

Protease-Activated Receptor-1 Mediates Thrombin-Induced Persistent Sodium Current in Human Cardiomyocytes

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ABBREVIATIONS: HEPES, N-(2-hydroxyethyl)-piperazine-N'-2-ethanesulfonic acid; EGTA, ethylene glycol-bis(beta-aminoethyl ether)-N,N,N',N'-tetra acetic acid; PLA₂, phospholipase A₂; DMSO, dimethyl sulfoxide; TTX, tetrodotoxine;

ABSTRACT

Following the thrombus formation in cardiac cavities or coronaries, the serine protease thrombin is produced and can therefore reach the myocardial tissue by the active process of extravasation and binds to the G protein-coupled protease-activated receptor-1 (PAR1) expressed in human myocardium. The role of PAR1 was investigated in the thrombin effect on sodium current (I_{Na}). I_{Na} was recorded in freshly isolated human atrial myocytes by the whole-cell patch-clamp method. Action potentials (AP) were recorded in guinea pig ventricular tissue by the conventional glass microelectrodes technique. Thrombin-activated PAR1 induced a tetrodotoxin-blocked persistent sodium current, I_{NaP} , in a concentration-dependent manner with an apparent EC_{50} of 28 U/ml. The PAR1 agonist peptide SFLLR-NH₂ (50 μ M) was able to mimic PAR1-thrombin action, whereas PAR1 antagonists, SCH 203099 (10 μ M) and ER 112787 (1 μ M), completely inhibited it. The activated-PAR1 involves calcium-independent phospholipase-A₂ signaling pathway since two inhibitors of this cascade, bromoenol lactone (50 μ M) and haloenol lactone suicide substrate (50 μ M), block PAR1-thrombin-induced I_{NaP} . As a consequence of I_{NaP} activation, in guinea pig right ventricle papillary muscle, action potential duration (APD) were significantly increased by 20% and 15% under the respective action of 32 U/ml thrombin and 50 μ M SFLLR-NH₂, and these increases in APD were prevented by 1 μ M TTX or markedly reduced by application of 1 μ M SCH 203099 or ER 112787. Thrombin, through PAR1 activation, increases persistent component of the Na current resulting in an uncontrolled sodium influx into the cardiomyocyte which can contribute to cellular injuries observed during cardiac ischemia.

Introduction

Thrombin is a serine protease released by thrombus with well-characterized roles in hemostasis, inflammation, and proliferative process. During cardiac ischemia-reperfusion, thrombin is an important mediator of myocardial injury (Erlich et al., 2000). Indeed, thrombin can permeate vessels or endocardial barrier by extravasation and therefore can affect the first layers of cardiomyocytes. Thus, thrombin levels may be locally elevated at site of vascular injury and thrombus formation, and reach an activity as high as 10 to 30 U/ml (Park et al., 1994). Then thrombin has multiples cellular effects mediated by a family of G-protein-coupled protease activated receptors (PARs) of which PAR1 is the prototype (Coughlin, 2000). PAR1 are expressed in human myocardium (Jiang et al., 1996) and activates a spectrum of biochemical signals leading to changes in contractile performance and alteration in gene expression (Glembotski et al., 1993), sarcomeric organization and cardiomyocyte morphology (Sabri et al., 2002). As shown recently, these phenomena are clearly involved in human coronary atherosclerosis and can lead to endothelial dysfunction and vascular inflammation (Lavi et al., 2007). Thus, thrombin stimulates phosphoinositide hydrolysis, for instance by converting phosphatidylcholine into lysophosphatidylcholine (LPC), *via* PLA₂ (Park et al., 1994; Sabri et al., 2000). Thrombin also activates the extracellular signal-regulated protein kinase (ERK) (Sabri et al., 2002), facilitates rapid sodium current (Pinet et al., 2002), modulates calcium homeostasis (Steinberg, 2005; Steinberg et al., 1991), increases arrhythmias (Goldstein et al., 1994) and hastens recovery from an imposed acid load by activating Na⁺-H⁺ exchange (Avkiran and Haworth, 2003). Taken together, these signal events profoundly alter electrophysiological

properties and contractile behavior, and could induce cardiomyocyte toxicity during the myocardial ischemia-reperfusion injuries.

Increase in intracellular sodium during ischemia plays a key role in intracellular calcium overload resulting in myocyte injuries. The involvement of thrombin-dependent activation of PAR1 in Na⁺ entry during ischemia has never been assessed. In a previous study, we reported that thrombin can reversibly act as direct agonist on I_{Na} . The main consequences of this action were the shift towards hyperpolarizing potentials of the activation-voltage relationship, the large increase in peak I_{Na} amplitude and the consequent increase in the window of sodium current, however this effect was independent of PAR1 activation (Pinet et al., 2002). The present study examines whether PAR1 signaling in human cardiomyocytes is involved in the effect of thrombin on sodium current specially the persistent sodium current (I_{NaP}) (Fedida et al., 2006; Haigney et al., 1994; Maltsev et al., 1998; Noble and Noble, 2006; Saint, 2006). The main result was that thrombin, by binding to PAR1 receptor, activates I_{NaP} .

Materials and Methods

Heart tissue samples. Protocols for obtaining human cardiac tissue were in conformation with the principles outlined in the Declaration of Helsinki. We used a total of 62 specimens of human right atrial appendages which were obtained from hearts of patients (51 to 74 years) undergoing heart surgery for coronary artery bypass graft (CABG) or valve replacement. 7% of patients have received β -blockers, 38% calcium antagonists, 45% antiulcer drugs, 8% diuretics, and 22% anti-thrombotics. Treatments were usually stopped 24 hours before operation. Patients with atrial dilation were avoided, and none had a history of supraventricular arrhythmias.

Cell isolation. Human atrial myocytes were isolated enzymatically as previously described (Antoine et al., 1998). The same chunk procedure was used to isolate myocytes from the right ventricle of guinea pig hearts with the difference that elastase was omitted in the chunks pre-digestion bath. Only quiescent rod-shaped myocytes with clear cross-striations, sharp edges and a well-delineated cell membrane were chosen for experiments. Small myocytes were preferred to optimize spatial voltage-clamp.

Solutions and drugs. For whole-cell current recordings, the intracellular pipette solution contained (in mM): NaCl 5, CsCl 130, MgCl₂ 2, CaCl₂ 1, EGTA 15, HEPES 10, MgATP 4, pH 7.2 with CsOH; the basal external solution contained: NaCl 25, CsCl 108.5, CoCl₂ 2.5, CaCl₂ 0.5, MgCl₂ 2.5, 4-aminopyridine 5, HEPES 10, glucose 10, pH 7.4 with CsOH. Hirudin (leech, \approx 2000 U/mg) was from Boehringer Mannheim. HELSS (haloenol lactone suicide substrate) from TEBU France, was dissolved in dimethyl sulfoxide (DMSO) which did not exceed 0.05%. The protease activated receptor (PAR1) agonist (SFLLR-NH₂) and antagonist Schering 203099 (SCH 203099) were from the Division of Medicinal Chemistry IV (Centre de Recherche Pierre Fabre, Castres, France), alpha-thrombin (human plasma \approx 1000 U/mg), and other chemicals were

purchased from Sigma. Commercially available lyophilized thrombin is provided as a sodium salt. The addition of 32 U/ml of thrombin to the basal perfusion medium containing 25 mM NaCl, increases the Na^+ activity to 32.1 ± 0.3 mM (n=10) (measured with a sodium ion-selective electrode). The control perfusion medium was therefore supplied with NaCl to keep the Na^+ activity equivalent.

