

RESEARCH PAPER

Ca²⁺ signalling by P2Y receptors in cultured rat aortic smooth muscle cells

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Background and purpose: P2Y receptors evoke Ca²⁺ signals in vascular smooth muscle cells and regulate contraction and proliferation, but the roles of the different P2Y receptor subtypes are incompletely resolved.

Experimental approach: Quantitative PCR was used to define expression of mRNA encoding P2Y receptor subtypes in freshly isolated and cultured rat aortic smooth muscle cells (ASMC). Fluorescent indicators in combination with selective ligands were used to measure the changes in cytosolic free [Ca²⁺] in cultured ASMC evoked by each P2Y receptor subtype.

Key results: The mRNA for all rat P2Y receptor subtypes are expressed at various levels in cultured ASMC. Four P2Y receptor subtypes (P2Y1, P2Y2, P2Y4 and P2Y6) evoke Ca²⁺ signals that require activation of phospholipase C and comprise both release of Ca²⁺ from stores and Ca²⁺ entry across the plasma membrane.

Conclusions and implications: Combining analysis of P2Y receptor expression with functional analyses using selective agonists and antagonists, we isolated the Ca²⁺ signals evoked in ASMC by activation of P2Y1, P2Y2, P2Y4 and P2Y6 receptors. *British Journal of Pharmacology* (2010) **160**, 1953–1962; doi:10.1111/j.1476-5381.2010.00763.x

Keywords: vascular smooth muscle; ATP; ADP; UTP; P2Y receptor; Ca²⁺ signal

Abbreviations: [Ca²⁺]_i, cytosolic free Ca²⁺ concentration; 2'-amino-UTP, 2'-amino-2'-deoxyuridine-5'-triphosphate; 2'-azido-UTP, 2'-azido-2'-deoxyuridine-5'-triphosphate; 2-MeSADP, 2-methylthio-adenosine-5'-diphosphate; 2-thio-UTP, 2-thio-uridine-5'-triphosphate; αβ-meATP, αβ-methylene-adenosine-5'-triphosphate; ASMC, aortic smooth muscle cell; DMEM, Dulbecco's modified Eagle's medium; HBSS, Hank's balanced salt solution; MRS2211, 2-[(2-chloro-5-nitrophenyl)azo]-5-hydroxy-6-methyl-3-[(phosphonoxy)methyl]-4-pyridinecarboxaldehyde; MRS2279, 2-chloro-N⁶-methyl-(N)-methanocarba-2'-deoxyadenosine-3',5'-bisphosphate; MRS2365, (N)-methanocarba-2-MeSADP; MRS2395, 2,2-dimethyl-propionic acid 3-(2-chloro-6-methylaminopurin-9-yl)-2-(2,2-dimethyl-propionyloxymethyl)-propyl ester; MRS2578, N,N'-1,4-butanediylbis [N'-(3-isothiocyanatophenyl)thio] urea; MRS2690, diphosphoric acid 1-α-D-glucopyranosyl ester 2-[(4'-methylthio)uridin-5''-yl] ester; PBS, phosphate-buffered saline; U73122, 1-[6-[(17β)-3-methoxyestra-1,3,5(10)-trien-17-yl] amino] hexyl]-1H-pyrrole-2,5-dione; U73343, 1-[6-[(17β)-3-methoxyestra-1,3,5(10)-trien-17-yl]amino] hexyl]-2,5-pyrrolidinedione; UDP-Glc, uridine-5'-diphospho-α-D-glucose

Introduction

P2Y receptors comprise a family of G protein-coupled receptors that are activated by adenine and uridine nucleotides, many of which evoke increases in cytosolic Ca²⁺ concentration (Abbraccio *et al.*, 2006) (Figure 1). The seven P2Y receptor subtypes known to be expressed in rats have overlapping ligand-recognition properties; they differ in the G proteins that they preferentially activate, and in whether they regulate phospholipase C, adenylyl cyclase or both. This complexity

and the limited range of subtype-selective antagonists make it difficult to discriminate the role of each P2Y receptor subtype in native tissues. In vascular smooth muscle, for example, many functional responses are regulated by the Ca²⁺ signals evoked by purinoceptors (Erlinge, 1998; Burnstock, 2002; Hou *et al.*, 2002; Vial and Evans, 2002). These receptors are activated by extracellular nucleotides released by many different cells, including activated platelets, and they can be further metabolized by extracellular nucleotidases (Abbraccio *et al.*, 2006). The Ca²⁺ signals regulate contraction (Horowitz *et al.*, 1996) and thereby contribute to control of blood pressure. They also regulate proliferation and are thereby important in mediating responses to vascular injury during restenosis (Inoue and Node, 2009) and atherosclerosis (Owens *et al.*, 2004). Analyses of the roles of specific P2Y receptors in evoking Ca²⁺ signals in vascular smooth muscle has, however, been hampered by the lack of selective ligands (Harper *et al.*,

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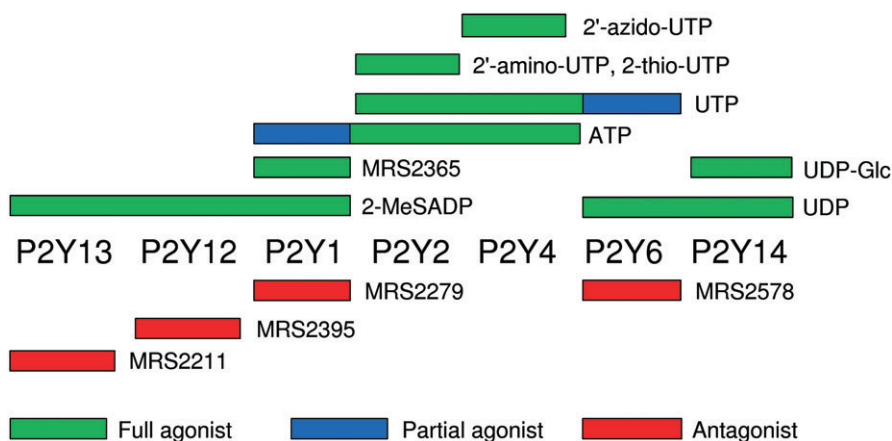


Figure 1 Selectivity of ligands for the P2Y receptors expressed in rat cultured ASMC. (Abbraccio *et al.*, 2006). The structures of the ligands are shown in Figure S1. 2'-amino-UTP, 2'-amino-2'-deoxyuridine-5'-triphosphate; 2'-azido-UTP, 2'-azido-2'-deoxyuridine-5'-triphosphate; 2-MeSADP, 2-methylthio-adenosine-5'-diphosphate; 2-thio-UTP, 2-thio-uridine-5'-triphosphate; MRS2211, 2-[(2-chloro-5-nitrophenyl)azo]-5-hydroxy-6-methyl-3-[(phosphonoxy)methyl]-4-pyridinecarboxaldehyde; MRS2279, 2-chloro-N⁶-methyl-(N)-methanocarpa-2'-deoxyadenosine-3',5'-bisphosphate; MRS2365, (N)-methanocarpa-2-MeSADP; MRS2395, 2,2-dimethyl-propionic acid 3-(2-chloro-6-methylaminopurin-9-yl)-2-(2,2-dimethyl-propionyloxymethyl)-propyl ester; MRS2578, N,N'-1,4-butanediylbis [N'-(3-isothiocyanatophenyl)thio] urea; UDP-Glc, uridine-5'-diphospho- α -D-glucose.

