

# Cloning, Up-Regulation, and Mitogenic Role of Porcine P2Y<sub>2</sub> Receptor in Coronary Artery Smooth Muscle Cells

Jianzhong Shen, Cheikh I. Seye, Meifang Wang, Gary A. Weisman, Peter A. Wilden, and Michael Sturek

*Departments of Medical Pharmacology and Physiology (J.S., M.W., P.A.W., M.S.), Biochemistry (C.I.S., G.A.W.), and Internal Medicine (M.S.), and the Center for Diabetes and Cardiovascular Health (J.S., M.W., P.A.W., M.S.), University of Missouri-Columbia, School of Medicine, Columbia, Missouri*

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## ABSTRACT

Previous work has shown up-regulation of a UTP-sensitive P2Y receptor in porcine coronary smooth muscle cells (CSMC) of organ-cultured arteries. However, the molecular identity and functional role of this putative receptor remained undefined. Here we report the cloning of the cDNA for this receptor that encodes an open reading frame for a protein of 373 amino acids with the highest homology to the human P2Y<sub>2</sub> receptor (84%). Heterologous expression of this receptor in 1321N1 cells revealed a novel pharmacology in that UTP and ITP were full agonists and UTP was more potent and efficacious than ATP for increasing intracellular [Ca<sup>2+</sup>]<sub>i</sub> and extracellular signal-regulated kinase phosphorylation. Stimulation of subcultured CSMC with UTP, ITP, or ATP induced a concentration-dependent increase in cellular DNA content, protein synthesis, cell number, and proliferating cell nuclear antigen expression, indicating a mitogenic role for P2Y<sub>2</sub> receptors. This was supported

by the finding that the treatment of CSMC with antisense oligonucleotides to the cloned cDNA sequence significantly inhibited UTP- and ATP-induced DNA and protein synthesis. In addition, reverse transcription-polymerase chain reaction analysis showed that P2Y<sub>2</sub> receptor mRNA was dramatically increased in cells of organ-cultured arteries compared with freshly harvested arteries, whereas the P2Y<sub>6</sub> receptor mRNA level was unchanged, and the P2Y<sub>4</sub> receptor mRNA was undetectable. This P2Y<sub>2</sub> subtype-specific up-regulation was confirmed in cells of coronary arteries stented in vivo. In conclusion, we have cloned the porcine P2Y<sub>2</sub> receptor with novel pharmacology and demonstrated that this receptor is up-regulated in CSMC of in vitro organ cultures or in vivo stented coronary arteries to mediate the mitogenic effects of nucleotides.

The acute effects of extracellular nucleotides in the regulation of vascular tone have been well documented. Specifically, ATP, ADP, UTP, and UDP can induce endothelium-dependent vasorelaxation via P2Y receptor-mediated nitric oxide, endothelium-derived hyperpolarizing factor, or prosta-

cyclin production (Wihlborg et al., 2003). Direct vasoconstriction in some blood vessels can also be triggered by these naturally occurring nucleotides via activation of P2X and/or P2Y receptors expressed in vascular smooth muscle cells (VSMC) (Matsumoto et al., 1997; Malmstroj et al., 2000a,b, 2003). In addition to these acute effects, recent studies have shown that extracellular nucleotides, in particular ATP and UTP, also have potent long-term effects on VSMC, i.e., proliferation (Erlinge, 1998; Burnstock, 2002) and migration (Pillois et al., 2002), both of which are important in the development of vascular diseases such as atherosclerosis and postangioplasty restenosis.

It is generally thought that P2X receptors, being ligand-gated ion channels, are not involved in the chronic effects of ATP and UTP, suggesting a role for G protein-coupled P2Y

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**ABBREVIATIONS:** VSMC, vascular smooth muscle cells; CSMC, coronary artery smooth muscle cells; ERK, extracellular signal-regulated kinase; DMEM, Dulbecco's modified Eagle's medium; FBS, fetal bovine serum; PCR, polymerase chain reaction; RACE, rapid amplification of cDNA ends; UTR, untranslated region; LPA, lysophosphatidic acid; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; PCNA, proliferating cell nucleus antigen; CCh, carbachol hydrochloride; RT-PCR, reverse transcription-polymerase chain reaction; bp, base pair(s); G3PDH, glyceraldehyde-3-phosphate dehydrogenase; OGN, oligonucleotide(s); RB-2, reactive blue 2; PPADS, pyridoxal phosphate-6-azophenyl-2',4'-disulfonic acid.

receptors, of which eight different subtypes (P2Y<sub>1, 2, 4, 6, 11, 12, 13, 14</sub>) have been cloned and pharmacologically characterized (Abbracchio et al., 2003; Dubyak, 2003). Interestingly, it turns out that, among the Gq-coupled P2Y receptors (P2Y<sub>1, 2, 4, 6, 11</sub>), only P2Y<sub>2</sub> was up-regulated in vascular injury models, including balloon-injured rat aorta (Seye et al., 1997) and collared rabbit carotid arteries (Seye et al., 2002). Studies on subcultured rat aortic VSMC have shown that ATP and UTP, agonists of P2Y<sub>2</sub> receptors, triggered cell proliferation (Ertinge et al., 1993, 1995; Malam-Souley et al., 1993, 1996). A similar mitogenic effect of ATP on subcultured porcine coronary artery smooth muscle cells (CSMC) has also been reported (Wilden et al., 1998). However, because of the lack of specific antagonists for most of the P2Y receptors, including P2Y<sub>2</sub>, the exact P2Y receptor subtype(s) responsible for the mitogenic effects of ATP and UTP, specifically in CSMC, has not been defined. Other complications are that P2Y receptor subtype expression in VSMC is highly heterogeneous and varies with species, type of vascular bed, and developmental state (Malmsjo et al., 2000a,b, 2003; Hill et al., 2001). Therefore, it is critical to choose the appropriate animal model to address the pathological significance of the individual P2Y receptor in the development of relevant diseases. This view is supported by the unexpected finding that UTP, an agonist of the P2Y<sub>2</sub> receptor, is a potent antimitogen in cultured human VSMC derived from internal mammary artery and saphenous vein (White et al., 2000). Thus, it remains to be determined whether P2Y<sub>2</sub> receptor up-regulation and contribution can be generalized to other vascular disease models, specifically coronary artery disease.