Current recordings. Ionic currents were recorded by the whole-cell patch-clamp technique with a patch-clamp amplifier (Axopatch 200B, Axon Instruments, Foster City, CA). Patch pipettes (Corning Kovar Sealing code 7052, WPI, FL) had resistances of 0.5-2.0 M Ω . Currents were filtered at 20 kHz (-3 dB, 8-pole low-pass Bessel filter) and digitized at 50 kHz (Digidata 1200, Axon Instruments). Cell membrane capacitance (C_m) was 48.2 ± 2.6 pF (n=67 cells, from 62 donors) for human atrial myocytes and 62.4 ± 7.5 pA/pF (n=18, from 10 hearts) for guinea pig right ventricular myocytes. Series resistance (R_s) was compensated at 80-95% resulting in voltage errors of <3 mV. Leakage current was compensated whereas cell membrane capacitive current was not. Peak I_{Na} and I_{NaP} amplitudes were monitored according to a steady-state pulse protocol: a 1000 ms depolarizing test pulse to -30 mV from a HP of -100 mV at 0.2 Hz. It is worth noting that the test pulse was chosen to be beyond the upper limit of the potential range inside which the sodium window current has been observed to increase under the direct effect of thrombin on sodium channel (Pinet et al., 2002). To avoid a putative overlap between PAR1-induced I_{NaP} and thrombin-increased sodium window current, experiments were also performed with test pulse to -10 mV. An equilibration period was allowed until peak I_{Na} reached steady-state and remained stable without evidence of a leftward shift of the availability-voltage relationship (h_{∞} - V_m). This protocol was designed to detect any such shifts. Following each sequence of five depolarizing pulses, HP was set to -140 mV. Consequently during the stabilization period when peak current

amplitude was higher following a HP at -140 mV compared to -100 mV, the recording was discarded. Similarly, since thrombin has no effect on the h_{∞} - V_m relationship (Pinet et al., 2002), recordings showing an irreversible hyperpolarizing shift of h_{∞} - V_m after thrombin application, were also discarded. The steady-state pulse protocol was applied in control, thrombin or when other substances were tested and washed out. Experiments were carried out at room temperature (22-25°C).

Action Potential recording. Male guinea pigs were anesthetized by intraperitoneal pentobarbital sodium injection and their hearts were quickly removed. The investigation conforms with the *Guide for the care and Use of Laboratory Animals* published by the US National Institutes of Health (NIH Publication No.85-23, revised 1996). A total of 21 hearts were used for the AP study. Papillary muscles were dissected from the right ventricle, and superfused (4-6 ml/min) with the Krebs solution maintained at $36.0^{\circ}\text{C} \pm 0.5^{\circ}\text{C}$ in a 5 mL tissue bath. The Krebs solution was oxygenated with 95% O_2 / 5% CO_2 and had the following composition (in mM): NaCl 113.1, KCl 4.6, MgCl_2 2.45, NaH_2PO_4 3.5, NaHCO_3 21.9, and glucose 5 (pH 7.4). Action potentials (AP) were recorded by conventional 'floating' glass microelectrodes (5-20 M Ω) filled with KCl 3M and were coupled to a high-input impedance preamplifier (VF 102 Biologic, Echirrolles, France). The preparations were electrically stimulated (PULSAR BP, FHC Brunswick, ME, U.S.A.) with 1 ms pulses at 1.5 times the threshold voltage through a bipolar Ag-electrode. APs were displayed on a dual-beam oscilloscope (TDS 2012, Tektronics, Heerenveen, The Netherlands) and simultaneously digitised and analysed with interactive software (NOTOCORD hem 3.4, Notocord Systems, France). The preparations were allowed to equilibrate for at least 1 h at a stimulation rate of 1Hz. A single impalement was maintained throughout control and compound superfusion periods. AP parameters measured were maximum

upstroke velocity $(dV/dt)_{\max}$, amplitude, resting membrane potential, and action potential duration (APD) at 50 and 90% repolarization levels (APD₅₀ and APD₉₀, respectively).

Data analysis and statistics. As much as possible, depending on the experimental protocol performed, only the TTX-inhibited PAR1-induced I_{NaP} was taken into account and determined by subtracting the current obtained under concomitant application of TTX and thrombin or SFLLR-NH₂ from that previously recorded under thrombin alone. The amplitude of TTX-blocked thrombin-induced I_{NaP} was the mean current calculated from 80 to 180 ms after the beginning of the depolarization (Valdivia et al., 2002). Peak I_{Na} amplitude was measured with respect to current amplitude at the end of the test pulse. Data are expressed as means \pm SEM of n determinations or myocytes. Statistical analysis were performed by using paired or unpaired Student's t -test, or ANOVA, as appropriate and the null hypothesis was rejected at the 0.05 level; *: $p < 0.05$; **: $p < 0.01$ and ***: $p < 0.001$.

Results

Thrombin application induces a persistent sodium current different from the thrombin-increased sodium window current. In order to be beyond the upper limit of the range of membrane potential (from -85 to -40 mV) in which sodium window current occurs, a test pulse to -30 mV was used to elicit I_{Na} (Pinet et al., 2002). Local perfusion of 32 U/ml thrombin on freshly isolated human atrial myocytes caused the already described marked increase in peak of I_{Na} (Figure 1 A) which occurred rapidly (Figure 1D) and was shown to be completely independent of the activation of PAR1 receptors (Pinet et al., 2002). In addition to this effect on peak I_{Na} , following a longer time of application, over 6 min, thrombin induced very slowly inactivating or persistent TTX-blocked sodium current (Figure 1B and 1D) termed as I_{NaP} . While the increase in peak I_{Na} was observed in all tested cells, the induction of I_{NaP} was observed in 51 out of 67 cells, most probably as a consequence of the lost and/or damage caused to PAR1 receptors by the enzymatic procedure used for cell dissociation. Thrombin (32 U/ml) increased peak I_{Na} , from -118.9 ± 12.6 to -191.7 ± 20.8 pA/pF ($n=23$; $p < 0.001$, paired t-test) and induced I_{NaP} in a concentration-dependent manner (Figure 1C). At this concentration of thrombin, maximum increase in I_{NaP} was 1.8% of control peak I_{Na} . The density of I_{NaP} was -0.02 ± 0.01 pA/pF, $n=3$ with 1 U/ml thrombin, -0.50 ± 0.02 pA/pF, $n=3$ with 10 U/ml thrombin, -2.23 ± 0.54 pA/pF, $n=6$ with 32 U/ml thrombin and -3.69 ± 0.82 pA/pF, $n=4$ with 100 U/ml thrombin. Estimated-maximal density of I_{NaP} was -3.98 pA/pF, the apparent EC_{50} and the Hill coefficient were respectively 28.4 U/ml and 1.98. This concentration-response relationship exhibits clear differences when compared to the reported concentration-response of the effect on peak I_{Na} (Pinet et al., 2002) where EC_{50} and Hill coefficient were respectively 91 U/ml and 0.75. Thrombin-induced I_{NaP} was not a residual current resulting from the thrombin-enhanced window sodium

current (Pinet et al., 2002) since the increase in I_{NaP} , with 32 U/ml thrombin, was observed with test pulses to -10 mV, a membrane potential at which the window current can not be activated (I_{NaP} density was -1.71 ± 0.83 pA/pF; $n=5$). These two distinct effects of thrombin on sodium current (increase in peak I_{Na} and in I_{NaP}) were suppressed by 50 μM of the specific sodium channel blocker TTX (Figure 1A and 1B). It is worth noting the difference in kinetics of thrombin effect on peak I_{Na} compared to the onset of I_{NaP} (Figure 1D), and that the time to reach maximum thrombin effects was much longer for I_{NaP} (416.0 ± 40.4 s, $n=15$), than for peak I_{Na} (61.7 ± 10.1 s, $n=12$; Figure 1E) suggesting distinct underlying mechanisms. This is also supported by the observation that after washout of thrombin, the amplitude of peak I_{Na} returned to the basal (Pinet et al., 2002) but not that of I_{NaP} (Figure 1D) in accordance with the irreversibility of the cleavage of PAR1 by thrombin. The panel below of Figure 1D exemplifies the prominence of thrombin effect on I_{NaP} over the effect on peak I_{Na} when the latter effect wore out.