1998; Pediani *et al.*, 1999; Kumari *et al.*, 2003). It is important to resolve the roles of each P2Y receptor subtype in mediating the Ca²⁺ signals evoked by nucleotides and so perhaps to identify receptors that might selectively contribute to control of contractility or proliferation.

In culture, vascular smooth muscle cells develop a proliferative phenotype that is more reminiscent of cells contributing to vascular remodelling than the contractile phenotype of fully differentiated cells (Schwartz *et al.*, 1986). Although cultured cells are not contractile, they continue to express several vascular smooth muscle-specific proteins, including smooth muscle α -actin (Chamley-Campbell *et al.*, 1979), and in common with intact tissues they express a variety of functional purinoceptors (Erlinge, 1998; Hou *et al.*, 2002; Vial and Evans, 2002; Burnstock, 2009). Cultured vascular smooth muscle cells are thus a useful model for the proliferative state and they afford opportunities for quantitative analysis of the Ca²⁺ signals evoked by different purinoceptors.

Methods

Isolation and culture of rat aortic smooth muscle cells

All animal care and experimental procedures complied with UK Home Office policy. Smooth muscle cells were isolated from the aortas of adult male Wistar rats (250–300 g) by enzymatic digestion (Orallo, 1997). Briefly, rats were humanely killed by cervical dislocation. The aorta was isolated under sterile conditions in Hank's balanced salt solution (HBSS) supplemented with penicillin (100 U·mL⁻¹) and streptomycin (100 μ g·mL⁻¹), and cleared of adhering fat, connective tissue and the inner endothelial layer. After the first digestion (37°C, 12 min) in HBSS supplemented with antibiotics and collagenase B (0.015 units·mL⁻¹), the adventitia were removed. After a second digestion (37°C, 90 min) in HBSS containing antibiotics, collagenase B (0.075 U·mL⁻¹) and elastase (8 U·mL⁻¹), single cells were dispersed by trituration

with a sterile Pasteur pipette. The cells were washed with Dulbecco's modified Eagle's medium (DMEM) (650 g, 5 min), resuspended in DMEM with antibiotics and fetal calf serum (10%) and grown at 37°C in a humidified atmosphere of 95% air and 5% CO₂. The aortic smooth muscle cells (ASMC) were either passaged or frozen when they reached confluence. Cells were used between passages 2 and 8, throughout which they retained the appearance of smooth muscle cells and the characteristic 'hill and valley' growth pattern (Chamley-Campbell *et al.*, 1979) and, assessed by immunostaining, they expressed smooth muscle α -actin (Figure 2A and B).

Quantitative PCR

Confluent cultures of cells in 96-well plates were lysed with Fastlane cell cDNA kit (50 μ L·well⁻¹, Qiagen, Crawley, West Sussex, UK), and cDNA was then synthesized from the extract (4 μ L) in a final volume of 20 μ L and diluted 10-fold in water. For quantitative PCR, each reaction included primers specific for P2Y receptor subtypes or, for calibration, primers for β -actin. The primers used are listed in Table S1. Each reaction (15 μ L) included cDNA (0.5 μ L), P2Y receptor-specific primers (Quantitect Primer Assay, Qiagen, 1.5 μ L) or β -actin primers (1.5 μ L, 500 nM final concentration), and Bioline Sensimix (Bioline, London, UK, 7.5 μ L). For PCR (Rotor-Gene 6000, Qiagen), an initial denaturation (95°C, 10 min) was followed by 40 cycles of amplification (95°C for 10 s, 60°C for 30 s and 72°C for 20 s), with fluorescence measured at the end of each cycle. For each PCR, amplification efficiency (E) was calculated as 10^m , where m is the average increase in fluorescence for four cycles after the cycle threshold (C_T). Expression levels relative to the housekeeping product (β -actin) were calculated from: relative expression = $E^{-C_T(P2Y)}/E^{-C_T(\beta\text{-actin})}$. Each QPCR reaction was performed in duplicate with samples from three different wells. The authenticity of each PCR product was assessed by melting-curve analysis (72°C for 2 min, then 1°C steps at 5 s intervals to 99°C). Each reaction produced a single

