope, culminate in the development of improved pharmacological reagents and further molecular pharmacology experiments that will provide the tools to address these questions in the coming years.

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## Conflict of interest

The authors declare no conflicts of interest.

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## **Appendix III**

# Drugs targeting protease-activated receptor-4 improve the anti-thrombotic therapeutic window

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## Introduction

Antiplatelet agents are the main pharmacotherapy for arterial thrombosis prevention and are central in the management of cardiovascular conditions such as myocardial infarction, transient ischaemic attack, and coronary and peripheral artery diseases. Yet despite their long history and extensive clinical use, antiplatelet agents appear to have reached a disappointingly low therapeutic ceiling—predominantly due to the narrow therapeutic window afforded by strategies targeting platelet function. Platelets are critical for normal hemostasis as well as pathological thrombosis. Inhibiting platelet function for protective benefit without causing unwanted bleeding limits the efficacy of current antiplatelet drugs. Aspirin and the thienopyridine class of drugs (e.g., clopidogrel, prasugrel, ticagrelor) are by far the most commonly prescribed antiplatelet agents, yet prevent just 15 and 17% of lethal cardiovascular events respectively (1,2). Combination therapy provides a marginal increase in efficacy (~7%), but also increases the risk of bleeding (2). More potent antiplatelet drugs, such as the glycoprotein  $\alpha_{IIb}\beta_3$  inhibitors, carry even more bleeding risk and are thereby limited to acute use settings such as periprocedural percutaneous coronary intervention (3,4). Therefore, the search to identify antiplatelet drugs that increase the therapeutic window of antithrombotic therapy continues. A recent study by Wong *et al.* (5) provides compelling evidence that targeting the platelet thrombin receptor, PAR4, may achieve

this goal.

In the setting of thrombosis, platelets are activated by a combination of endogenous agonists, some of which are blocked by existing drugs. For example, aspirin prevents production of the platelet activator thromboxane  $A_2$  while the thienopyridines block the major platelet ADP receptor, P2Y<sub>12</sub>. Thrombin is the most potent platelet activator, which it achieves predominantly via two cell surface GPCRs, protease-activated receptor 1 (PAR1) and PAR4. PAR1 has greater affinity for thrombin than PAR4 and has therefore been the focus of drug development targeting thrombin-induced platelet activation. The first PAR1 antagonist, vorapaxar, was approved by the US FDA in 2014 for the prevention of thrombotic events in patients with a history of myocardial infarction or peripheral artery disease. Yet because it must be administered in addition to standard-of-care antiplatelet therapy (aspirin and/or a thienopyridine), vorapaxar provides only limited therapeutic benefit to a small group of patients without significantly increasing major bleeding (6,7). In line with the clinical experience of other combination antiplatelet therapies, the narrow therapeutic window of vorapaxar in the presence of standard-of-care antiplatelet drugs has translated to limited clinical utility. As a result, there has been much renewed interest in targeting the ‘second’ platelet thrombin receptor, PAR4, for antithrombotic therapy. Although previous studies have rationalised PAR4 as a viable antithrombotic target (8–11), the work

**Table 1** Characteristics of the PAR4 antagonist BMS-986120

Test	Experimental system	Parameter measured	Value for BMS-986120
Binding	HEK293T membranes	$K_d$	0.098 nM
		$K_{on}$	$0.12 \text{ nM}^{-1} \cdot \text{min}^{-1}$
		$K_{off}$	$0.008 \text{ min}^{-1}$
Specificity	HEK293T	Calcium mobilisation $IC_{50}$ vs. PAR4-AP	0.56 nM
	HEK293T	Calcium mobilisation $IC_{50}$ vs. PAR1-AP	>5,000 nM
	CHO	Calcium mobilisation $IC_{50}$ vs. PAR2-AP	>42,000 nM
Efficacy (inhibition of PAR4 signaling)	HEK293T	$IC_{50}$ : activation of $G_{\alpha 11}$	3.4 nM
		$IC_{50}$ : activation of $G_{\alpha q}$	3.9 nM
		$IC_{50}$ : activation of $G_{\alpha 14}$	31 nM
		$IC_{50}$ : $\beta$ -arrestin 2 recruitment	7.2 nM
		$IC_{50}$ : ERK1/2 activation	47 nM
Efficacy (inhibition of platelet aggregation)	Platelet-rich plasma (human)	$IC_{50}$ vs. $\gamma$ -thrombin	7.3 nM
	Whole blood (human)	$IC_{50}$ vs. PAR4-AP	9.5 nM
	Whole blood (monkey)	$IC_{50}$ vs. PAR4-AP	2.1 nM
Efficacy (prevention of <i>in vivo</i> thrombosis)	Monkey	Carotid artery occlusion time (fold-increase vs. vehicle)	2.7 $\times$ , 0.2 mg/kg; 3 $\times$ , 0.5 mg/kg; no occlusion, 1 mg/kg
		Thrombus weight (% reduction vs. vehicle)	36%, 0.2 mg/kg; 50%, 0.5 mg/kg; 82%, 1 mg/kg
Safety (impact on <i>in vivo</i> hemostasis)	Monkey	Kidney bleeding time (fold increase vs. vehicle)	1.4 $\times$ , 0.2 mg/kg; 1.9 $\times$ , 0.5 mg/kg; 2.2 $\times$ , 1 mg/kg
		Mesenteric bleeding time (fold increase vs. vehicle)	1.4 $\times$ , 0.2 mg/kg; 1.7 $\times$ , 0.5 mg/kg; 1.8 $\times$ , 1 mg/kg

PAR4, protease-activated receptor-4.

by Wong and colleagues expands on this to describe the development of a potent and specific small molecule PAR4 antagonist with a markedly improved therapeutic window over one standard antiplatelet drug (clopidogrel) in a preclinical model.