Previous work in our lab has shown that CSMC dispersed from freshly harvested porcine coronary arteries do not express a functional Gq-coupled uridine nucleotide P2Y receptor, as evidenced by the lack of intracellular Ca<sup>2+</sup> mobilization in response to UTP (Hill et al., 2001; Hill and Sturek, 2002). However, UTP triggered robust Ca<sup>2+</sup> mobilization in dispersed CSMC from organ-cultured coronary arteries, indicating that a UTP-sensitive P2Y receptor(s) was up-regulated under organ culture conditions (Hill et al., 2001; Hill and Sturek, 2002). This notion was further supported by the pharmacological inhibition of UTP-induced Ca<sup>2+</sup> mobilization by P2Y receptor antagonists and inhibitors of G proteins and the sarcoplasmic reticulum Ca<sup>2+</sup> pump (Hill and Sturek, 2002). Therefore, we attempted to determine the molecular identity and the (patho)-physiological role(s) of this UTP-sensitive P2Y receptor. The identification of the mitogenic P2Y receptor that is up-regulated in porcine CSMC of diseased coronary arteries is important, because these animals are excellent models for human cardiovascular disease (Johnson et al., 1999).

Thus, the aims of this study were 1) to clone and characterize this porcine UTP-sensitive P2Y receptor and 2) to determine whether the activation of this receptor is responsible for the nucleotide-induced proliferation of CSMC. Data presented herein show that the cloned porcine receptor has extensive homology with the human P2Y<sub>2</sub> receptor but a different agonist potency profile in that the porcine P2Y<sub>2</sub> receptor is more sensitive to UTP than ATP with respect to nucleotide-induced increases in the intracellular [Ca<sup>2+</sup>] and mitogen-activated protein kinase (ERK) phosphorylation. These relative agonist potencies are similar to the UTP-sensitive receptor previously characterized in organ-cultured

porcine coronary arteries (Hill et al., 2001; Hill and Sturek, 2002) and in monolayer cultures of porcine CSMC (Seiler et al., 1999). Finally, a mitogenic role for the cloned P2Y<sub>2</sub> receptor in porcine CSMC was confirmed by the inhibition of mitogenesis with the P2Y<sub>2</sub> receptor antisense oligonucleotide, suggesting that these receptors may play a role in the development of coronary artery disease.

## Materials and Methods

**Isolation and Culture of Porcine CSMC.** Porcine hearts were obtained from domestic farm or Yucatan pigs. Coronary arteries dissected from the heart and denuded of endothelium using aseptic techniques were placed in a physiological buffer, and smooth muscle cells were isolated enzymatically (Hill et al., 2001; Hill and Sturek, 2002). Dispersed cells were recovered and subcultured at 37°C with 5% CO<sub>2</sub> in DMEM/high-glucose medium containing 10% fetal bovine serum (FBS). Smooth muscle cell lineage was confirmed by  $\alpha$ -smooth muscle actin immunocytochemistry. Stock cell cultures were maintained in a subconfluent state and used before passage 10 (Wilden et al., 1998).

**PCR Cloning and RACE.** The cDNA for the porcine P2Y<sub>2</sub> receptor was cloned by a PCR cloning strategy. Total RNA was extracted from cultured CSMC using the RNeasy Total RNA Mini Kit (QIAGEN, Valencia, CA). For the synthesis of first-strand cDNA, 1  $\mu$ g of total RNA was reverse transcribed using a cDNA synthesis kit (BD Biosciences Clontech, Palo Alto, CA). The cDNA was then amplified by PCR using 2.5 U of *Taq* DNA polymerase (Promega, Madison, WI). The initial PCR reaction was performed with a pair of degenerate primers based on the conserved regions of human and rat P2Y<sub>2</sub> receptor sequences (forward, 5'-TGC/TCGCTTCAACGAGGACTTCAAGTA-3' and reverse, 5'-AG/CGCCATGTTGATGGCGTTGAGGGT-3'). After sequencing several of the obtained PCR products of the expected size, new sets of primers specific to the putative porcine P2Y<sub>2</sub> receptor were synthesized and used to obtain the full-length cDNA.

To obtain the 3'-end of the coding sequence of the porcine P2Y<sub>2</sub> receptor cDNA, we used a SMART RACE cDNA amplification kit (BD Biosciences Clontech). In brief, total RNA was reverse transcribed using an oligo(dT)<sub>20</sub> primer with adaptor sequence. The first round of PCR was carried out with a porcine P2Y<sub>2</sub> receptor cDNA-specific forward primer (5'-CGTCACCCGACCCCTCTACTACTCTT-3') and a reverse adaptor primer provided in the kit. The PCR product was diluted 1/1000 in diethyl pyrocarbonate-treated water and used as a template for a second round of PCR. The second round of PCR was performed as seminested PCR, in which the same reverse primer was used with a different gene-specific forward primer (5'-CCTGAGCTACGGCGTGGTGTGCGTG-3') downstream of the original primer. The PCR conditions were as follows: jump start for 1 min at 94°C, denaturation for 1 min at 94°C, annealing for 1 min at 60°C, and extension at 72°C for 1 min for 35 cycles.

To obtain the 5'-end of the coding sequence of the porcine P2Y<sub>2</sub> receptor cDNA, we designed a degenerate forward primer based on the one conserved sequence located in the 5'-untranslated region (5'-UTR) of human and rat P2Y<sub>2</sub> receptor cDNAs. The degenerate forward primer (5'-GGA/GACCTGTTT/CTTCTGTTTCC-3') and a gene-specific reverse primer (5'-CGGCACAGGAAGATGTAGAGCGCC-3') were used to amplify a new PCR product. After sequencing several products of the anticipated size, partial cloning of the 5'-UTR sequence was achieved. Then a final set of primers (forward, 5'-GAGCATCCTGACAGCGAGAGCAG-3' and reverse, 5'-CTACAGCCGGATGTCTTCGCACAG-3'), based on the 5'-UTR and 3'-UTR sequence flanking the open reading frame, was used for amplification of the full-length porcine P2Y<sub>2</sub> receptor cDNA. The PCR products resolved on an agarose gel were isolated using a QIAquick gel extraction kit (QIAGEN), and the purified cDNA was ligated into the plasmid vector pCR3.1 (a modified form of pcDNA3.1) (Invitrogen, Carlsbad, CA). The ligation product was used to transform One Shot TOP10F'-competent *Escherichia coli* cells from the

TA cloning kit (Invitrogen). The cDNA inserts from at least three different clones were sequenced in both directions using T7 and bovine growth hormone primers. DNA sequencing was performed by the DNA Core Facility of the University of Missouri-Columbia.