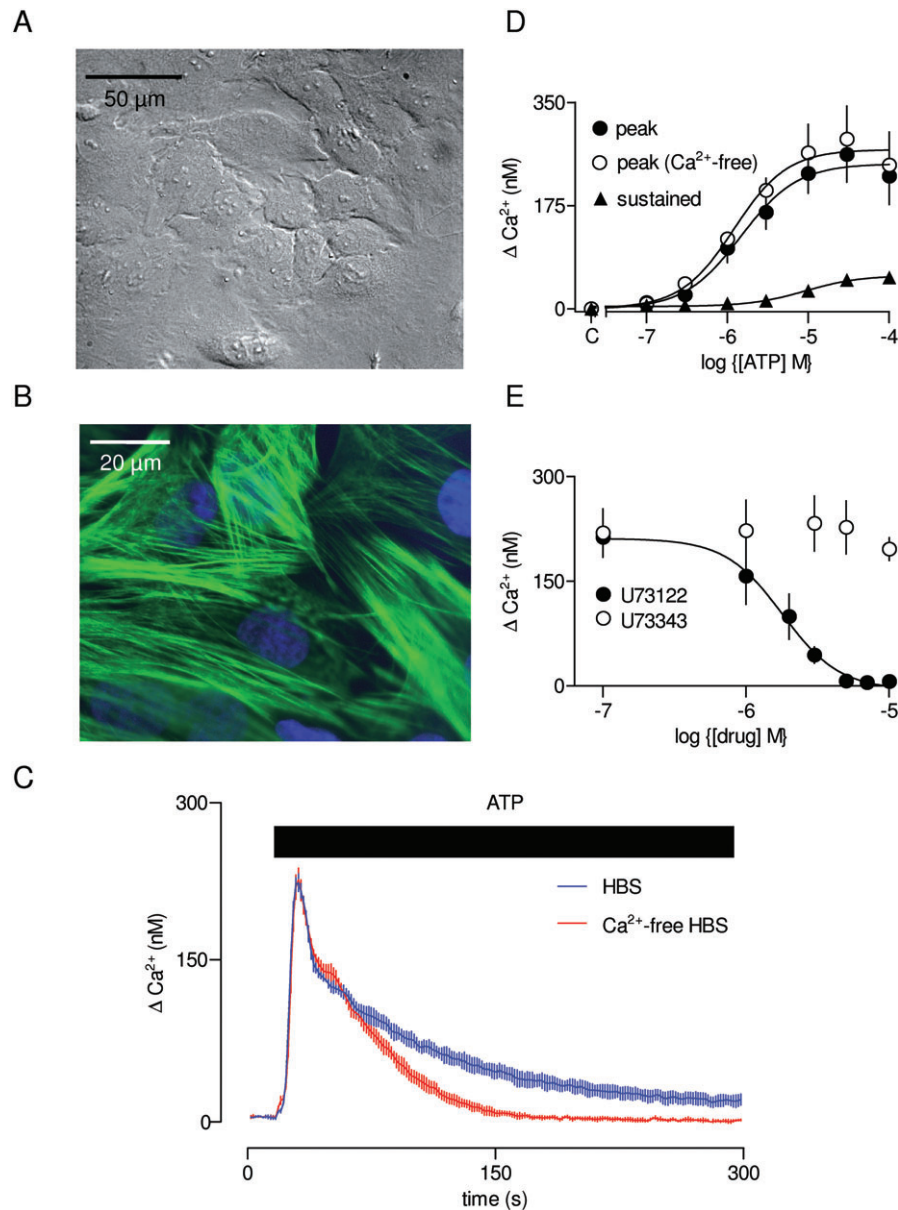


Figure 2 Ca²⁺ signals evoked by ATP in cultured ASMC. Typical field of cultured rat ASMC at passage 5 (A). Field of cells immunostained for smooth muscle α -actin (B). Ca²⁺ signals recorded from a population of fluo-4-loaded ASMC stimulated with ATP (100 μ M, bar) in the presence or absence of extracellular Ca²⁺ (C). Results are means \pm SEM from three wells on a single plate and are typical of results from three independent plates. Concentration-dependent effects of ATP on the peak increase in [Ca²⁺]_i in the presence or absence of extracellular Ca²⁺, and on the sustained Ca²⁺ entry signal (measured 250–300 s after addition of ATP) (D). C denotes control. Peak responses to ATP (100 μ M) with the indicated concentrations of U73122 or U73343 (each added for 5 min before and then during the stimulation) (E). [Ca²⁺]_i, cytosolic free Ca²⁺ concentration; ASMC, aortic smooth muscle cell; HBS, HEPES-buffered saline; U73122, 1-[6-[[[(17 β)-3-methoxyestra-1,3,5 (10)-trien-17-yl] amino] hexyl]-1H-pyrrole-2,5-dione]; U73343, 1-[6-[[[(17 β)-3-methoxyestra-1,3,5 (10)-trien-17-yl)amino] hexyl]-2,5-pyrrolidinedione.

amplification product with a melting point above 80°C. Control reactions in which the template cDNA was omitted from the PCR reaction, or reverse-transcriptase omitted from the reverse transcription reaction, failed to produce any specific amplification products. Similar methods were used for analyses of cells freshly isolated from aorta (30 mg) immediately before the second enzyme digestion, as described above. For these cells, RNA was extracted using an Illustra RNA spin mini-RNA isolation kit (GE Healthcare, Little Chalfont, Buckinghamshire, UK), and cDNA was synthesized using Bioscript reverse transcriptase (Bioline).

Measurement of [Ca²⁺]_i in cell populations

Confluent cultures of cells grown in 96-well plates were loaded with fluo-4 by incubation in HEPES-buffered saline (HBS) supplemented with probenecid (2.5 mM) and fluo-4 acetoxyethyl ester [4 μ M in fresh dimethyl sulphoxide (DMSO)]; final DMSO concentration, 0.4%. After 1 h at 20°C, cells were washed with HBS and after 30 min (to allow complete de-esterification of fluo-4AM), they were used for experiments. HBS had the following composition: 135 mM NaCl, 5.9 mM KCl, 1.2 mM MgCl₂, 1.5 mM CaCl₂, 11.5 mM glucose and 11.6 mM HEPES, pH 7.3, at 20°C. All experiments were

Table 1 Ca²⁺ signals evoked by activation of P2Y receptors

	$\Delta[\text{Ca}^{2+}]_i$ (nM)		pEC_{50} (h)		
	Peak	Sustained	Peak (Ca ²⁺ -free)	Peak (with Ca ²⁺)	Ca ²⁺ entry
ATP	248 ± 48	64 ± 20	5.95 ± 0.10 (1.1 ± 0.2)	5.83 ± 0.03 (1.4 ± 0.2)	5.01 ± 0.13 (1.8 ± 0.4)
MRS2365	154 ± 5	18 ± 7	8.56 ± 0.06 (0.9 ± 0.1)	8.68 ± 0.16 (0.8 ± 0.2)	8.11 ± 0.32 (0.7 ± 0.2)
UTP	254 ± 47	67 ± 5	5.39 ± 0.15 (1.1 ± 0.1)	5.23 ± 0.22 (1.0 ± 0.2)	4.64 ± 0.32 (0.7 ± 0.2)
UDP	94 ± 20	25 ± 6	6.16 ± 0.43 (0.6 ± 0.2)	6.45 ± 0.45 (0.6 ± 0.2)	6.83 ± 0.35 (0.7 ± 0.1)
2'-amino-UTP	301 ± 34	84 ± 36	ND	5.58 ± 0.37 (1.1 ± 0.1)	5.40 ± 0.23 (1.1 ± 0.1)
2-thio-UTP	327 ± 42	88 ± 26	ND	5.89 ± 0.29 (1.1 ± 0.3)	5.11 ± 0.18 (1.1 ± 0.2)
2'-azido-UTP	142 ± 29	55 ± 20	ND	4.67 ± 0.32 (1.3 ± 0.1)	4.67 ± 0.06 (1.1 ± 0.1)
2-MeSADP	122 ± 30	18 ± 4	ND	7.23 ± 0.53 (0.8 ± 0.1)	5.58 ± 0.16 (1.8 ± 0.7)

The results in the Table show the peak and sustained increase (measured between 250 and 300 s) in [Ca²⁺]_i in populations of cultured ASMC stimulated with the indicated ligands. pEC_{50} values with Hill coefficients (*h*) are shown for the responses in HBS and Ca²⁺-free HBS, and for the Ca²⁺ entry. [Ca²⁺]_i, cytosolic free Ca²⁺ concentration; 2'-amino-UTP, 2'-amino-2'-deoxyuridine-5'-triphosphate; 2'-azido-UTP, 2'-azido-2'-deoxyuridine-5'-triphosphate; 2-MeSADP, 2-methylthio-adenosine-5'-diphosphate; 2-thio-UTP, 2-thio-uridine-5'-triphosphate; HBS, HEPES-buffered saline; MRS2365, (N)-methanocarba-2-MeSADP; ND, not determined.

performed in HBS (or nominally Ca²⁺-free HBS) at 20°C. For fluorescence measurements, the 96-well plate containing fluo-4-loaded cells was mounted in a FlexStation III (MDS Analytical Technologies, Wokingham, Berkshire, UK), which allows up to three automated additions to each well, while recording fluorescence (excitation at 485 nm; emission at 525 nm). Six simultaneous readings were taken from each well at intervals of 1.52 s. At the end of each experiment, fluorescence of Ca²⁺-saturated indicator (F_{max}) was determined for each well by addition of Triton X-100 (0.05%) and CaCl₂ (10 mM). Fluorescence from cells containing only Ca²⁺-free indicator (equivalent to background fluorescence because Ca²⁺-free fluo-4 is not fluorescent) was measured in parallel wells treated with Triton X-100 (0.05%) and BAPTA (10 mM). This background fluorescence was subtracted from all measurements. The corrected fluorescence (*F*) was then calibrated to cytosolic free Ca²⁺ concentration ([Ca²⁺]_i) from: [Ca²⁺]_i = $K_D F / (F_{\text{max}} - F)$, where the K_D of fluo-4 for Ca²⁺ is 345 nM (Gee *et al.*, 2000).