### Discovery of a potent and specific small molecule PAR4 antagonist

In proof-of-concept work that supports previous studies (8-11), the team at BMS first used function-blocking anti-PAR4 antibodies in a guinea pig model to show *in vivo* antithrombotic efficacy and relative safety of selective PAR4 blockade. To shift to the highly desired small molecule approach, they then embarked on an impressive drug discovery program. The unique activation mechanism of

PARs has provided a major hurdle for the development of efficacious antagonists. Thrombin cleavage of PARs reveals an endogenous tethered ligand which then binds to and self-activates the receptor. Therefore, antagonists must overcome an agonist that is intrinsic to the receptor and presumably has considerable steric advantage. Wong and colleagues screened a library of over 1 million compounds to identify a lead candidate that was then subject to iterative rounds of medicinal chemistry and testing to result in BMS-986120—a potent and selective PAR4 antagonist with impressive oral bioavailability and antithrombotic efficacy (Table 1).

Selective inhibition of PAR4 over PAR1 has been elusive in previous efforts developing small molecule PAR4 antagonists. For example, the indazole-derivative YD-3 (12) and its derivative ML354 (13) exhibit cross-



reactivity toward PAR1 (12,13). Here though, BMS-986120 demonstrated specificity in both HEK293 cells transfected with PAR4 and human platelets, with no effect observed on platelet activation by a PAR1 activating peptide, collagen, ADP, or thromboxane A<sub>2</sub> (Table 1). To demonstrate efficacy of BMS-986120, the authors examined inhibitory profiles of platelet aggregation against two isoforms of thrombin— $\alpha$  and  $\gamma$ , thought to preferentially activate PAR1 and PAR4 respectively. BMS-986120 effectively suppressed platelet aggregation in response to  $\gamma$ -thrombin but required concomitant PAR1 inhibition to do so against  $\alpha$ -thrombin. Whether complete blockade of thrombin-induced platelet activation will be required for effective antithrombotic therapy, or whether partial inhibition will be sufficient, remains to be determined.

For PAR antagonists to be efficacious against endogenous enzymatic activation of the receptor by thrombin, they must exhibit strong binding affinity. Yet in the clinical context it is highly desirable for an anti-platelet agent to have the potential to be rapidly reversed should any unwanted bleeding challenges occur. Wong *et al.* used (<sup>3</sup>H)-BMS-986120 binding to PAR4-expressing cell membranes to reveal that BMS-986120 is a high affinity and reversible binder of PAR4 (Table 1). In studies performed on platelets isolated from monkeys dosed with BMS-986120, this translated to normalised aggregation 24 h after a single dose of 0.2 mg/kg. A more detailed time course will be required to determine the half-life of BMS-986120. Although the dissociation constant is relatively fast, competition studies would be useful to demonstrate whether there was potential for an antidote if required. Regardless, the pharmacodynamic profile is considerably advantageous in comparison to other antiplatelet drugs. For example, platelet inhibition by the PAR1 antagonist vorapaxar is retained 4–8 weeks after a single loading dose in humans (14) while aspirin and clopidogrel are both irreversible protein modifiers with long-term effects.

Most importantly though, BMS-986120 appeared to provide an impressive therapeutic window, with a single oral dose of BMS-986120 providing marked antithrombotic effects and a low bleeding profile in a series of *in vivo* models in the cynomolgus monkey (Table 1). Appropriate examination of *in vivo* platelet PAR function is limited to primates since the traditional small animal models (e.g., mice, rats, guinea-pigs, rabbits, dogs) have a different PAR expression profile to that of humans. Therefore, Wong and colleagues used an electrolytic model

of carotid artery thrombosis in the cynomolgus monkey for preclinical evaluation of their PAR4 antagonist. BMS-986120 alone prevented occlusive thrombus formation at the highest dose (1 mg/kg) and significantly reduced thrombosis at lower doses (0.2–0.5 mg/kg). Hemostasis, measured either by kidney or mesenteric bleeding time, was increased by just over 2-fold at the 1 mg/kg dose and much less at the lower doses. Importantly, when this dose-response of BMS-986120 on hemostasis and thrombosis was compared directly with that of clopidogrel, there was a clear separation provided by BMS-986120 that was not evident with clopidogrel. At doses of these two agents that caused equivalent anti-thrombotic effects, markedly more bleeding was observed with clopidogrel compared with BMS-986120. For example, at a dose that caused a 50% reduction in thrombus weight, clopidogrel induced a 7.3- to 8.1-fold increase in bleeding compared with a 1.7- to 1.9-fold increase for BMS-986120. This was more pronounced at doses that caused a near 100% reduction in thrombus weight, with clopidogrel inducing a >10-fold increase in bleeding versus a 1.8- to 2.2-fold increase for BMS-986120. Given this stark difference, it would be interesting to know how the therapeutic window changed when used in combination with aspirin, P2Y<sub>12</sub> antagonists and/or vorapaxar.