**Heterologous Expression.** The purified pCR3.1 plasmids harboring the coding sequence of the porcine P2Y<sub>2</sub> receptor (pCR3.1-pP2Y<sub>2</sub>) were transfected into human 1321N1 astrocytoma cells in DMEM with 10% FBS using Effectene Transfection Reagent (QIAGEN). Stable transfectants were obtained by selection in medium containing 0.5 mg/ml G418 (geneticin) (Calbiochem, San Diego, CA) for ~2 to 3 weeks. The efficiency of transfection was determined by monitoring UTP-induced Ca<sup>2+</sup> mobilization in transfected cells.

**[<sup>3</sup>H]Thymidine Incorporation.** The incorporation of [<sup>3</sup>H]thymidine into DNA was carried out as described previously (Wilden et al., 1998). Cells were cultured in 12-well culture plates as described above. Cells near confluence were treated with or without nucleotides or LPA for 24 h at 37°C in DMEM/high-glucose medium containing 0.5% FBS, followed by the addition of 1 µCi of [methyl-<sup>3</sup>H]thymidine (PerkinElmer Life and Analytical Sciences, Boston, MA) for an additional 24 h at 37°C. The cells were washed three times in ice-cold PBS and solubilized in 0.1% SDS. Trichloroacetic acid was added to a final concentration of 10%, and the precipitate was collected by filtration on glass-fiber disks for the determination of radioactivity by liquid scintillation counting.

**Protein Synthesis.** [<sup>3</sup>H]Leucine incorporation was used for measuring total cellular protein synthesis. The method is similar to that described under [<sup>3</sup>H]Thymidine Incorporation, except that 1 µCi of [<sup>3</sup>H]leucine instead of [<sup>3</sup>H]thymidine was added to the medium in the presence or absence of UTP, ATP, UDP, or LPA.

**Colorimetric Determination of Cell Number.** Cultured CSMC were suspended by trypsinization, counted, replated in 96-well plates at a density of 2000 cells/well in cell culture medium containing 10% FBS, and then serum-starved for 48 h. Agonists were added and present for 4 days. By the end of cell stimulation, 20 µl of the tetrazolium compound 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) in 100 µl of fresh DMEM was added for 4 h at 37°C according to the instructions in the MTT Cell Growth Assay Kit (Chemicon International, Temecula, CA). MTT reduced to formazan by intracellular dehydrogenases was determined at 490 nm with an enzyme-linked immunosorbent assay plate reader and was directly proportional to the number of living cells.

**Determination of p-ERK and PCNA by Western Analysis.** Cells were serum-deprived for 24 h (48 h for PCNA) before stimulation with agonists at the indicated concentration for 5 min (ERK assay) or 24 h (PCNA assay). Then cells were washed with ice-cold phosphate-buffered saline, solubilized in Laemmli sample buffer containing 200 mM dithiothreitol, and boiled. Lysates were sonicated to disrupt DNA, and proteins were separated on 10% SDS-polyacrylamide gel electrophoresis gels. The proteins were electrophoretically transferred to nitrocellulose in 25 mM Tris, 192 mM glycine, 20% methanol, and 0.02% SDS. The nitrocellulose was blocked with 5% nonfat milk in 20 mM Tris, pH 7.4, 150 mM NaCl, and 0.01% Tween 20. The membranes were probed with the primary antibody anti-p-ERK1/2 (1:2000; Cell Signaling Technology Inc., Beverly, MA) or anti-PCNA (1:1000; Cell Signaling Technology Inc.) overnight in 20 mM Tris, pH 7.4, 150 mM NaCl, 3% bovine serum albumin, and 0.01% Tween 20. The blots were washed in 20 mM Tris, pH 7.4, 150 mM NaCl, and 0.01% Tween 20, and the bound antibody was detected by horseradish peroxidase-conjugated anti-rabbit IgG and enhanced chemiluminescence (Pierce, Rockford, IL). As a loading control, membranes were stripped of original antibodies and re-probed with primary anti-actin antibody (1:1000; Cytoskeleton, Denver, CO).

**Organ Culture of Coronary Arteries and Dispersion of Smooth Muscle Cells.** Pig hearts were isolated and prepared as described previously (Hill et al., 2000, 2001; Hill and Sturek, 2002). Right coronary arteries (distal end) were either stored for 4 days at 5°C or organ-cultured for 4 days at 37°C in a 95% O<sub>2</sub>/5% CO<sub>2</sub>

incubator. Arterial segments were denuded of endothelium, opened longitudinally to expose the lumen, and placed in a 100-mm Petri dish containing 30 ml of RPMI 1640 (Invitrogen) without serum, which was changed every 2 days. CSMC were enzymatically isolated as described previously (Hill et al., 2000, 2001; Hill and Sturek, 2002) and identified morphologically (Hill et al., 2000).

**Porcine Model of Coronary Artery Stenting.** Protocols conformed to the Animal Care and Use Guidelines of the University of Missouri-Columbia. A baseline coronary angiogram for the pigs was performed, and the circumflex artery was selected for the implantation of an appropriately sized stent (3.0 mm in diameter and 13 mm long) centered on a 3.0 × 15-mm balloon (Guidant Corporation, Indianapolis, IN). To avoid overinflation injury, a stent-to-artery ratio of 1:1 compared with the baseline segment diameter was assessed with angiography and intravascular ultrasound. The 10 mm of artery distal to the stenting site served as a control segment. After recovery for 3 weeks, pigs were anesthetized and sacrificed. The isolation of coronary artery and dispersion of CSMC are described above.