Involvement of the PAR1 in thrombin-induced I_{NaP} . To test further for the requirement of protease activity of thrombin in the activation of I_{NaP} , we investigated the effect of hirudin, a direct protease inhibitor of thrombin. The induction of I_{NaP} by 32 U/ml thrombin was markedly reduced in presence of 320 U/ml hirudin (Figure 2B, 2C and 2D, I_{NaP} density was, under thrombin, -2.27 ± 0.51 pA/pF, ($n=8$) vs -0.43 ± 0.19 pA/pF under thrombin in presence of hirudin, $n=9$; $p<0.001$) whereas the effect of thrombin on peak I_{Na} was unchanged (Figure 2A and 2C). This result confirms that the protease activity of thrombin is necessary to activate I_{NaP} probably through the cleavage of still undefined target proteins. Among the proteins cleaved by thrombin, the most likely target protein is the well known PAR1 receptor. In order to study the involvement of this receptor, selective PAR1-antagonist, SCH 203099 (Ahn and Chackalamannil, 2001), have been used. Following human atrial cell-preincubation with 10 μM SCH 203099 for at

least 15 min, thrombin failed to induce I_{NaP} (Figure 3B, 3C and 3D); -0.30 ± 0.07 pA/pF (n=14) vs -2.27 ± 0.51 pA/pF, n=8 in absence of SCH 203099, $p < 0.001$), whereas the thrombin effect on peak I_{Na} was unaffected (Figure 3A and 3C; from -103.7 ± 14.1 pA/pF under SCH 203099 to -158.4 ± 21.6 pA/pF (n=14) after addition of 32 U/ml thrombin; ($p < 0.001$)). These results suggest that PAR1 activation constitutes a key mechanism for the thrombin-induced I_{NaP} . Then, the PAR1 agonist peptide, SFLLR-NH₂, was tested in order to putatively mimic the role of thrombin in induction of I_{NaP} via PAR1 (Figure 4). SFLLR-NH₂ stimulated I_{NaP} (mean increased: -1.46 ± 0.63 pA/pF, n=5, $p < 0.05$ vs control (Fig 4B and 4C), exhibiting a fair reversibility upon washout (Fig. 4B and 4D), but failed to increase peak I_{Na} (Figure 4A and 4D). Taken together, the present results demonstrate that the thrombin-induced I_{NaP} is mediated by PAR1 activation.

Involvement of Ca²⁺-independent phospholipase-A₂ pathways consecutively to PAR1 activation. PAR1 is a member of the G-protein coupled receptors family, and therefore its activation stimulates numerous of intracellular signaling pathways (Coughlin, 2000). Several studies have reported that in cardiomyocytes, thrombin stimulates the Ca²⁺-independent phospholipase-A₂ (PLA₂) via PAR1, which produced an intracellular lysophosphatidyl choline (LPC) accumulation (Park et al., 1994; Yan et al., 1995). Therefore, two inhibitors of PLA₂ (bromo-enol lactone (BEL) and halo-enol lactone suicide substrate (HELSS)) were used to investigate whether PLA₂ activity is involved in the enhancement of I_{NaP} , induced by PAR1 stimulation. Following cell-incubation with BEL during at least 30 min, thrombin had its usual effect on peak I_{Na} (Figure 5A) but failed to stimulate I_{NaP} (Figure 5B: -0.22 ± 0.11 pA/pF, n=5 vs -2.27 ± 0.51 pA/pF, n=8, in absence of BEL; $p < 0.01$). A similar result has been obtained with HELSS (Figure 5C). The density of I_{NaP} activated in presence of thrombin and HELSS was -0.39

± 0.14 pA/pF, $n=5$; ($p < 0.01$). These results indicate that the PAR1-induced I_{NaP} requires the activation of Ca^{2+} -independent PLA_2 .

Thrombin and PAR1 agonist peptide SFLLR-NH₂ increase action potential duration. To examine the consequence of activation of I_{NaP} by thrombin and PAR1 agonist peptide SFLLR-NH₂ on cardiac tissue electrical activity, we recorded action potentials in right ventricle papillary muscle from guinea pigs. We used the classical microelectrode technique which allows a stable and reliable AP recording during prolonged period as required for the study of the delayed effect of thrombin on I_{NaP} . We first checked that, in this specie and in ventricular cells, thrombin also activates I_{NaP} by performing patch-clamp experiments in isolated guinea pig ventricular myocytes. Test pulses to -10 mV from a holding potential of -80 mV were used to elicit current. Immediate application of thrombin (32 U/ml) increased peak I_{Na} (Figure 6Aa and 6Ab) from -112.6 ± 11.5 pA/pF to -166.9 ± 16.4 pA/pF, $n=11$, ($p < 0.001$; paired t-test) blocked by application of 50 μ M TTX. Moreover, following 7 minutes of application, thrombin induced a TTX-sensitive I_{NaP} of -2.91 ± 1.1 pA/pF, $n=9$ (Figure 6Ab and 6Ad). The increase in I_{NaP} was of 0.026% of control peak I_{Na} . This effect was blocked by the PAR1 antagonist, ER 112787 (Barry et al., 2006; Chackalamannil and Xia, 2006) (Figure 6B). Following myocyte-preincubation with 1 μ M ER 112878 for at least 5 min, thrombin did not induced I_{NaP} (Figure 6Bb, 6Bc and 6Bd; -0.076 ± 0.05 pA/pF ($n=5$) vs -2.91 ± 0.68 pA/pF, in absence of drug, $n=9$; $p < 0.001$), whereas the effect of thrombin on peak I_{Na} was unaffected (Figure 6Aa and 6Ac; -87.3 ± 6.1 pA/pF, $n=5$, under ER 112787 to -123.4 ± 10.3 pA/pF, $n=9$, after addition of thrombin; $p < 0.05$).

Figure 7Aa shows the effects of thrombin (32 U/ml) and SFLLR-NH₂ on action potentials recorded from right papillary ventricle. Following 10-minutes application, thrombin and SFLLR-NH₂ increased significantly action potential duration (APD) measured at 50 and 90% of

repolarisation (Figure 7A and Table 1A). At these two levels of repolarisation, the increases in APD were respectively 15.3 and 12.9 % for thrombin, and 12.4 and 12.1% for SFLLR-NH₂ (Table 1A). The increases in APD were hardly reversible (Table 1A). Importantly, in presence of TTX (1 μM), applications of thrombin (32 U/ml) or SFLLR-NH₂ (100 μM) failed to increase the action potential duration (Figure 7Ab, Table 1B). We used a low concentration of TTX to minimize the consequences of the inhibition of peak I_{Na} on AP upstroke.

To investigate the involvement of PAR1 on thrombin- and SFLLR-induced APD lengthenings, two selective PAR1 antagonists, SCH 203099 and ER 112787, were then evaluated. Figure 7Ba and 7Bb show that both SCH 203099 and ER 112787, in concentration-dependent manner, reduced the thrombin- and SFLLR-NH₂-induced APD prolongations. Thus, SCH 203099 (10 μM) and ER 112787 (1 μM) both fully blocked the APD prolongation induced by 100 μM SFLLR-NH₂ (Fig 7B), a result which is in a complete agreement with the potency of this antagonist against PAR1 (Chackalamannil and Xia, 2006). Taken together these results indicate that stimulation of the inward persistent Na⁺ current by protease activated receptors following PAR1 activation can modify the AP duration.

Discussion

The present results demonstrate for the first time that thrombin induces a PAR1-mediated persistent sodium current component, I_{NaP} , in cardiomyocytes. Because thrombin is a serine protease formed at the site of coronary vascular wall injury, this effect on Na current might have important consequences in the setting of myocardial ischemia.

Involvement of PAR1 signaling pathway in the thrombin effect on I_{NaP} . The conclusion that thrombin enhances I_{NaP} through the proteolytic cleavage of PAR1 is supported by the findings that this thrombin effect on current is mimicked by the PAR1 peptide agonist, SFLLR-NH₂, and is fully blocked by the PAR1 antagonists, SCH 203099 and ER 112787 (Ahn and Chackalamannil, 2001). Cleavage of the amino-terminal PAR1 exodomain unmasks a tethered ligand that binds the receptor body to trigger intracellular signaling. Consequently, PAR1 is irreversibly activated by this proteolytic cleavage, contributing to keep on its signal-transducing capability. This could explain the low reversibility of the thrombin effect on I_{NaP} upon washout (see Fig. 1). This was not the case for SFLLR-NH₂ (see Fig. 4), an exogenous peptide that does not cleave the PAR1, and which exhibited an effect more sensitive to washout. Another evidence that thrombin effect on I_{NaP} is mediated by PAR1 signaling pathways is provided by the observation that this effect is suppressed by the inhibition of PLA₂ by BEL and HELSS. There are previous studies showing that the increase in intracellular LPC levels plays an important role in the thrombin effect (Park et al., 1994; Yan et al., 1995; Undrovinas et al., 1992). For instance, the cleavage of PAR1 by thrombin stimulates the amphipathic lipids catabolism leading to an accumulation of LPC in the intracellular medium (Park et al., 1994; Undrovinas et al., 1992). However, the precise level of the signaling pathway at which PLA₂ inhibitors were able to inhibit thrombin-induced I_{NaP} remains to be clarified.