Immunostaining

Cells grown on poly-L-lysine-coated coverslips under conditions identical to those used for measurements of [Ca²⁺]_i were washed twice with phosphate-buffered saline (PBS: 137 mM NaCl, 2.7 mM KCl, 10.1 mM Na₂HPO₄ and 1.76 mM KH₂PO₄, pH 7.4) and fixed with paraformaldehyde (4%) in PBS for 20 min at 20°C. Cells were then permeabilized with Triton X-100 (0.1%) in PBS for 5 min. Non-specific binding was blocked by incubating with BSA (5%) in PBS for 10 min at 20°C, before staining with a FITC-conjugated monoclonal antibody to anti-smooth muscle α -actin (1:200) for 2.5 h at 20°C. Cells were then washed three times with PBS, and the nuclei stained with Hoechst 33258 (1 $\mu\text{g}\cdot\text{mL}^{-1}$). Coverslips were mounted in Prolong Gold (Invitrogen, Paisley, Scotland) and stored at -20°C before confocal imaging.

Statistical analysis

Results are presented as means ± SEM from *n* independent experiments (*n* = 3, unless otherwise stated). Statistical com-

parisons used paired or unpaired Student's *t*-test, as appropriate, with *P* < 0.05 considered significant (GraphPad Prism, version 5, La Jolla, CA, USA).

Materials

Collagenase B and hexokinase were from Roche Applied Science (Burgess Hill, West Sussex, UK). Elastase and FITC-conjugated α -actin antiserum were from Sigma (Poole, Dorset, UK). MRS2211 (2-[(2-chloro-5-nitrophenyl)azo]-5-hydroxy-6-methyl-3-[(phosphonooxymethyl)4-pyridinecarboxaldehyde), MRS2279 (2-chloro-N⁶-methyl-(N)-methanocarba-2'-deoxyadenosine-3',5'-bisphosphate), MRS2365 [(N)-methanocarba-2-MeSADP], MRS2578 (N,N''-1,4-butanediylbis [N'-(3-isothiocyanatophenyl)thio] urea), MRS2690 (diphosphoric acid 1- α -D-glucopyranosyl ester 2-[(4'-methylthio)uridin-5'-yl] ester), 2-MeSADP (2-methylthio-adenosine-5'-diphosphate) and UDP were from Tocris (Bristol, Avon, UK). Cell culture materials were from Invitrogen, except for fetal bovine serum (Sigma). 2'-amino-UTP (2'-amino-2'-deoxyuridine-5'-triphosphate), 2'-azido-UTP (2'-azido-2'-deoxyuridine-5'-triphosphate) and 2-thio-UTP (2-thio-uridine-5'-triphosphate) were from Trilink Biotechnologies (San Diego, CA, USA). The structures of the ligands are shown in Figure S1. Fluo-4AM was from Invitrogen. *Pertussis* toxin was from List Biological Laboratories (Campbell, CA, USA). Other reagents, including MRS2395 (2,2-dimethyl-propionic acid 3-(2-chloro-6-methylaminopurin-9-yl)-2-(2,2-dimethyl-propionyloxymethyl)-propyl ester), were from Sigma unless stated otherwise. The nomenclature of receptors and ligands follows Alexander *et al.* (2009).

Results

Ca²⁺ signals evoked by ATP in rat aortic smooth muscle cells

ATP evoked a concentration-dependent increase in [Ca²⁺]_i in populations of rat ASMC. Both the initial peak response and its sensitivity to ATP were unaffected by removal of extracellular Ca²⁺, but a small sustained Ca²⁺ signal was abolished in Ca²⁺-free medium (Figure 2C and D; Table 1). There was

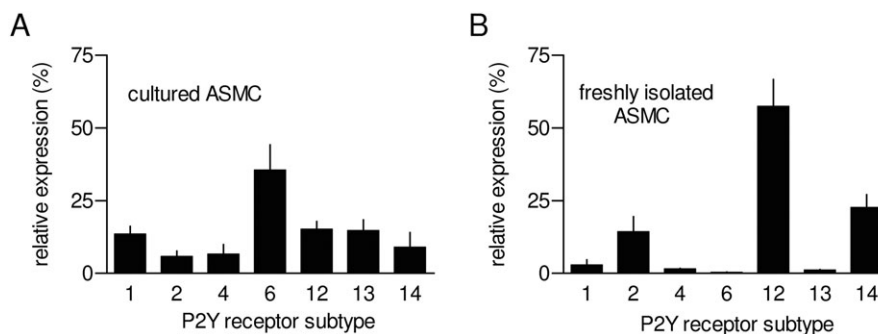


Figure 3 Expression of P2Y receptor subtypes in aortic smooth muscle cell (ASMC). Quantitative PCR was used to measure relative levels of mRNA encoding the subtypes of P2Y receptors in either cultures of ASMC (A, passage 3–7) or in freshly isolated ASMC (B). Results, expressed as percentages of total mRNA encoding P2Y receptors, are means \pm SEM from three experiments, each performed in duplicate.

no response to $\alpha\beta$ -meATP ($\alpha\beta$ -methylene-adenosine-5'-triphosphate), a selective agonist of P2X1 and P2X3 receptors (not shown); the former, with P2X7 receptors, predominate in vascular smooth muscle (North, 2002). The lack of responses mediated by P2X receptors is consistent with previous studies indicating that P2X receptors are down-regulated in cultured vascular smooth muscle cells (Erlinge *et al.*, 1998). The Ca²⁺ signals and their sensitivity to ATP were unaffected by treatment of the ATP and cells with an ATP-regenerating system (2 U·mL⁻¹, creatine phosphokinase, 1 mM phosphocreatine and 10 mM LiCl in HBS for 10 min) (Figure S2A). This confirms that the Ca²⁺ signals are evoked by ATP rather than by contaminating ADP. U73122 (1-[6-[[[(17 β)-3-methoxyestra-1,3,5 (10)-trien-17-yl] amino] hexyl]-1H-pyrrole-2,5-dione), an inhibitor of phospholipase C (Bleasdale *et al.*, 1990), caused a concentration-dependent and complete inhibition (pIC₅₀ = 5.78 \pm 0.07) of the Ca²⁺ signals evoked by ATP; the inactive analogue, U73343 (1-[6-[[[(17 β)-3-methoxyestra-1,3,5 (10)-trien-17-yl]amino] hexyl]-2,5-pyrrolidinedione), had no effect (Figure 2E). These results demonstrate that activation of P2Y receptors by ATP in rat ASMC evokes both Ca²⁺ release and Ca²⁺ entry via activation of phospholipase C.