**Intracellular [Ca<sup>2+</sup>] Measurements.** Intracellular free Ca<sup>2+</sup> concentration ([Ca<sup>2+</sup>]<sub>i</sub>) in transfected 1321N1 cells and cultured CSMC was measured with the InCa<sup>2+</sup> calcium imaging system (Intracellular Imaging, Inc., Cincinnati, OH). Cells were grown on glass coverslips for 2 days, serum-starved for another 24 h, and then used at about 40% confluence. After incubation with 2.5 µM Fura-2/acetoxymethyl ester at 37°C for 30 min, the cells were superfused in a constant-flow superfusion chamber mounted on an inverted epifluorescence microscope (model TMD; Nikon, Melville, NY) using physiological salt solution containing 138 mM NaCl, 5 mM KCl, 2 mM CaCl<sub>2</sub>, 1 mM MgCl<sub>2</sub>, 10 mM HEPES, and 10 mM glucose, pH 7.4, in the presence or absence of the indicated concentrations of nucleotides, LPA, or carbachol hydrochloride (CCh). Fura-2 was excited at 340 and 380 nm, and emitted fluorescence (510 nm) was collected using a monochrome charge-coupled device camera (Cohu Electronics, San Diego, CA). Data were collected as a ratio of emitted light intensity at 340 and 380 nm and then converted to [Ca<sup>2+</sup>]<sub>i</sub> based on a standard curve (Hill et al., 2000, 2001; Hill and Sturek, 2002).

**Semiquantitative RT-PCR.** Total RNA was isolated from cultured or dispersed cells, and RT-PCR protocols were performed as described in *PCR Cloning and RACE*. Sets of primers used for detecting and semiquantifying mRNA for individual P2Y receptor subtypes were porcine P2Y<sub>2</sub> receptor (GenBank accession no. AY620400): forward, 5'-CGTCACCCGCACCTCTACTACTCCTT-3' and reverse, 5'-CTACAGCCGGATGTCTTCGCC-ACG-3', yielding a 340-bp product; porcine P2Y<sub>6</sub> receptor (GenBank accession no. AY620399): forward, 5'-CTGCATCAGCTTCCAGCGCTACCT-3' and reverse, 5'-GTCCAAGACGCTGTTGGCACTGGC-3', yielding a 538-bp product; porcine P2Y<sub>4</sub> receptor (GenBank accession no. AY662405): forward, 5'-GTTTGATGAGGATTT-CAAGTTCATCC-3' and reverse, 5'-CAGACAGCAAAGACAGTCAGCAC-3', yielding a 686-bp product; and human G3PDH (GenBank accession no. X01677): forward, 5'-TGAAGGTCGGAGTCAACGGATTGTTGGT-3' and reverse, 5'-CATGTGGGCCATGAGGTCCACCAC-3', yielding a 983-bp product. The PCR conditions were as described in *PCR Cloning and RACE*, except for the P2Y<sub>4</sub> receptor detection, for which the annealing temperature was 56°C, and 100 ng of porcine genomic DNA was used as a positive control. The resulting PCR products were resolved on a 2% agarose ethidium bromide gel. The amplified bands were visualized with ultraviolet light, and the relative densities of individual bands were normalized to G3PDH using the Quantity One computer program.

**Antisense Oligonucleotide Experiments.** Phosphorothioate-modified oligonucleotides (OGN) designed according to the cDNA sequence of the porcine P2Y<sub>2</sub> receptor cloned in this study were antisense OGN (5'-TCGGGGCCTGTAGCCATCAC-3') and sense OGN (5'-GTGATGGCTACAGGCCCGCA-3'). Sequence uniqueness was checked across multiple databases using BLAST. An optimized concentration of 3 µM sense or antisense OGN was added to the culture medium 24 h before and after serum starvation. Cellular



uptake of OGN was verified using fluorescein isothiocyanate-conjugated OGN under the same experimental conditions. Nucleotide-induced  $\text{Ca}^{2+}$  mobilization was assayed to verify inhibition of porcine P2Y<sub>2</sub> receptor expression by antisense OGN.

**Materials.** Fura 2-AM was purchased from Molecular Probes (Eugene, OR). Cell culture media were obtained from Invitrogen. Unless indicated, all other reagents and compounds were purchased from Sigma-Aldrich (St. Louis, MO). PCR primers and sense/antisense OGN were synthesized and purified by Integrated DNA Technologies (Coralville, IA). All nucleoside diphosphates were prepared in stock solutions (1 mM) that contained 50 U/ml hexokinase and 110 mM glucose to eliminate the potential contamination of the corresponding nucleoside triphosphate.

**Data Analysis.** Data are expressed as means  $\pm$  S.E.M. All concentration-response curves were fitted by logistic (Hill equation), nonlinear regression analysis using Prism 4.0 (GraphPad Software Inc., San Diego, CA). Means of two groups were compared using Student's *t* test (unpaired, two-tailed), and one-way analysis of variance was used for the comparison of more than two groups, with *p* < 0.05 considered to be statistically significant.