Nature of the I_{NaP} current. The intracellular accumulation of LPC could mediate the effect of thrombin on I_{NaP} . The LPC is well known to maintain sodium channel in bursting activity thus giving rise to non-inactivating sodium current (Undrovinas et al., 1992). The exact nature of this non-inactivating sodium current is not yet fully elucidated. There are evidences that this persistent component of sodium current, I_{NaP} , could be generated by a small fraction of the "normal" transient-mode sodium channel population, that undergoes burst or/and late scattered mode(s). Such modes have been recently described by Maltsev and Undrovinas (Maltsev and Undrovinas, 2005) for the persistent Na^+ current in human ventricular myocytes. Thus, thrombin could favor this bursting activity of voltage-gated sodium channels, by cleaving PAR1 and stimulating the conversion of phosphatidylcholine into LPC which, *via* PLA₂ (Park et al., 1994; Sabri et al., 2000), increases intracellular LPC (Park et al., 1994; Undrovinas et al., 1992). In a previous study, we reported that thrombin stimulates the peak I_{Na} and the sodium-window current. However, this effect which is fast and fully reversible is not mediated by PAR1 (Pinet et al., 2002) as further demonstrated in the present study. Moreover, thrombin was able to induce I_{NaP} at membrane potentials as high as -10 mV which is clearly not in membrane potential range of window current (from -85 to -40 mV). Finally, there are clear differences between the concentration-response curves of the effect of thrombin on peak I_{Na} (Pinet et al., 2002) and on I_{NaP} . One hypothesis currently tested concerning the direct effect of thrombin on peak I_{Na} is the involvement of a β -subunit (Herfst et al., 2003). The majority of Na^+ channels in the heart correspond to the expression of the TTX-resistant Nav1.5 isoform and this is the case for human atrial myocytes (Makielski et al., 2003). However, it cannot be excluded that thrombin-PAR1 activation targets another population of sodium channels (Brette and Orchard, 2006). The effect

of thrombin on I_{NaP} that develops slowly is likely to have the more significant consequences on cardiac myocyte.

Pathophysiological consequences of the activation by thrombin of I_{NaP} . Both persistent and window sodium current are known to participate to action potential duration. While the window sodium current which activates in a restricted membrane potential range from -85 to -40 mV is involved in the late phase of repolarisation and in the control of resting membrane potential (Pinet et al., 2002), I_{NaP} , activated from -10 mV, is found to contribute to the regulation of the early phase of APD (Fedida et al., 2006; Kiyosue and Arita, 1989; Maltsev et al., 1998; Noble and Noble, 2006; Sakmann et al., 2000; Wu et al., 2006). It has been previously reported that blocking the I_{NaP} with TTX caused a 10-20% decrease of APD though this current is of very small density in control condition (Kiyosue and Arita, 1989; Maltsev et al., 1998; Sakmann et al., 2000). Therefore, the progressive several fold increase in I_{NaP} by thrombin, is likely to have a significant effect on APD, whereas, in spite of a fast increase in peak I_{Na} , (Fig 1), thrombin is devoid of significant effect on $(dV/dt)_{max}$. Indeed, thrombin and SFLLR increase markedly the duration of action potentials of guinea pig papillary muscle. This effect of thrombin or SFLLR-NH₂ on AP duration can be largely explained by an increase of I_{NaP} because it was abolished by TTX. However, it cannot be excluded that the thrombin/PAR1 pathway can affect other ionic currents indirectly *via* changes in $[Na]_i$ and $[Ca]_i$. Previous studies have shown that thrombin lengthens APD and increases cesium-induced early after depolarizations (EAD) and pro-arrhythmic events in canine Purkinje fibers (Steinberg et al., 1991) and in intact adult rat hearts during early reperfusion (Jacobsen et al., 1996; Woodcock et al., 1998). It has been shown also that suppression of the late sodium current can suppress EADs of myocytes isolated from failing hearts (Maltsev et al., 1998; Undrovinas et al., 2002; Valdivia et al., 2005; Fedida et al., 2006).

Thus, blocking I_{NaP} by inhibiting PAR1 – PLA₂ pathway may be a new pharmacological target to reduce thrombin-induced arrhythmic activity.

Yan et al. (Yan et al., 1995) have demonstrated that activation of PAR1 by SFLLR-NH₂ peptide induced a rapid and dramatic elevation in $[Na^+]_i$ which was associated with a concomitant increase in LPC content in isolated, blood-perfused rabbit hearts in response to ischemia (Lavi et al., 2007). The increase in $[Na^+]_i$ could contribute to myocyte injury during ischemia as the result of intracellular calcium overload and the activation of Ca-dependent signaling cascades. This hypothesis is supported by the recent finding of the cardioprotective effects of the late sodium current inhibitor ranolazine (Belardinelli et al., 2006). Moreover, *in vivo* and *in vitro* studies, Strande et al. (Strande et al., 2007), have shown that a preventive and a curative treatments with a selective PAR1 antagonist reduced the infarct size and increased ventricular function recovery following ischemia reperfusion in an isolated heart model. It remains to determine whether the increase in $[Na^+]_i$ induced by SFLLR-NH₂ or by thrombin during ischemia is mediated by the activation of I_{NaP} .

In conclusion, this study describes a new regulatory mechanism of sodium current involving PAR1-PLA₂ signaling pathway which could be evoked by thrombin during cardiac ischemia and thrombus formation (Haigney et al., 1994; Maltsev et al., 1998). Selective antagonists of PAR1 receptor which are very efficient to suppress this effect of thrombin, might represents a novel cardioprotective strategy in the clinical setting of myocardial ischemia and reperfusion (Strande et al., 2007).

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Footnotes

Equal contribution note

The two first authors contributed equally to this study.

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Legends for figures

Fig. 1. Rapid sodium current was modified following thrombin application to human atrial myocytes. Panel A exemplifies that thrombin (32 U/ml) was able to increase the TTX-blocked fast sodium current I_{Na} , as previously described. In B the vertical scale shows a higher gain than in A and the time base is slowed to demonstrate the effect of thrombin on the non-inactivating persistent component of the current (I_{NaP}) which was blocked by 50 μ M TTX. Arrows indicate zero current level. C, concentration-response curve for thrombin-induced I_{NaP} . Normalized mean increase in I_{NaP} amplitude is reported against thrombin activity. Each point represents the mean \pm SEM of n measurements indicated. Data points were fitted by the equation $Y / Y_{max} = 1 / [1 + (EC_{50} / \text{thrombin activity})^{nH}]$. Y_{max} (maximum mean induced- I_{NaP}), EC_{50} (activity inducing half-maximal effect), and Hill parameter nH were respectively 3.98 pA/pF, 28.4 U/ml and 1.98 for I_{NaP} . D, kinetic of the effect of thrombin on I_{NaP} and on peak I_{Na} . Time courses of I_{NaP} (upper panel) and of peak I_{Na} (middle panel) densities, and percentage of induced I_{NaP} over peak I_{Na} , (lower panel) under conditions indicated by horizontal bars. Currents were elicited by depolarization from HP = -140 to -30 mV. E, Bar graphs correspond to the time necessary to obtain the thrombin maximal effect on the two currents; in parenthesis: number of measurements.

Fig. 2. The effect of hirudin, a direct inhibitor of the protease activity of thrombin, was tested on thrombin-induced I_{NaP} . Typical current recordings (represented as in Figure 1) showing that co-application of hirudin (320 mM) leaves unchanged the increase in peak I_{Na} by thrombin (32 U/ml) (A), but prevents the thrombin-induced I_{NaP} (B). C, kinetic comparisons of the action of thrombin in presence of hirudin on I_{NaP} and on peak I_{Na} . Time courses of I_{NaP} (upper trace) and of

peak I_{Na} (lower trace) densities, under conditions indicated by horizontal bars. Currents were elicited as in Figure 1. D, bar graphs confirm that hirudin prevented the induction of I_{NaP} by thrombin.