### A potential mechanism for an improved therapeutic window

How is it that PAR4 inhibition provides such strong separation between impacting on thrombosis and hemostasis? One clue comes from recent work indicating that PAR4 performs distinct functions to other key platelet receptors. PAR4 activation elicits a slower, but significantly more sustained, intracellular calcium response than that elicited by PAR1 (15). This prolonged calcium signal mediates later-stage platelet activation events, such as the platelet procoagulant response involving phosphatidylserine exposure on the platelet membrane and consequent assembly of coagulation factors leading to thrombin generation and fibrin formation. Indeed, selective inhibition of PAR4 but not PAR1 significantly inhibits thrombin activity and fibrin deposition in human thrombi *ex vivo* (8). One explanation for the improved therapeutic window of BMS-986120 reported by Wong *et al.* is that PAR4 inhibition is blocking platelet function at a distinct time and place to all existing approaches.

### What does the future hold for PAR4 antagonists?

BMS-986120 was evaluated in a phase 1 dosing study, yet despite efficacy and a lack of adverse events no phase 2 studies of this compound were undertaken. Rather, BMS are investigating the related compound, BMS-986141, which also underwent a phase 1 study (NCT02341638) and a subsequent phase 2 trial for the prevention of mini-stroke (NCT02671461). The trial (A Phase 2, Placebo Controlled, Randomized, Double-Blind, Parallel-Arm Study to Evaluate Efficacy and Safety of BMS-986141 For the Prevention of Recurrent Brain Infarction in Subjects Receiving Acetylsalicylic Acid Following Acute Ischemic Stroke or Transient Ischemic Attack) had a primary efficacy endpoint of a composite of symptomatic ischemic stroke or unrecognized brain infarction, and a primary safety endpoint of a composite of adjudicated major bleeding and adjudicated clinically relevant non-major bleeding during the treatment period. It was completed in April 2017 but has not yet been reported.

It is far too early to predict the likely clinical success and/or usefulness of PAR4 antagonists, and several key questions remain. How well will PAR4 antagonism combine with current standard-of-care agents? This is a central point, since any trial will be conducted in the presence of standard-of-care, which frequently involves dual antiplatelet therapy. With the PAR1 antagonist vorapaxar, for example, the increased bleeding observed is believed to be due to poor compatibility with clopidogrel. Indeed, sub-study analyses show no additional bleeding in patients receiving aspirin plus vorapaxar versus those receiving aspirin alone (6,16). Here, it is interesting to note that BMS chose to investigate a patient group being treated with aspirin alone in its first phase 2 trial of its lead PAR4 antagonist.

What specific indications will be best served by a PAR4 antagonist? Again, sub-study analyses of the vorapaxar trials may provide pointers. These trials showed the most efficacy in reducing the rate of spontaneous myocardial infarction as well as in prevention of vascular complications associated with peripheral artery disease. This is perhaps unsurprising given the well-known role of thrombin generation in acute myocardial infarction, particularly in patients with a background of unstable angina and/or coronary artery disease (17). Whether PAR4 antagonism will similarly demonstrate superior efficacy in these clinical situations where thrombin-induced platelet activation are implicated is an obvious place to start for future clinical trials.

Finally, one emerging issue for PAR4 antagonism is

that of population genetics. Recent studies have revealed a commonly expressed genetic variant of PAR4 (rs773902; encoding either Ala120 or Thr120) that appears to significantly alter receptor pharmacology. Specifically, the Thr120 PAR4 variant, expressed in 20–80% of people depending on the population, renders the receptor hypersensitive to agonists and hypo-sensitive to antagonists (18,19). The mechanism behind this change in PAR4 pharmacology remains unknown, as does whether all PAR4 antagonists, including BMS-986120 and BMS-986141, will be similarly affected. Studies directly addressing these points will be critical in determining whether the approach proposed by Wong *et al.* will afford consistent antithrombotic benefit across the population.

### Conclusions

The recent preclinical study by Wong *et al.* (5) details the development and preclinical evaluation of the first PAR4 antagonist to enter a clinical trial and represents a potentially important breakthrough in the treatment of arterial thrombosis. While further insights are still to be gained regarding the utility of PAR4 antagonism in clinical settings, this study has contributed an important reagent to help study this previously under-appreciated platelet activation mechanism, and has identified a potentially useful approach for the safe and effective prevention of arterial thrombosis.

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### Footnote

**Conflicts of Interest:** The authors have no conflicts of interest to declare.

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