## Results

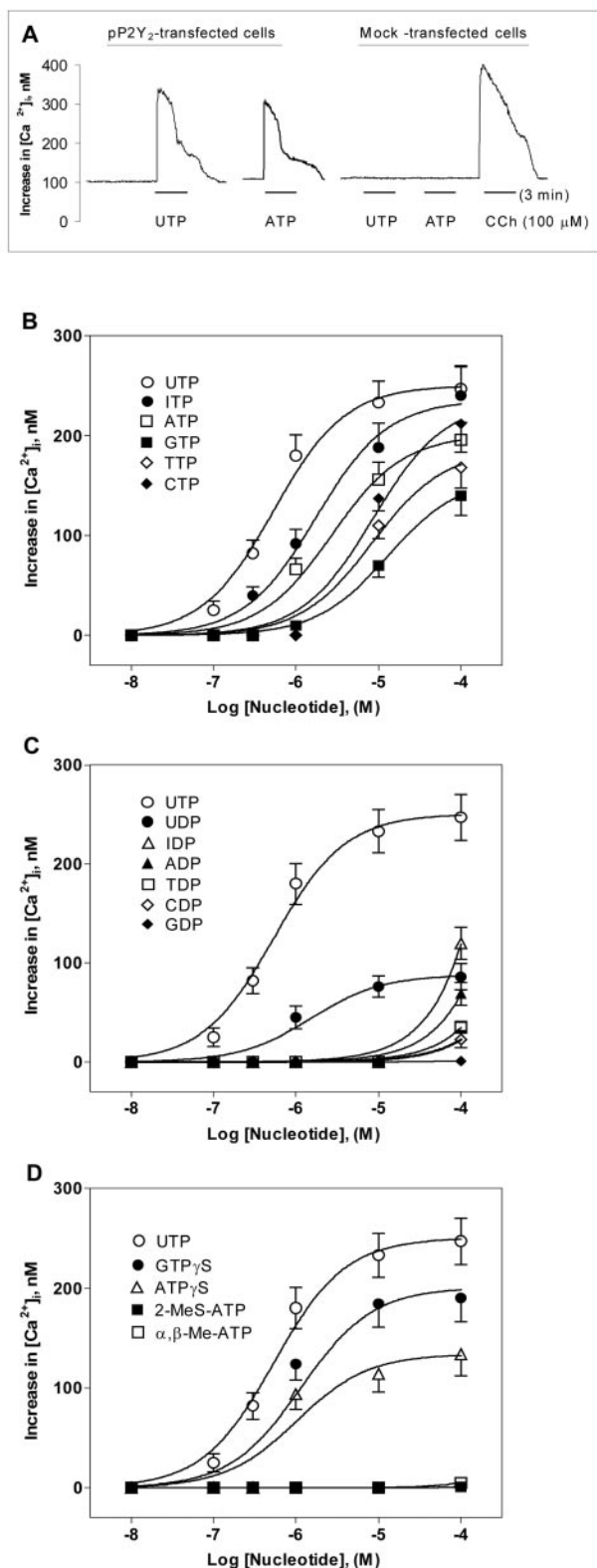
**Cloning of the Porcine P2Y<sub>2</sub> Receptor.** Based on the conserved regions of coding sequences for human and rat P2Y<sub>2</sub> receptors, degenerate primers were constructed and used for initial RT-PCR experiments, with total RNA isolated from either cultured CSMC or coronary artery denuded of endothelium. After sequencing PCR products of anticipated size, sets of new gene-specific forward and reverse primers were designed that enabled the isolation of full-length porcine P2Y<sub>2</sub> receptor cDNA with an open reading frame of 1119 bp (GenBank accession no. AY620400). Hydrophobicity anal-

ysis showed a deduced primary sequence for the porcine P2Y<sub>2</sub> receptor of 373 amino acids containing seven hydrophobic domains characteristic of G protein-coupled receptors (Fig. 1). Computer sequence alignment revealed that the porcine P2Y<sub>2</sub> receptor had the highest amino acid identity with human P2Y<sub>2</sub> receptors (84%) and had four amino acids deleted at two different positions of the C terminus (Fig. 1). In addition, several V-to-A/G transitions are notable and scattered in various extracellular loops and transmembrane regions. Furthermore, a putative *N*-glycosylation site and most of the putative phosphorylation sites for protein kinase C, protein kinase A, and G protein-coupled receptor kinase (Zamboni et al., 2000) are conserved (annotated in Fig. 1). Interestingly, an integrin-binding Arg-Gly-Asp acid motif found in the first extracellular loop of the human P2Y<sub>2</sub> receptor (Erb et al., 2001) is also conserved in the porcine P2Y<sub>2</sub> receptor. However, the porcine P2Y<sub>2</sub> receptor does not contain the "PXXP" SH3 binding domain, recently identified in the C terminus of the human P2Y<sub>2</sub> receptor (Liu et al., 2004). RT-PCR analysis showed a wide tissue distribution for porcine P2Y<sub>2</sub> receptor mRNA, including heart, kidney, intestine, skeletal muscle, and adrenal glands (data not shown).

**Stable Expression and Functional Characterization of the Cloned Porcine P2Y<sub>2</sub> Receptor.** To determine whether the cloned cDNA sequence of the porcine P2Y<sub>2</sub> receptor encodes a functional receptor, a recombinant mammalian expression vector harboring the porcine P2Y<sub>2</sub> receptor cDNA (pCR3.1-pP2Y<sub>2</sub>) was stably transfected into human 1321N1 astrocytoma cells devoid of any endogenous P2Y receptors (Parr et al., 1994). Figure 2A shows that 100  $\mu\text{M}$

Human	MAADLGPWND	TINGTWDGDE	LGYRCRFNED	FKYVLLPVSY	GVVCVPGICL	50
Porcine	MATGPDLLWG	TVNGTSDGDD	WGYRCRFHED	FKYVLLPLSY	GVVCVLGLSL	
Consensus	MA....WN.	T+NGT.DGD+	.GYRCRF.ED	FKYVLLP+SY	GVVCV.GL.L	
		*			TM I	
	NAVALYIFLC	RLKTWNASTT	YMFHLAVSDA	LYAASLPLLV	YYYARGDHWP	100
	NAGALYIFLC	RLKTWNASTT	YMFHLAVSDA	LYAASLPLLV	YYYARGDHWP	
	NA.ALYIFLC	RLKTWNASTT	YMFHLAVSDA	LYAASLPLLV	YYYARGDHWP	
					TM II	!!!
	FSTVLCKLVR	FLFYTNLYCS	ILFLTICISVH	RCLGVLRLPLR	SLRWGRARYA	150
	FSTALCKLVR	FLFYTNLYCS	ILFLTICISVH	RCLGVLRLPLR	SLRWGHARYA	
	FST.LCKLVR	FLFYTNLYCS	ILFLTICISVH	RCLGVLRLPLR	SLRWG.ARYA	
					TM III	
	RRVAGAVVWL	VLACQAPVLY	FVTTSARGGR	VTCHDTSAP	LFSRFVAYSS	200
	RRVAAAVWGL	VLACQAPALY	FITTTAQGGR	ITCHDTSAP	LFSHFVAYSL	
	RRVA+AVW.L	VLACOAP.LY	F+TT+A.GGR	+TCHDTSAP	LFS.FVAYS.	
					TM IV	
	VMLGLLFAVP	FAVILVCYVL	MARRLLKPAY	GTSGGLPRAK	RKSVRTIAVV	250
	VMLSVLFAAP	FAVILVCYAL	MARRLLRPAY	GTAGGLPRAK	RKSVRTIAVV	
	VML.+LFA.P	FAVILVCY.L	MARRLL+PAY	GT+GGLPRAK	RKSVRTIAVV	
					TM V	
	LAVFALCFPL	FHVTRTLYYS	FRSLDLSCHT	LNAINMAYKV	TRPLASANSC	300
	LAVFALCFPL	FHVTRTLYYS	FRTLDLSCHT	LDAINMAYKI	TRPLASANSC	
	LAVFALCFPL	FHVTRTLYYS	FR+LDLSCHT	L.AINMAYK+	TRPLASANSC	
					TM VI	TM VII
	LDPVLYFLAG	QRLVRFARDA	KPPTGPSPAT	PARRRLGLRR	SDRTDMQRIE	350
	LDPVLYFLAG	QRLVRFARDA	KPPTDATPTA	QACRRRLGLRR	SHGTDTKRTE	
	LDPVLYFLAG	QRLVRFARDA	KPPT..+P..	.A.RRLGLRR	S..TD..R.E	
	DVLGSSEDSR	RTESTPAGSE	NTKDIRL			377
	DS-ASSEDSR	RTEITPARGE	---DIRL			373
	D...SSEDSR	RTE.TPA..E	...DIRL			
	▲▲	●	●			