Expression of P2Y receptor subtypes in rat aortic smooth muscle cells

ATP is an agonist of several subtypes of P2Y receptor (Abbraccio *et al.*, 2006) (Figure 1). We therefore used PCR to quantify the transcripts for all known rat P2Y receptor subtypes in cultures of ASMC before attempting to delineate the subtypes responsible for ATP-evoked Ca²⁺ signals. The results demonstrate the presence of transcripts for seven different P2Y receptor subtypes, including three for which ATP is an agonist (Figure 3A). These results establish which P2Y receptors may be expressed in rat ASMC and thereby dictate our choice of ligands to resolve, in functional assays, the receptor subtypes that mediate Ca²⁺ signals (Figure 1). Similar analyses of freshly isolated ASMC established a different pattern of expression with mRNA for only three P2Y receptors predominating (P2Y2, P2Y12 and P2Y14) (Figure 3B). It is noteworthy that the predominant P2Y receptor subtype in cultured ASMC is P2Y6, which is involved in regulating proliferation (Hou *et al.*, 2002), whereas P2Y12 is the major subtype in freshly isolated ASMC (Figure 3). The substantial effects of cell culture on the

expression of mRNA for P2Y6 receptors differ from a previous study of a more limited range of P2Y receptors, in which relative levels of expression of the P2Y receptor subtypes were similar in fresh and cultured ASMC (Erlinge *et al.*, 1998). Here, we focus on analyses of cultured ASMC because they are more amenable than primary cells to quantitative analyses of the Ca²⁺ signals evoked by different P2Y receptors.

Some of the P2Y receptor subtypes identified by PCR analysis are activated by UTP and/or UDP, rather than ATP (Figure 1). We therefore examined the effects of UTP and UDP on Ca²⁺ signals in ASMC. Responses to UTP were unaffected by treatment with a regenerating system (as described above) (Figure S2B), but responses to the highest concentrations of UDP (>10 μ M) were reduced after the UDP was treated (1 h, 37°C) with hexokinase (50 U·mL⁻¹, pH 7.0 in 0.1 M phosphate buffer with 6.5 mM MgCl₂) and glucose (110 mM) (Figure 4A). This indicated that the UDP was contaminated with UTP (Harden *et al.*, 1997; Kumari *et al.*, 2003). We therefore treated UDP with glucose/hexokinase before all assays. Both UTP and UDP evoked Ca²⁺ release and Ca²⁺ entry (Figure 4B and C; Table 1). These responses, like those to ATP, were abolished by inhibition of phospholipase C with U73122 (not shown). The peak increases in [Ca²⁺]_i were similar for ATP and UTP, but clearly smaller for UDP (Table 1). The results so far established that ATP, UTP and UDP, via P2Y receptors, evoked both Ca²⁺ release and Ca²⁺ entry via their ability to activate phospholipase C in rat cultured ASMC.

Activation of P2Y1 receptors evokes Ca²⁺ signals

MRS2365 is a selective full agonist of P2Y1 receptors (Chhatriwala *et al.*, 2004), while MRS2279 is a selective high-affinity ($K_D \sim 8$ nM) competitive antagonist (Boyer *et al.*, 2002; Waldo *et al.*, 2002). MRS2365 evoked concentration-dependent increases in [Ca²⁺]_i (pEC₅₀ is shown in Table 1) arising from both release of Ca²⁺ from intracellular stores and Ca²⁺ entry (Figure 5A). Increasing concentrations of MRS2279 caused rightward parallel shifts in the concentration–effect relationships for the Ca²⁺ signals evoked by MRS2365 (Figure 5B). The pK_D of MRS2279 for the receptor through which MRS2365 evokes Ca²⁺ signals was 8.16 \pm 0.26 (Figure 5C). These results and the K_D for MRS2279 are consistent with expression of functional P2Y1 receptors in rat cultured ASMC.

2-MeSADP is a full agonist for three of the P2Y receptor subtypes identified in ASMC by PCR (P2Y1, P2Y12 and P2Y13)

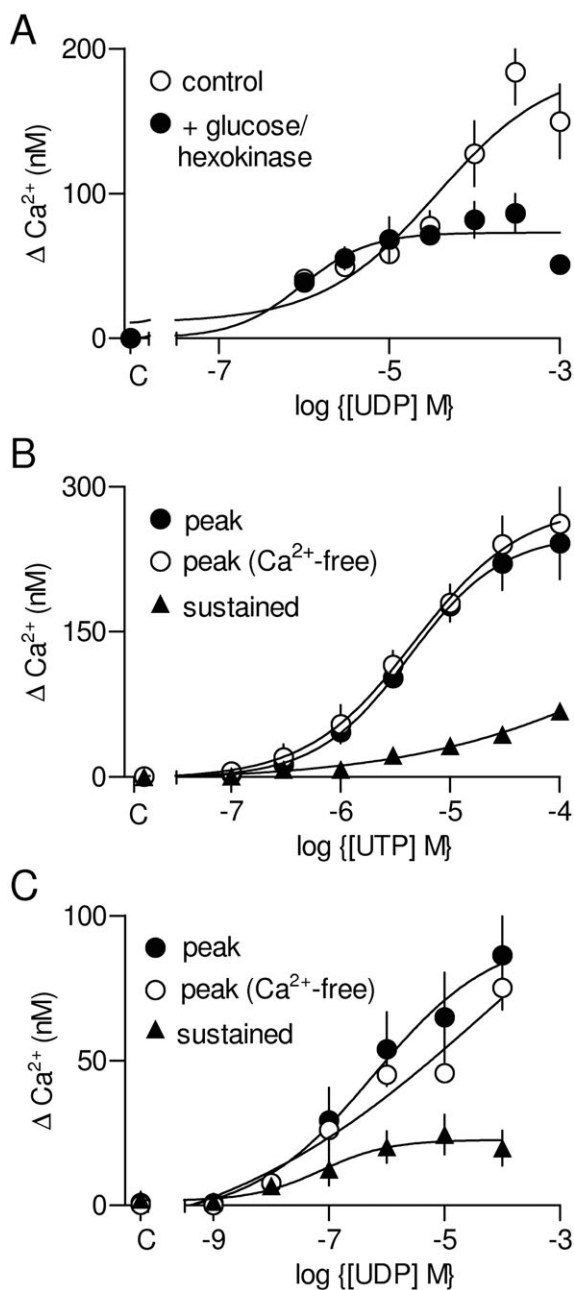


Figure 4 Ca²⁺ signals evoked by UTP and UDP. Concentration-dependent effects of UDP on the peak Ca²⁺ signal with or without prior treatment of the UDP with glucose and hexokinase (see text for details) (A). These results (A) are shown as means \pm SEM from a single experiment with three replicates. Concentration-dependent effects of UTP (B) or glucose/hexokinase-treated UDP (C) on the peak increase in [Ca²⁺]_i (cytosolic free Ca²⁺ concentration) in the presence or absence of extracellular Ca²⁺, and on the sustained Ca²⁺ entry signal. C denotes control.