Fig. 3. As the induction of I_{NaP} depends on the protease activity of thrombin, the implication of the protease-activated-receptor 1 (PAR1), was tested using the PAR1 antagonist SCH 203099. In panels A: current traces exemplifying that SCH 203099 do not prevent the increase in peak I_{Na} by thrombin, whereas it blocks the thrombin-induced I_{NaP} (panels B). C, time courses of I_{NaP} and I_{Na} densities, under conditions indicated by horizontal bars. In D, bar graphs summarize the effect of SCH 203099 vs thrombin on I_{NaP} . Thrombin was used at 32 U/ml, SCH 203099 at 10 μ M.

Fig. 4. SFLLR-NH₂, a synthetic PAR1 agonist peptide, was able to reproduce the potent induction of I_{NaP} by thrombin, but not the increase in peak I_{Na} . Typical current traces of I_{Na} (A) showing activation of I_{NaP} and that peak I_{Na} remains unchanged under SFLLR-NH₂. B, the corresponding traces with different amplitude- and time-scales on right hand. C and D, the effect of SFLLR-NH₂ on I_{Na} was compared to thrombin-activation of I_{NaP} . In C, typical traces of I_{NaP} were obtained under control, SFLLR-NH₂ and thrombin conditions. In panel D, comparison of I_{NaP} (upper panel) and I_{Na} (middle panel) time courses, and percentage of I_{NaP} over I_{Na} (lower panel), under conditions indicated by horizontal bars, showing the activation of I_{NaP} by SFLLR-NH₂, followed by its partial reversion upon washout, and the usual effects of thrombin. SFLLR-NH₂ was used at 50 μ M and thrombin at 32 U/ml.

Fig. 5. Thrombin-induced I_{NaP} requires activation of Ca^{2+} -independent phospholipase A_2 . Effects of bromoenol lactone (BEL), a PLA_2 inhibitor, upon thrombin-increased peak I_{Na} (A) and thrombin-induced I_{NaP} (B). A, sample current traces showing no effect of BEL on thrombin-increased peak I_{Na} , in B, the corresponding traces with different amplitude- and time-scales demonstrating that the presence of BEL prevented the induction of I_{NaP} . Cell was previously pre-incubated with 50 μM BEL during 30 min. C, bar graphs summarizing the inhibition of thrombin-induced I_{NaP} following BEL incubation and haloenol lactone suicide substrate (HELSS) application. Thrombin was used at 32 U/ml and HELSS at 10 μM .

Fig. 6. Guinea pig ventricular myocytes also exhibited a thrombin-increased I_{Na} and a thrombin-induced I_{NaP} . In A, panel (a) exemplifies that thrombin (32 U/ml) was able to increase the TTX-blocked fast sodium current I_{Na} . In (b), the corresponding traces with different amplitude- and time-scales showing the TTX-blocked thrombin-induced- I_{NaP} recorded after 7 min application of thrombin (32 U/ml). TTX was used at 50 μM . In (c) and (d), bar graphs summarizing the effects of thrombin and TTX on I_{Na} and I_{NaP} . In B, with this cell type, another PAR1-antagonist, ER 112787, has been tested. In panel (a): current traces exemplifying that ER 112787 did not prevent the increase in peak I_{Na} by thrombin, whereas it blocked the thrombin-induced I_{NaP} (panel b). In (c), time courses of I_{NaP} and I_{Na} densities, under conditions indicated by horizontal bars. In (d), bar graphs summarize the effect of ER 112787 vs thrombin on I_{NaP} . Thrombin was used at 32 U/ml and ER 112787 at 1 μM .

Fig. 7. A, direct applications of thrombin or of the synthetic PAR1 agonist peptide SFLLR-NH₂ prolonged the action potential duration (a), prolongations prevented in presence of TTX (b). Action potentials were recorded from guinea pig right ventricular papillary muscles before and after 15 minutes following the applications of thrombin (32 U/ml) or SFLLR-NH₂ (100 μM) in a, and when those applications were performed in presence of 1 μM TTX, in b. Arrows indicate zero potential level. B, effects of two selective PAR1 antagonists, SCH 203099 and ER 112787, on thrombin- and SFLLR-induced APD-lengthenings (panels a and b). Bar graphs showing the concentration-responses inhibition effect of SCH 203099 and ER 112787 on thrombin- and SFLLR-lengthened APD of action potentials recorded from guinea pig right ventricular papillary muscles.

Table 1

A, effects of thrombin and SFLLR on action potential parameters

Thrombin 32 U/ml

	n	RP (mV)	APD ₅₀ (ms)	APD ₉₀ (ms)	(dV/dt) _{max} (V/s)
Control	6	-91.6 ± 0.9	134.5 ± 10.3	159.5 ± 11.2	244 ± 29
Thrombin	6	-90.2 ± 0.6	155.1 ± 14.5*	180.2 ± 16.3*	252 ± 33
Washout	4	-91.9 ± 0.4	143.7 ± 16.7	167.6 ± 18.1	264 ± 45

SFLLR-NH₂ 100 μM

Control	6	-92.3 ± 0.4	125.2 ± 19.5	153.7 ± 22.7	219 ± 12
SFLLR-NH ₂	6	-89.4 ± 0.9	140.7 ± 21.8*	172.4 ± 27.1*	213 ± 16
Washout	4	-90.5 ± 0.6	150.3 ± 25.4	181.6 ± 31.6	225 ± 13

B, TTX (1 μM) prevents effects of thrombin and SFLLR-NH₂ on action potential

Thrombin 32 U/ml

TTX	5	-92.1 ± 1.4	162.9 ± 7.6	195.7 ± 5.5	205 ± 19
TTX & Thrombin	5	-90.9 ± 1.3	163.9 ± 11.3	195.1 ± 10.2	206 ± 28

SFLLR-NH₂ 100 μM

TTX	4	-91.6 ± 1.1	161.4 ± 16.4	192.9 ± 16.1	165 ± 8
TTX & SFLLR-NH ₂	4	-92.1 ± 0.8	166.7 ± 16.1	198.4 ± 16.3	158 ± 10

Guinea pig right ventricular papillary muscle action potential parameters determined in absence (A) and presence of TTX (B). In both procedures, parameters were determined at least 15 minutes following the applications of thrombin or SFLLR-NH₂. RP: resting membrane potential; APD₅₀ and APD₉₀: action potential duration measured respectively at 50 and 90 % of repolarisation. (dV/dt)_{max}: maximum upstroke velocity. Data are means ± SEM. n: number of tested animals. *: p < 0.05.