**Fig. 1.** Deduced amino acid sequence of the cloned porcine P2Y<sub>2</sub> receptor and alignment with the human P2Y<sub>2</sub> receptor sequence. Putative transmembrane-spanning domains are indicated with solid lines and marked as TM I to TM VII. Sequence matches are shaded only, and conservative substitutions are shaded and marked as +. Gaps (-) are introduced to best fit the alignment. A potential phosphorylation site for G protein-coupled receptor kinase is denoted with ▲▲, and potential phosphorylation sites for protein kinase A or protein kinase C are indicated by ●. !!!, conserved integrin-binding Arg-Gly-Asp acid motif; \*, and a consensus site for *N*-linked glycosylation.



**Fig. 2.** Nucleotide-induced changes in  $[Ca^{2+}]_i$  in human 1321N1 astrocytoma cells expressing porcine P2Y<sub>2</sub> receptors. Human 1321N1 cells, transfected with recombinant plasmid pCR3.1-pP2Y<sub>2</sub> or mock plasmid pCR3.1, were used for  $Ca^{2+}$  mobilization assays with UTP, ATP, or CCh as agonists (A). Concentration-response relationships for nucleoside triphosphates (B), diphosphates (C), and synthetic nucleotides (D) for the recombinant pP2Y<sub>2</sub> receptor expressed in 1321N1 cells. Data points in panels B through D indicate the average peak increase in  $[Ca^{2+}]_i$  for 8 to 16 cells on each of five coverslips and represent the means  $\pm$  S.E.M.

UTP or ATP increased  $[Ca^{2+}]_i$  in 1321N1 cells expressing the porcine P2Y<sub>2</sub> receptor, whereas both nucleotides were inactive in mock plasmid-transfected cells. CCh, a muscarinic M receptor agonist, triggered robust  $Ca^{2+}$  responses in the mock-transfected cells. We also tested the potential coupling of the cloned porcine P2Y<sub>2</sub> receptor to the Gs/Gi adenylyl cyclase cAMP signaling pathway and found no significant change in intracellular cAMP levels induced by UTP (data not shown), suggesting that the porcine P2Y<sub>2</sub> receptor is coupled primarily to phospholipase C via the Gq protein.

The cloned porcine P2Y<sub>2</sub> receptor was activated by all the nucleoside triphosphates, albeit with varying potencies (UTP > ITP > ATP > CTP  $\approx$  TTP > GTP) and efficacies (UTP  $\approx$  ITP  $\approx$  CTP > ATP  $\approx$  TTP > GTP), with ATP, TTP, and GTP acting as partial agonists (Fig. 2B; Table 1). In contrast, with the exception of UDP, which was a partial agonist (Fig. 2C), most of the nucleoside diphosphates were inactive at 10  $\mu$ M. Some synthetic nucleotides, including ATP $\gamma$ S and GTP $\gamma$ S, were also active with potencies and efficacies comparable with that of ATP, whereas 2-methylthio-ATP and  $\alpha,\beta$ -methyl-ATP had no effect (Fig. 2D). We also investigated whether the cloned porcine P2Y<sub>2</sub> receptor expressed in 1321N1 cells could activate ERK, a component of the mitogen-activated protein kinase signaling pathway activated by many G protein-coupled receptors, including P2Y receptors (Burnstock, 2002; Liu et al., 2004). In pP2Y<sub>2</sub>-transfected 1321N1 cells, UTP induced robust phosphorylation of ERK1/2 with an EC<sub>50</sub> of  $\sim$ 10 nM, whereas ATP was a partial agonist with an EC<sub>50</sub> of  $\sim$ 100 nM. UDP also induced phosphorylation of ERK1/2 at concentrations  $\geq$  1  $\mu$ M (Fig. 3). The nonspecific P2Y receptor antagonists suramin, RB-2, and PPADS at 10  $\mu$ M caused 50% inhibition of the UTP-induced increase in peak  $[Ca^{2+}]_i$  in pP2Y<sub>2</sub>-1321N1 cells (data not shown), consistent with our previous results with organ-cultured CSMC (Hill and Sturek, 2002).

**Effects of UTP, ATP, and ITP on Proliferation of CSMC.** Treatment of CSMC with UTP, ATP, or ITP caused a concentration-dependent increase in cellular DNA (Fig. 4A) and protein synthesis (Fig. 4B) and cell number (Fig. 4C), whereas UDP (100  $\mu$ M) only caused a small increase in protein synthesis (Fig. 4B). Intriguingly, ATP was much more potent and efficacious than UTP, ITP, and UDP in increasing cellular DNA synthesis (Fig. 4A) and expression of PCNA, a protein marker of cell proliferation (Fig. 4D), suggesting that another receptor may contribute to the proliferative response.