(Figure 1). 2-MeSADP caused a concentration-dependent increase in [Ca²⁺]_i (Figure 5D; Table 1). The response to 2-MeSADP was competitively antagonized by MRS2279 with an affinity ($pK_D = 8.64 \pm 0.26$; Figure 5D and E) not significantly different from that for its antagonism of responses to the P2Y1-selective ligand, MRS2365 ($pK_D = 8.16 \pm 0.26$). Because 2-MeSADP interacts with three P2Y receptor sub-

types, but MRS2279 binds only to P2Y1 receptors (Boyer *et al.*, 2002; Waldo *et al.*, 2002; Marteau *et al.*, 2003), these results established that the Ca²⁺ signals evoked by 2-MeSADP were entirely mediated by P2Y1 receptors. That conclusion was further substantiated by results with MRS2395 (30 μ M) and MRS2211 (10 μ M), selective antagonists of P2Y12 and P2Y13 receptors, respectively (Xu *et al.*, 2002; Kim *et al.*, 2005), neither of which affected the Ca²⁺ signals evoked by any concentration of 2-MeSADP (Figure S3). A concentration of MRS2279 (3 μ M) sufficient to abolish responses to a maximally effective concentration of MRS2365 had no effect on the Ca²⁺ signals evoked by any concentration of UTP (Figure 5F) or ATP (Figure S4). This is consistent with ATP being only a weak partial agonist of P2Y1 receptors (Palmer *et al.*, 1998) and with the lack of interaction between UTP and P2Y1 receptors (Waldo *et al.*, 2002). We conclude that of the three P2Y receptors expressed in ASMC that were stimulated by 2-MeSADP (P2Y1, P2Y12 and P2Y13), only the P2Y1 receptors evoke Ca²⁺ signals. The peak increase in [Ca²⁺]_i evoked by maximal activation of P2Y1 receptors was only $62 \pm 2\%$ of that evoked by ATP (Table 1).

ATP evokes Ca²⁺ signals by activating P2Y2 and P2Y4 receptors

The only remaining P2Y receptors expressed in cultured ASMC that are activated by ATP are P2Y2 and P2Y4 receptors (Figure 1). ATP and UTP are full agonists for both rat receptors, and each ligand has similar potency for each receptor (Wildman *et al.*, 2003). The peak amplitudes of the Ca²⁺ signals evoked by ATP and UTP in ASMC were similar (Table 1), consistent with each agonist similarly activating the same population of receptors. UTP analogues have been used to differentiate between human P2Y2 and P2Y4 receptors. 2'-amino-UTP and 2-thio-UTP are more selective for human P2Y2 receptors, while 2'-azido-UTP is more selective for human P2Y4 receptors (Jacobson *et al.*, 2006); each analogue is a full agonist for its respective receptor, but neither is entirely selective. There are no antagonists that discriminate between rat P2Y2 and P2Y4 receptors (Abbracchio *et al.*, 2006). Each of the three UTP analogues evoked both Ca²⁺ release and Ca²⁺ entry in ASMC (Figures 6A–C; Table 1). The analogues likely to be selective for P2Y2 receptors (2'-amino-UTP and 2-thio-UTP) evoked substantially larger Ca²⁺ signals than did the P2Y4-selective agonist (2'-azido-UTP) (Figure 6D). We noted that the peak Ca²⁺ signal evoked by activation of both P2Y2 and P2Y4 receptors by UTP (273 ± 40 nM) was no larger than that evoked by activation of only the P2Y2 receptor by 2'-amino-UTP (301 ± 34 nM) (Table 1). The likely explanation is that maximal responses to full agonists may be limited by the finite size of the inositol 1,4,5-trisphosphate (IP₃)-sensitive Ca²⁺ stores. Our results allow us to conclude that in cultured ASMC, both P2Y2 and P2Y4 receptors evoke Ca²⁺ signals, with the P2Y2 receptors contributing most to the responses evoked by ATP and UTP.

P2Y6 receptors evoke Ca²⁺ signals

From the seven P2Y receptor subtypes expressed in cultured ASMC (Figure 3A), we have so far shown that P2Y1 (Figure 5), P2Y2 and P2Y4 receptors (Figure 6) evoke Ca²⁺ signals, while

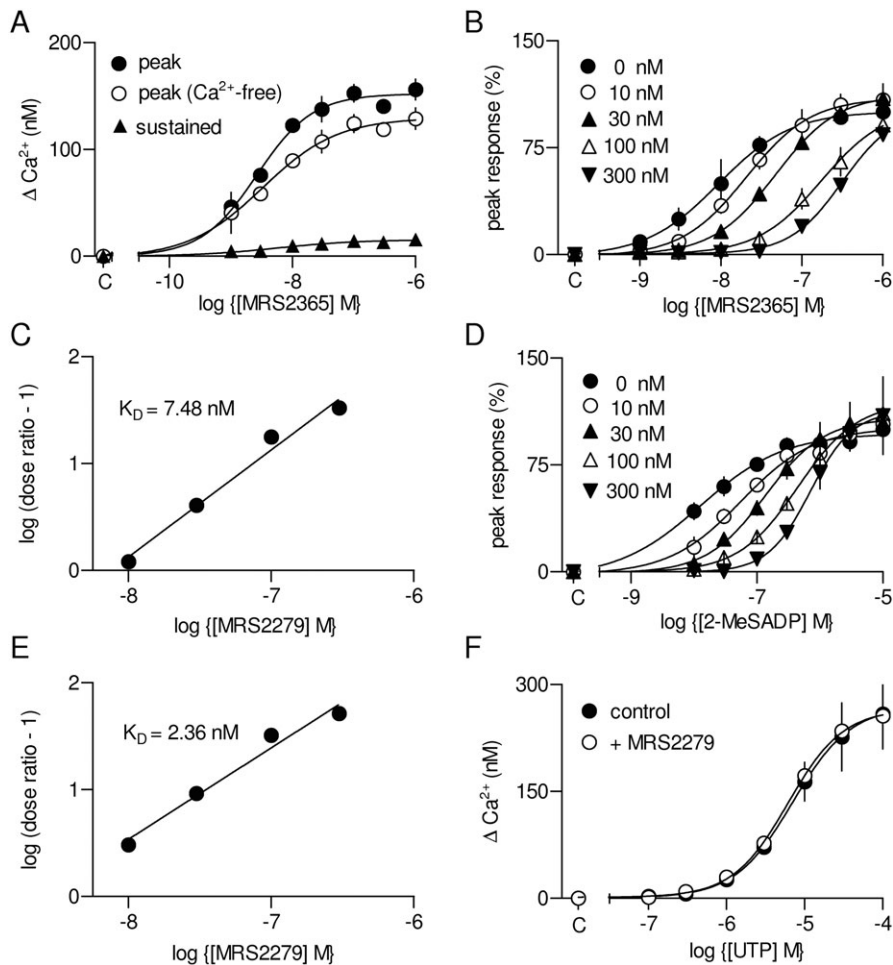


Figure 5 Ca²⁺ signals evoked by activation of P2Y1 receptors. Concentration-dependent effects of MRS2365 [(N)-methanocarba-2-MeSADP] on the peak Ca²⁺ signal in the presence or absence of extracellular Ca²⁺, and on the sustained Ca²⁺ entry (A). Concentration-dependent effects of MRS2365 (B) and 2-MeSADP (2-methylthio-adenosine-5'-diphosphate) (D) on the peak Ca²⁺ signal in the presence of 0–300 nM MRS2279 (2-chloro-N⁶-methyl-(N)-methanocarba-2'-deoxyadenosine-3',5'-bisphosphate) (as indicated). Results are plotted as percentages of the maximal response in the absence of antagonist. The results from (B and D) are shown as Schild plots (C and E). Calculated affinities of the antagonist (K_D) are shown. Concentration-dependent effects of UTP, alone or with MRS2279 (3 μ M), on the peak Ca²⁺ signal (F). C denotes control (in A, B, D and F).