Figure 1

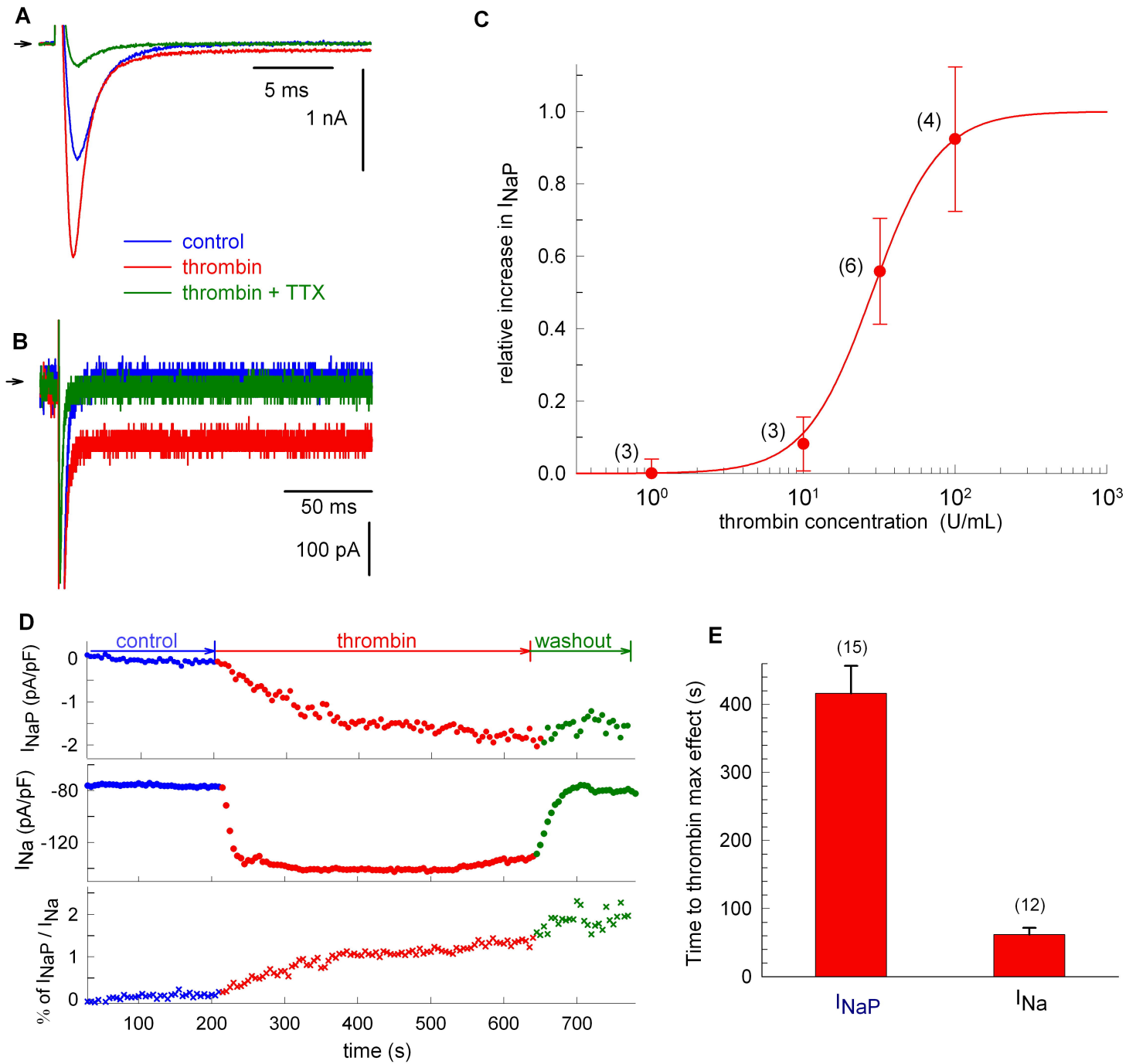


Figure 2

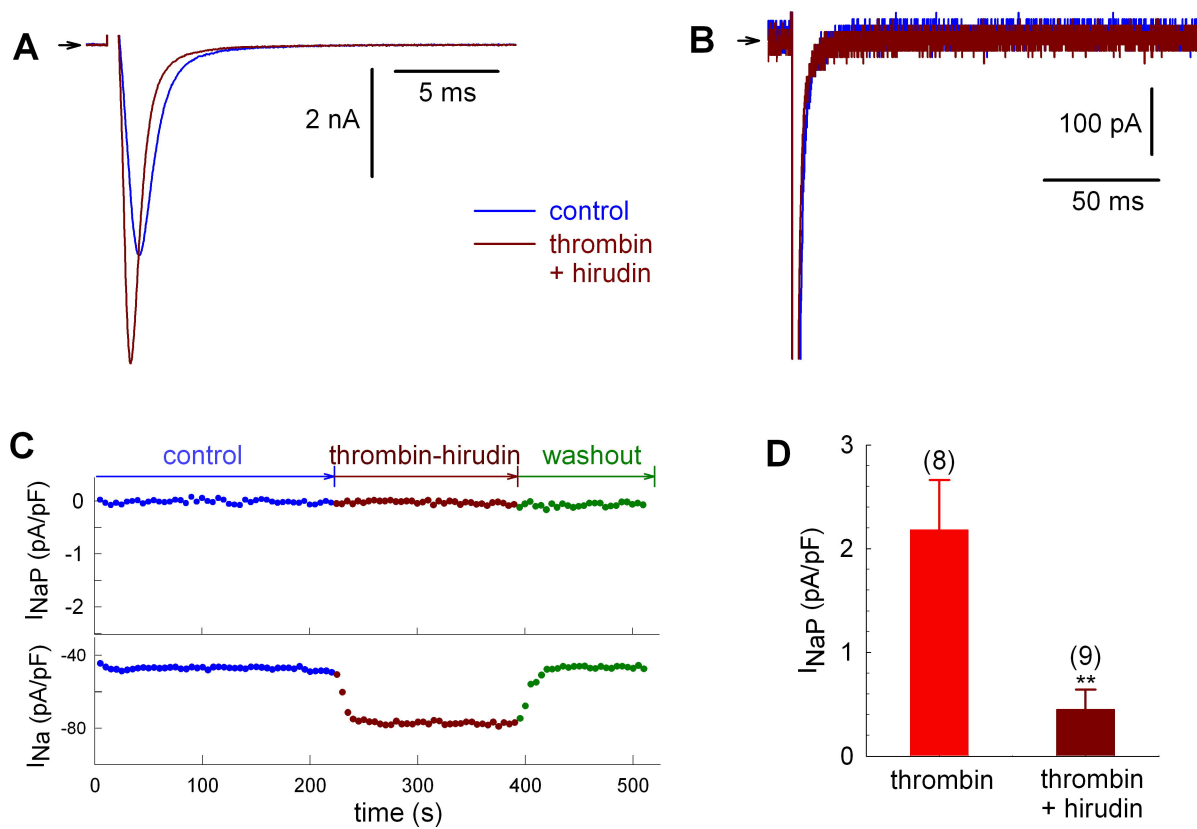


Figure 3

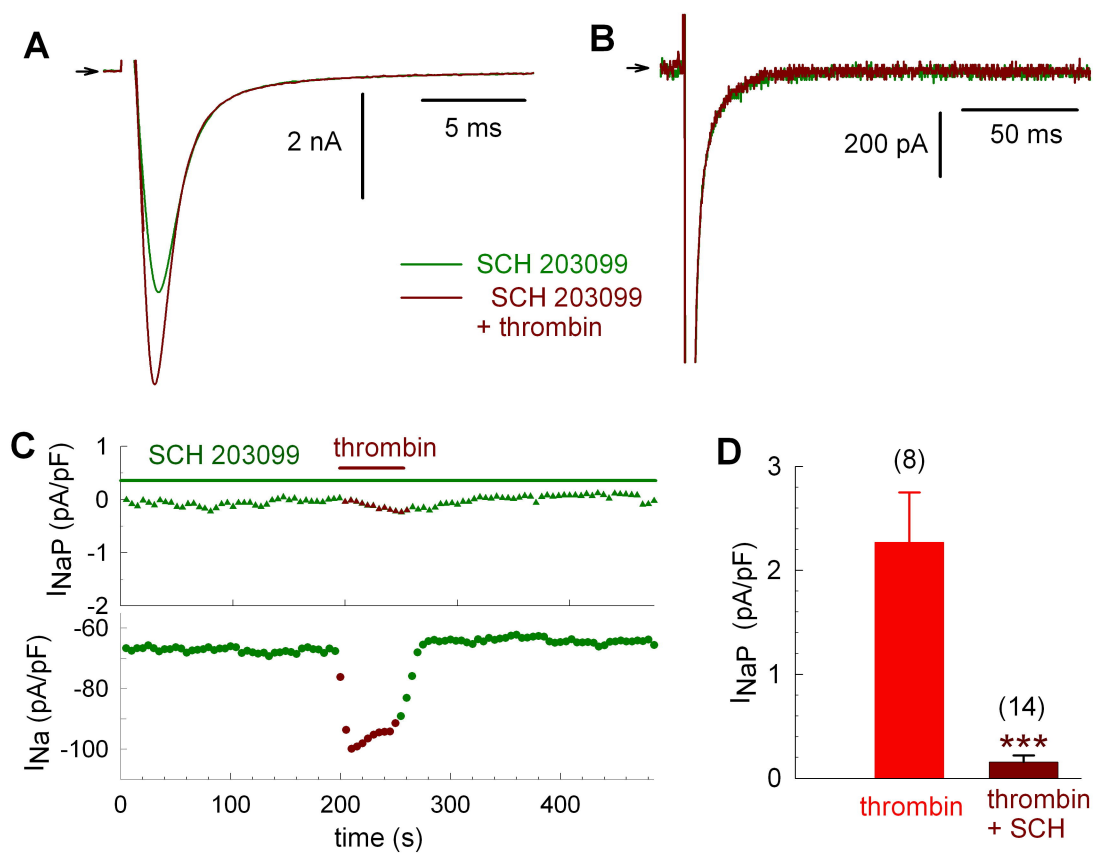


Figure 4

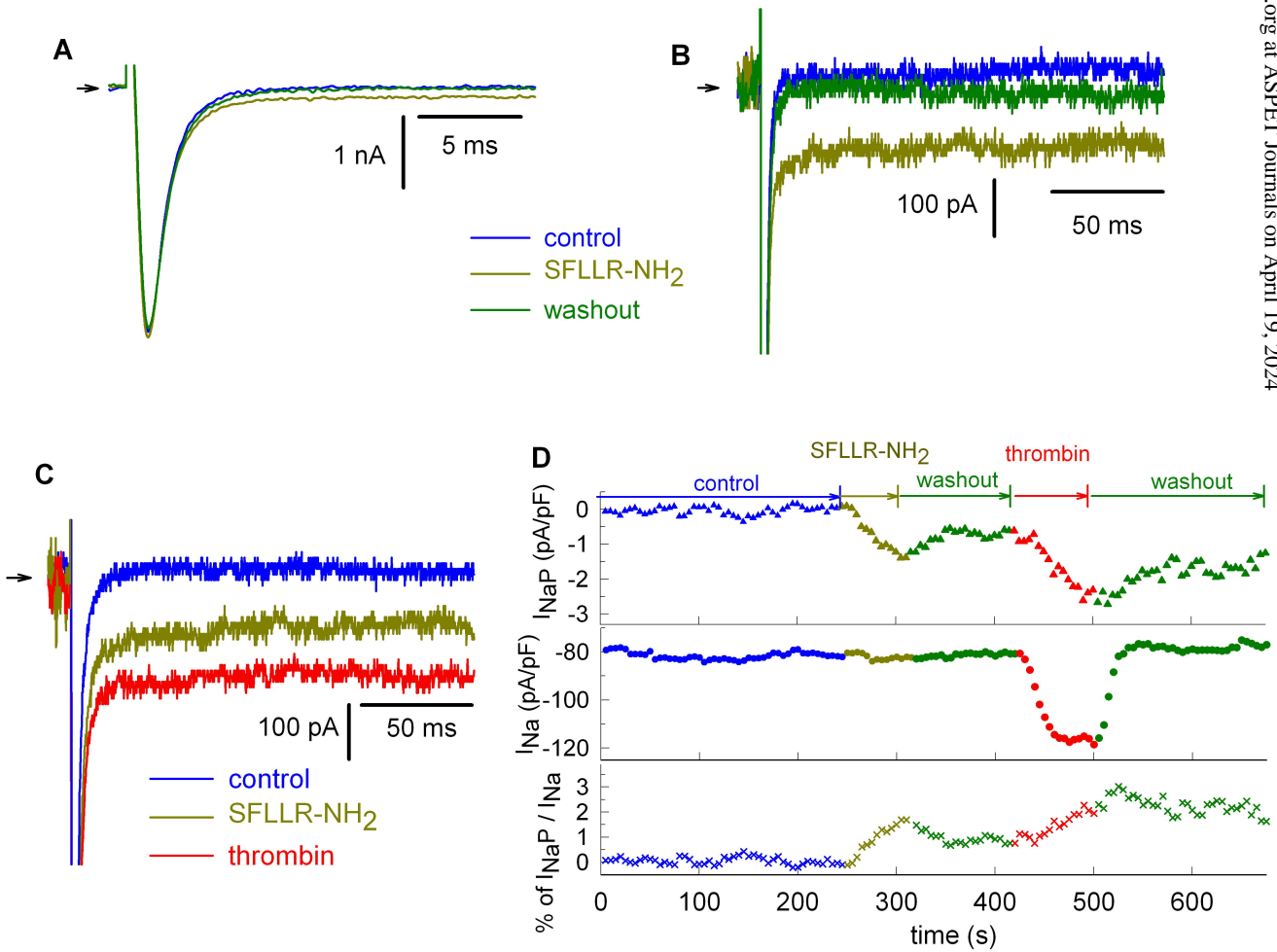


Figure 5