**Effect of Antisense OGN on Porcine P2Y<sub>2</sub> Receptor-Mediated Increases in  $[Ca^{2+}]_i$  and Proliferation in Cultured CSMC.** To confirm the involvement of the cloned P2Y<sub>2</sub> receptor in mediating ATP- and UTP-induced proliferation of CSMC, we employed antisense OGN selective for P2Y<sub>2</sub> receptor mRNA (see *Materials and Methods*). Results obtained indicate that pretreatment of the cultured CSMC with pP2Y<sub>2</sub> antisense, but not sense, OGN dramatically suppressed (85%) the increase in  $[Ca^{2+}]_i$  induced by 100  $\mu$ M UTP (Fig. 5A). In contrast, pP2Y<sub>2</sub> antisense OGN did not affect the increase in  $[Ca^{2+}]_i$  induced by LPA, an agonist of another family of G protein-coupled receptors. In subcultured CSMC, pP2Y<sub>2</sub> antisense OGN also inhibited (90%) the increase in  $[Ca^{2+}]_i$  induced by 100  $\mu$ M ATP (Fig. 5B). Consistent with a role for an additional receptor, pP2Y<sub>2</sub> antisense OGN only partially suppressed ATP-induced DNA and protein synthesis, whereas UTP-induced DNA and protein synthesis were

TABLE 1

Agonist potencies and efficacies for the porcine P2Y<sub>2</sub> receptor

Human 1321N1 cells transfected with recombinant plasmid pCR<sup>3.1</sup>-pP2Y<sub>2</sub> were used for Ca<sup>2+</sup> mobilization assays with indicated nucleotide agonists. Data shown are the means ± S.E.M. summarized from the dose-response curves in Fig. 2.

Nucleotide	EC <sub>50</sub> μM	Efficacy % of UTP
UTP	0.53 ± 0.08	100
ITP	1.7 ± 0.5	94.2 ± 6.8
ATP	2.7 ± 0.4	80.1 ± 4.9
GTP	1.3 ± 0.4	63.3 ± 5.4
TTP	8.6 ± 1.6	74.4 ± 6.0
CTP	8.4 ± 1.3	93.6 ± 5.5
UDP	1.5 ± 0.4	35.2 ± 8.7
IDP	35.7 ± 10.8	N.D.
ADP	29.3 ± 9.8	N.D.
TDP	31.4 ± 11.0	N.D.
CDP	87.1 ± 20.3	N.D.
GDP	N.E.	N.E.
Guanosine 5'-O-(3-thio)triphosphate	1.2 ± 0.4	80.2 ± 5.4
Adenosine-5'-O-(3-thio)triphosphate	1.0 ± 0.3	53.6 ± 6.7
2-Methylthio-ATP	N.E.	N.E.
α,β-Methyl-ATP	N.E.	N.E.

N.E., no effect at 100 μM; N.D., not determined because no maximal response was reached.

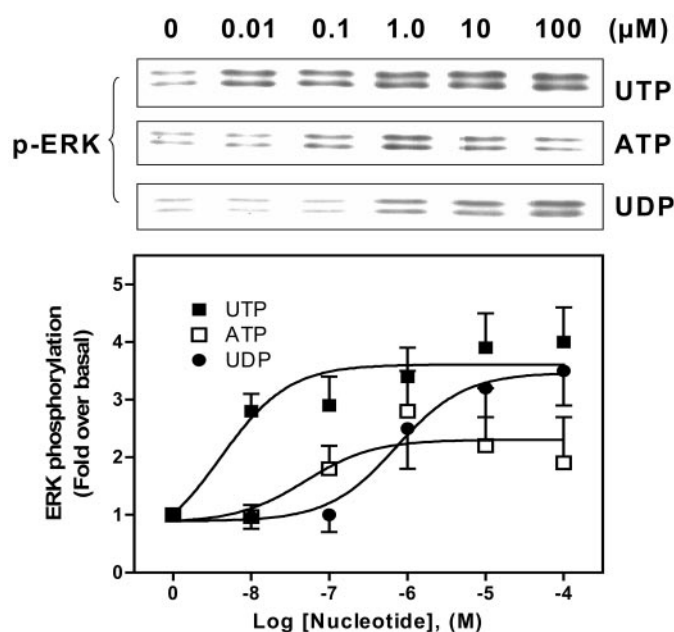
nearly completely inhibited, and responses to LPA were unaffected (Fig. 5, C and D).

**Up-Regulation of Porcine P2Y<sub>2</sub> Receptor mRNA in Cells from in Vitro Organ Cultures and in Vivo Stented Coronary Arteries.** Because our previous work demonstrated the up-regulation of a functional UTP-sensitive P2Y receptor in organ-cultured coronary artery smooth muscles (Hill et al., 2001; Hill and Sturek, 2002), we determined whether the increased activity was associated with the increased expression of mRNA to the P2Y<sub>2</sub> receptor subtype that we have cloned. RT-PCR analysis showed that CSMC isolated from organ-cultured porcine coronary arteries expressed P2Y<sub>2</sub> receptor mRNA to a greater extent than CSMC isolated from freshly harvested or cold-stored porcine coronary arteries (Fig. 6A). In contrast, P2Y<sub>6</sub> receptor mRNA levels were similar in CSMC isolated from organ-cultured and freshly harvested arteries, whereas P2Y<sub>6</sub> receptor mRNA was expressed at lower levels in cold-stored arteries (Fig. 6B). We were unable to detect mRNA for the UTP-sensitive P2Y<sub>4</sub> receptor in either subcultured porcine CSMC or organ-cultured porcine coronary arteries (Fig. 6C).

To determine whether P2Y<sub>2</sub> receptor up-regulation occurs in clinically relevant arterial injury, we assayed for P2Y<sub>2</sub> receptor mRNA expression in a porcine model of coronary artery stent angioplasty. Interestingly, we found that P2Y<sub>2</sub> receptor mRNA levels were significantly increased in CSMC dispersed from stented segments of coronary arteries 3 weeks after stent angioplasty compared with CSMC from unstented segments (Fig. 6D). Again, no significant difference was observed for levels of P2Y<sub>6</sub> receptor mRNA in the stented and unstented artery segments (data not shown), whereas P2Y<sub>4</sub> receptor mRNA was undetectable (Fig. 6C).