P2Y12 and P2Y13 receptors do not (Figure 5). The physiological ligands of the receptors are likely to be ADP for P2Y1 receptors, and ATP or UTP for P2Y2 and P2Y4 receptors (Abbracchio *et al.*, 2006). The remaining subtypes of rat P2Y receptors (P2Y6 and P2Y14) are each activated by UDP (Nicholas *et al.*, 1996; Fricks *et al.*, 2008), which evokes smaller Ca²⁺ signals than either ATP or UTP in ASMC (Figure 4A). MRS2578 has been reported to inhibit rat P2Y6 receptors selectively and irreversibly (Mamedova *et al.*, 2004), although it is both sparingly soluble and unstable in aqueous solution (Abbracchio *et al.*, 2006). In our hands, MRS2578 (10 μ M) had no effect on the Ca²⁺ signals evoked by any concentration of UDP in ASMC (not shown). There are no selective antagonists of P2Y14 receptors, although MRS2690 (Ko *et al.*, 2007) and UDP-glucose (Chambers *et al.*, 2000) are selective agonists. Neither MRS2690 (10 μ M) nor UDP-glucose (10 μ M) evoked Ca²⁺ signals in cultured ASMC (not shown).

We attempted further to resolve the roles of P2Y6 and P2Y14 receptors by exploiting their preferences for coupling to different G proteins (Abbracchio *et al.*, 2006). P2Y14 recep-

tors activate G $\alpha_{i/o}$ (Moore *et al.*, 2003), whereas P2Y6 receptors are expected to couple to phospholipase C via G $\alpha_{q/11}$ (Chang *et al.*, 1995; Robaye *et al.*, 1997). We used *Pertussis* toxin to uncouple P2Y14 receptors from G $\alpha_{i/o}$ and so resolve the relative roles of P2Y6 and P2Y14 receptors in initiating the Ca²⁺ signals evoked by UDP. ASMC were incubated with *Pertussis* toxin (100 ng·mL⁻¹) for 24 h prior to examining the effects of nucleotides on [Ca²⁺]. This treatment significantly reduced the response to ATP: in paired comparisons, the maximal response was reduced by 24 \pm 6% and the pEC₅₀ shifted from 5.63 \pm 0.09 to 5.30 \pm 0.08 (Figure 7A). This is consistent with evidence that P2Y2 receptors (via which ATP exerts its major Ca²⁺-mobilizing effect in ASMC, Figure 6) stimulate phospholipase C via both G α_i and G $\alpha_{q/11}$ (and/or G α_{16}) (Baltensperger and Porzig, 1997; Murthy and Makhlof, 1998). However, under identical conditions responses to UDP were unaffected by *Pertussis* toxin (Figure 7B). These results, together with the lack of response to the P2Y14-selective agonists (MRS2690 and UDP-glucose), suggest that responses of ASMC to UDP are mediated by P2Y6 rather than P2Y14 receptors.

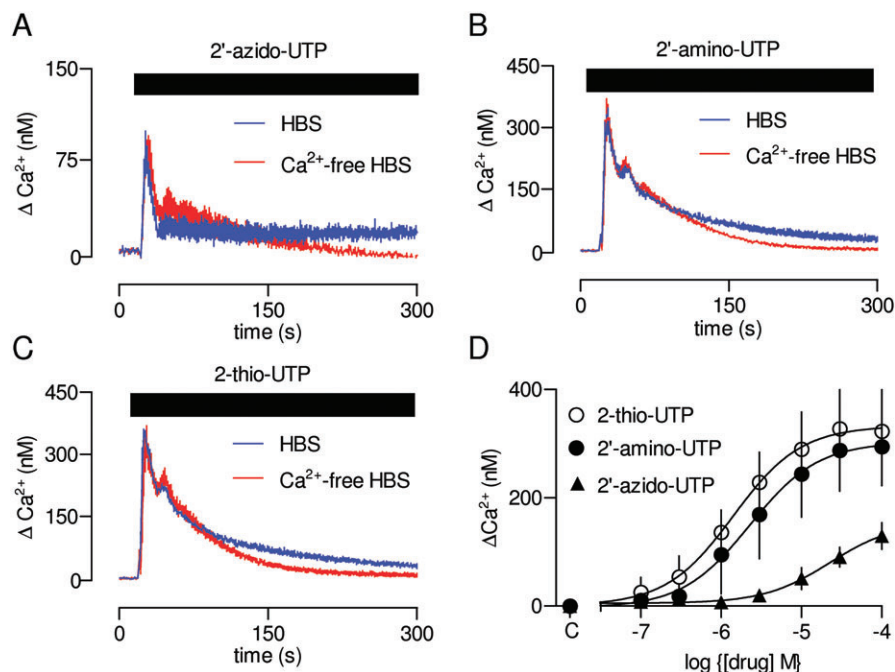


Figure 6 Ca²⁺ signals evoked by activation of P2Y2 and P2Y4 receptors. Typical results from populations of cultured ASMC stimulated with 2'-azido-UTP (100 μ M, A), 2'-amino-UTP (100 μ M, B) or 2-thio-UTP (100 μ M, C). Results are means \pm SEM from three wells on a single plate and are typical of results from three independent plates. Summary results showing concentration-dependent effects of the three analogues on the peak Ca²⁺ signal (D). C denotes control. 2'-amino-UTP, 2'-amino-2'-deoxyuridine-5'-triphosphate; 2'-azido-UTP, 2'-azido-2'-deoxyuridine-5'-triphosphate; 2-thio-UTP, 2-thio-uridine-5'-triphosphate; ASMC, aortic smooth muscle cell; HBS, HEPES-buffered saline.

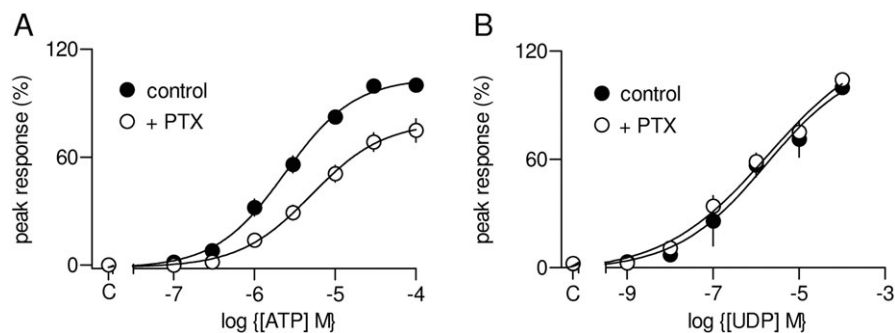


Figure 7 Ca²⁺ signals evoked by activation of P2Y6 receptors. Concentration-dependent effects of ATP (A) or UDP (B) on the peak Ca²⁺ signal with or without pretreatment with *Pertussis toxin* (PTX, 100 ng·mL⁻¹, 24 h). Results are expressed as percentages of the peak Ca²⁺ signal evoked by a maximally effective concentration of the agonist alone. C denotes control.