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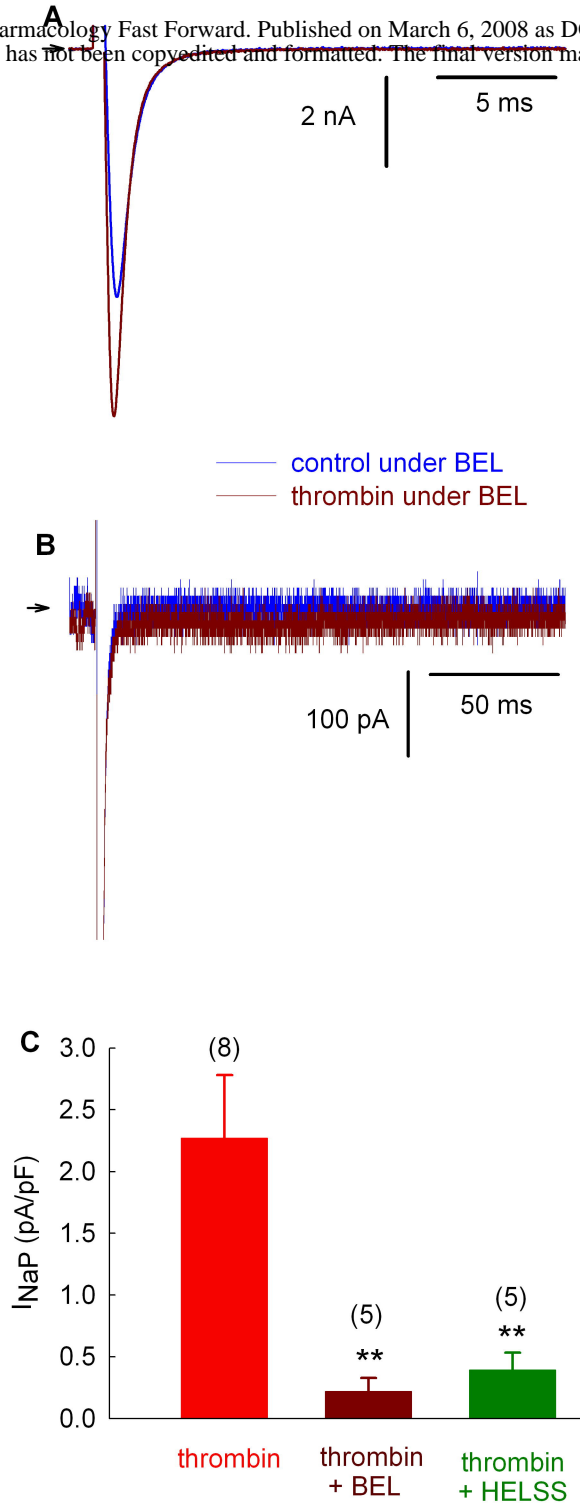


Figure 6A

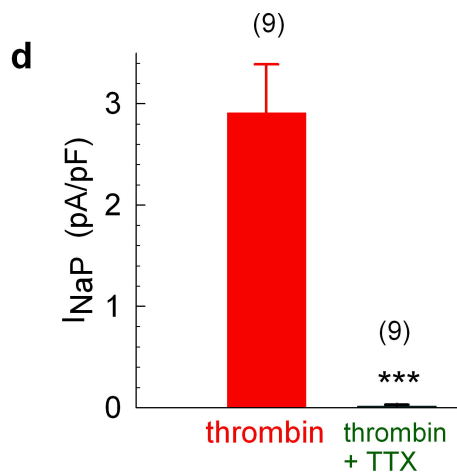
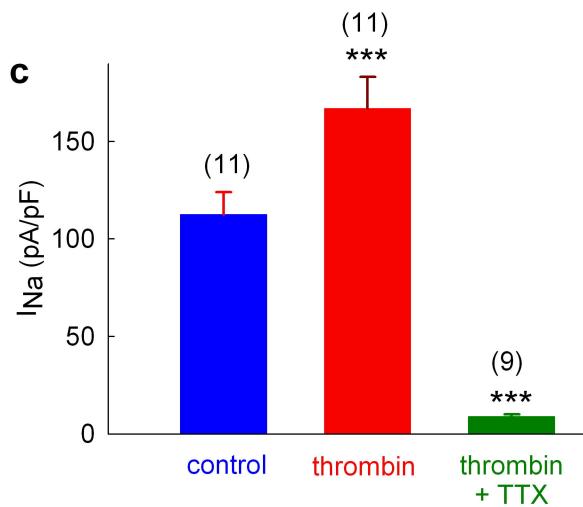
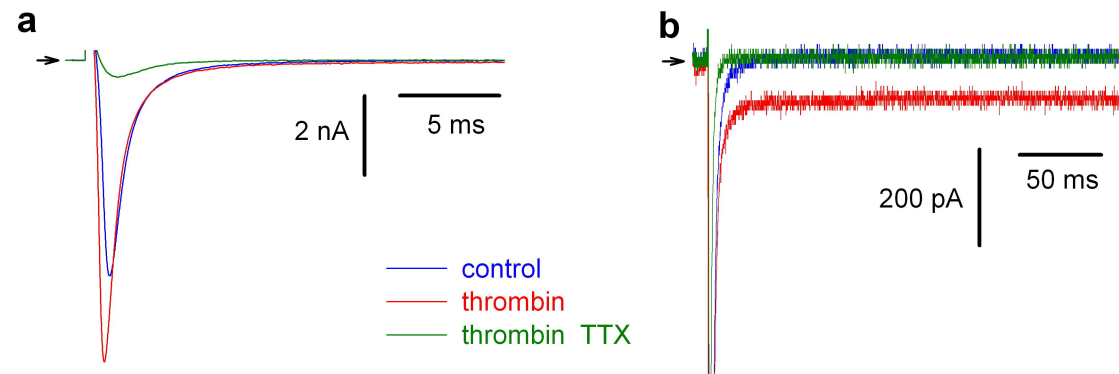