Discussion

Rat (Erlinge *et al.*, 1998) and human (Wang *et al.*, 2002) vascular smooth muscle cells express several subtypes of P2Y receptors, but with the limited range of selective ligands available it has proven difficult to resolve fully the contributions of each to functional responses. We have shown that mRNA encoding each of the seven rat P2Y receptor subtypes is expressed in cultured ASMC, and a subset of these (P2Y2, P2Y12 and P2Y14) is expressed in freshly isolated ASMC (Figure 3). Our finding that P2Y6 predominates in cultured cells, whereas P2Y12 is the major subtype in fresh cells, is consistent with earlier evidence that P2Y6 receptors regulate proliferation (Hou *et al.*, 2002).

In cultured ASMC, ATP, UTP and UDP evoke Ca²⁺ signals via activation of phospholipase C, and each stimulates both

release of Ca²⁺ from intracellular stores and Ca²⁺ entry (Figures 2C–E and 3). We have not further investigated the identity of the Ca²⁺ entry pathway, although published evidence suggests that it may be via store-operated Ca²⁺ entry (Potier *et al.*, 2009), a diacylglycerol-regulated TRP protein (Inoue *et al.*, 2001) or by a Na⁺/Ca²⁺ exchanger (Lemos *et al.*, 2007). By combining analyses of ligands that are either entirely or partially subtype-selective, we have established that four P2Y receptors evoke Ca²⁺ signals and the conditions that allow each to be activated selectively (Figure 8). P2Y1 receptors, for which ADP is the most likely physiological stimulus, evoke Ca²⁺ signals that appear to desensitize rapidly. P2Y2 receptors, which are likely to be activated endogenously by ATP and UTP, evoke the largest Ca²⁺ signals, and these are sustained by Ca²⁺ entry during prolonged stimulation. P2Y4 receptors are also likely to be activated by ATP and UTP, but

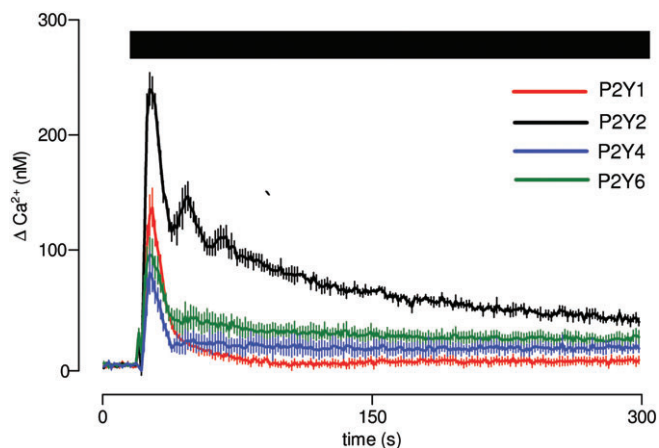


Figure 8 Different Ca²⁺ signals evoked by selective activation of different P2Y receptors. Typical results from populations of cultured ASMC stimulated in HBS with agonists that allow selective activation of P2Y1 (1 μ M MRS2365), P2Y2 (100 μ M 2'-amino-UTP), P2Y4 (100 μ M 2'-azido-UTP) or P2Y6 receptors (100 μ M UDP). Results are means \pm SEM from three wells on a single plate and are typical of results from three independent plates. 2'-amino-UTP, 2'-amino-2'-deoxyuridine-5'-triphosphate; 2'-azido-UTP, 2'-azido-2'-deoxyuridine-5'-triphosphate; ASMC, aortic smooth muscle cell; MRS2365, (N)-methanocarba-2-MeSADP.

they evoke considerably smaller Ca²⁺ signals than do P2Y2 receptors, although again the sustained phase of the response is maintained without obvious desensitization. The relative size of the Ca²⁺ signals evoked by selective activation of P2Y2 and P2Y4 receptors is consistent with an earlier suggestion that P2Y2 receptors mediate most of the response to UTP and ATP (Kumari *et al.*, 2003). In the absence of better ligands, that conclusion derived from use of suramin as a competitive antagonist of P2Y2 receptors (Wildman *et al.*, 2003), but suramin is now known to interact also with P2Y1 and P2Y6 receptors (Abbraccio *et al.*, 2006), both of which evoke Ca²⁺ signals in cultured ASMC (Figure 8). Finally, P2Y6 receptors, for which the endogenous ligand is probably UDP, evoke Ca²⁺ signals with a large sustained Ca²⁺ entry phase (Figure 8). This sustained response is consistent with the lack of desensitization reported for human P2Y6 receptors (Robaye *et al.*, 1997). We also detected mRNA encoding P2Y12, P2Y13 and P2Y14 receptors in cultured ASMC (Figure 3A). Of these, P2Y14 receptors have been shown to be capable of evoking Ca²⁺ signals (Fumagalli *et al.*, 2003), but we found no evidence to suggest that any of these P2Y receptors evokes Ca²⁺ signals in cultured ASMC (Figures 5 and 7).

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Conflicts of interest

None.

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Supporting information

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

- Figure S1** Structures of the ligands used.
- Figure S2** Effects of an ATP/UTP-regenerating system on responses to ATP and UTP. Concentration-dependent effects of ATP (A) and UTP (B) on [Ca²⁺]_i (cytosolic free Ca²⁺ concentration) in aortic smooth muscle cell in HEPES-buffered saline alone or after pretreatment of cells and ligands with an ATP/UTP-regenerating system (as described in the text).
- Figure S3** Effects of MRS2395 (2,2-dimethyl-propionic acid 3-(2-chloro-6-methylaminopurin-9-yl)-2-(2,2-dimethyl-propionyloxymethyl)-propyl ester) and MRS2211 (2-[(2-chloro-5-nitrophenyl)azo]-5-hydroxy-6-methyl-3-[(phosphonoxy)methyl]-4-pyridinecarboxaldehyde) on the Ca²⁺ signals evoked by 2-MeSADP (2-methylthio-adenosine-5'-diphosphate). Concentration-dependent effects of 2-MeSADP on [Ca²⁺]_i (cytosolic free Ca²⁺ concentration) alone or in the presence of MRS2395 (30 μ M, A) or MRS2211 (10 μ M, B). Results are expressed as percentages of the maximal response to 2-MeSADP alone.
- Figure S4** Effects of MRS2279 (2-chloro-N⁶-methyl-(N)-methanocarba-2'-deoxyadenosine-3',5'-bisphosphate) on the Ca²⁺ signals evoked by ATP. Concentration-dependent effects of ATP on [Ca²⁺]_i (cytosolic free Ca²⁺ concentration) alone or with MRS2279 (3 μ M).
- Table S1** Primers used for quantitative PCR

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