Figure 6 B

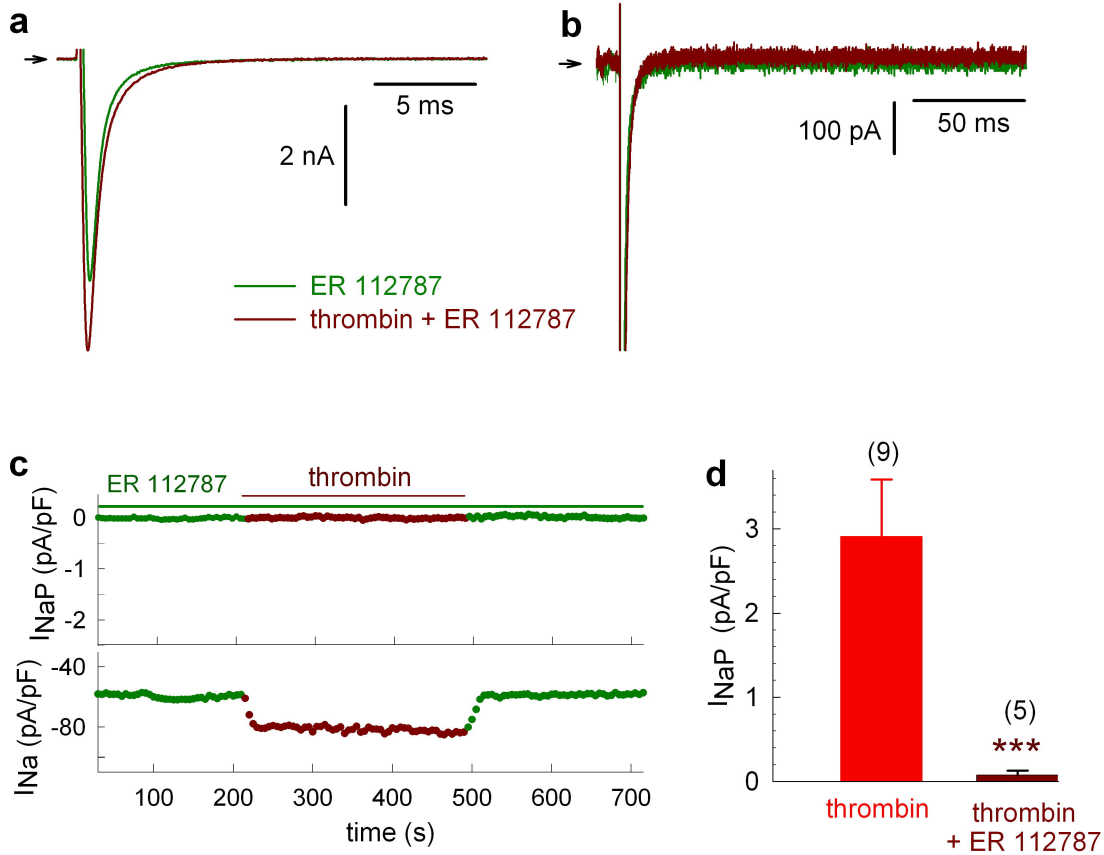


Figure 7A

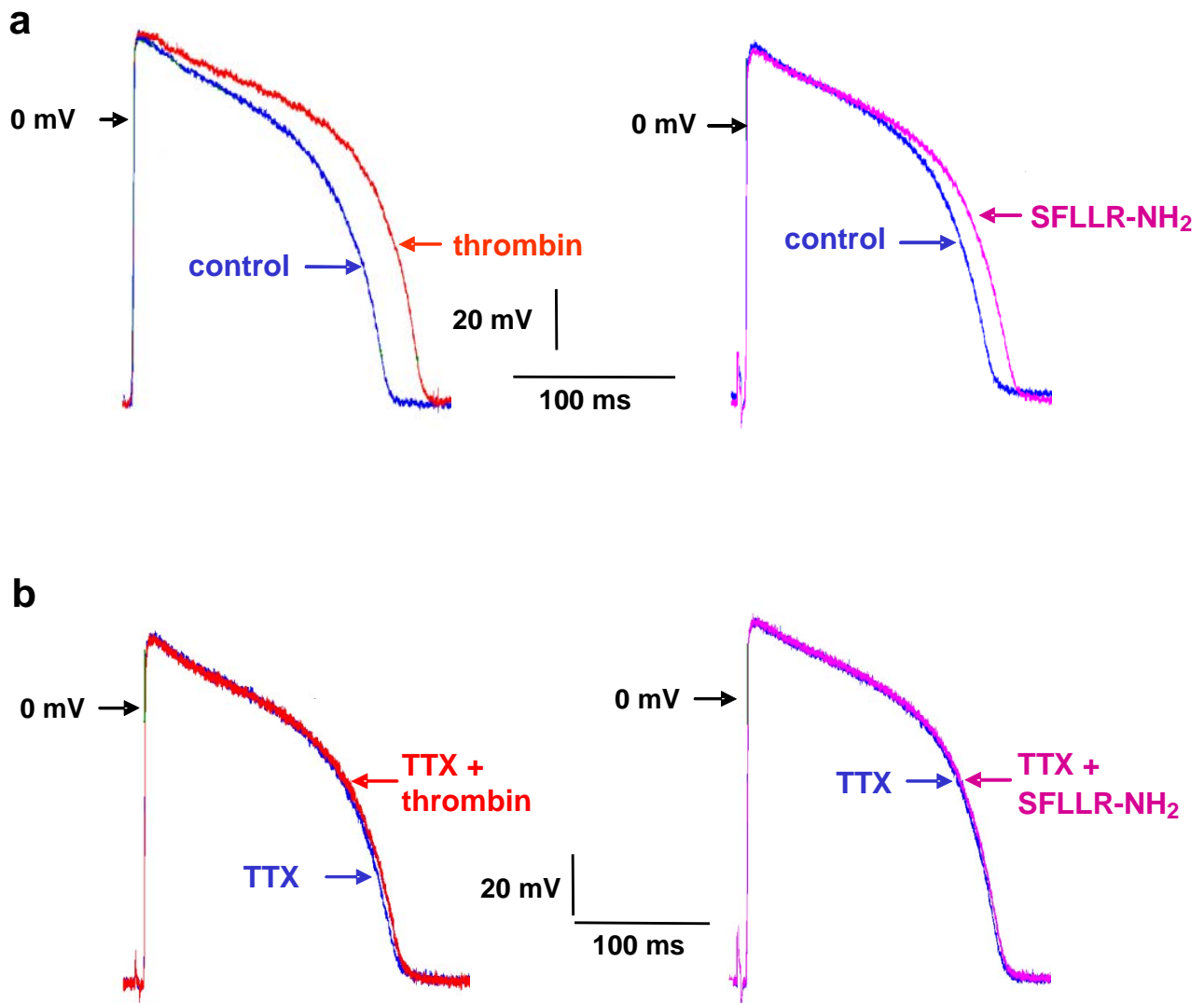


Figure 7B