## Discussion

In the present study, we report the cloning of the porcine P2Y<sub>2</sub> receptor for the first time and have demonstrated that the porcine P2Y<sub>2</sub> receptor is the first species ortholog that shows a significant difference between the agonist potencies of UTP and ATP. We also have demonstrated that ATP is more potent than UTP in stimulating the proliferation of

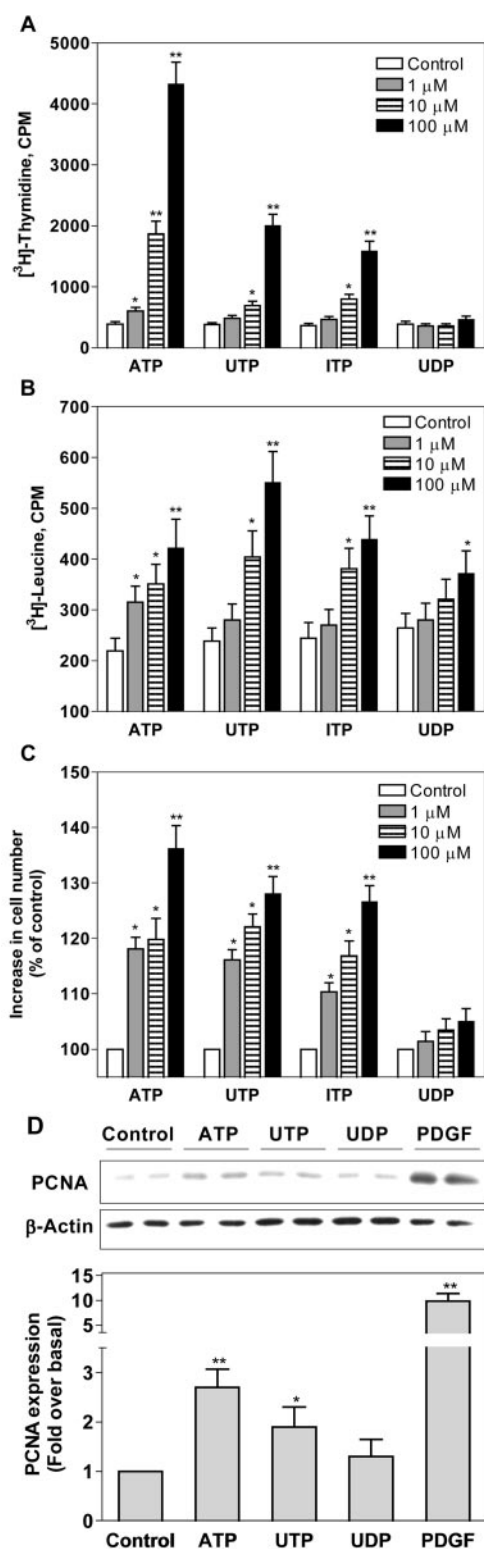


**Fig. 3.** Dose-response of nucleotide-stimulated tyrosine phosphorylation of ERK1/2 in 1321N1 cells expressing porcine P2Y<sub>2</sub> receptors. Confluent cultures of 1321N1 cells expressing pP2Y<sub>2</sub> receptors were serum-starved for 24 h and then stimulated for 5 min with the indicated concentrations of UTP, ATP, or UDP. After treatment, cells were lysed, and phospho-ERK1/2 was detected by Western analysis. Data shown are the means ± S.E.M. of results from three independent experiments.

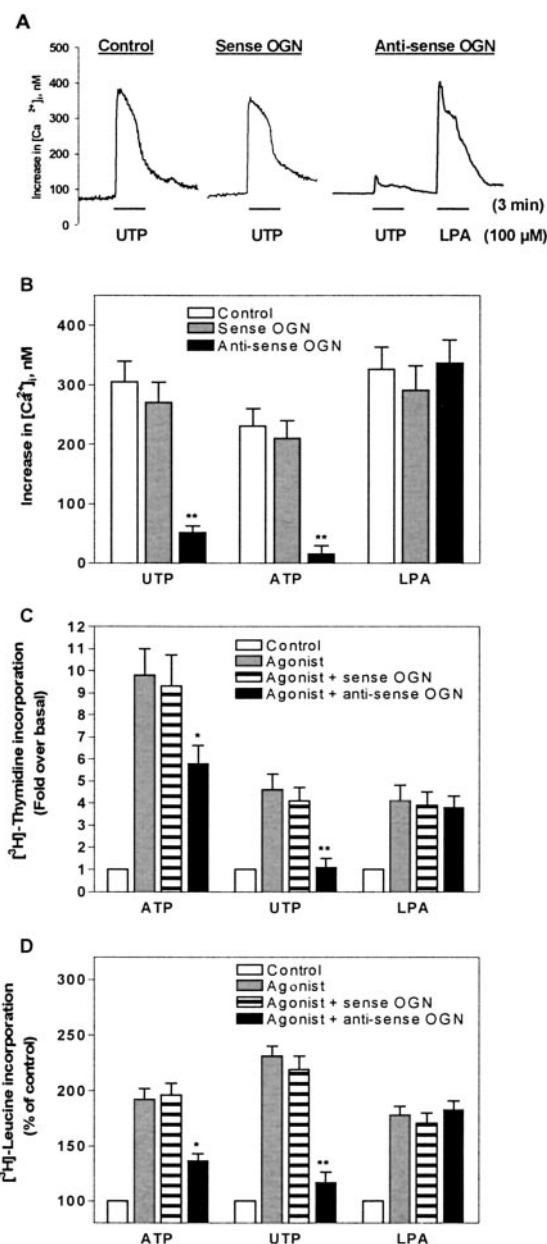
cultured CSMC, suggesting that the P2Y<sub>2</sub> receptor participates with another ATP/ADP/adenosine-related receptor(s) to regulate cell proliferation. These studies further indicated that up-regulation of P2Y<sub>2</sub> receptors occurred in CSMC of organ-cultured porcine coronary arteries and in the clinically relevant model of in vivo coronary artery stent angioplasty.

**Porcine P2Y<sub>2</sub> Receptor Pharmacology.** A P2Y receptor subtype that is more sensitive to UTP than ATP has been reported in cultures of porcine CSMC (Seiler et al., 1999) and in CSMC from organ-cultured porcine arteries (Hill et al., 2001; Hill and Sturek, 2002), and this agonist potency relationship does not match any cloned P2Y receptors. To clone this novel receptor, we used PCR cloning and cDNA amplifi-



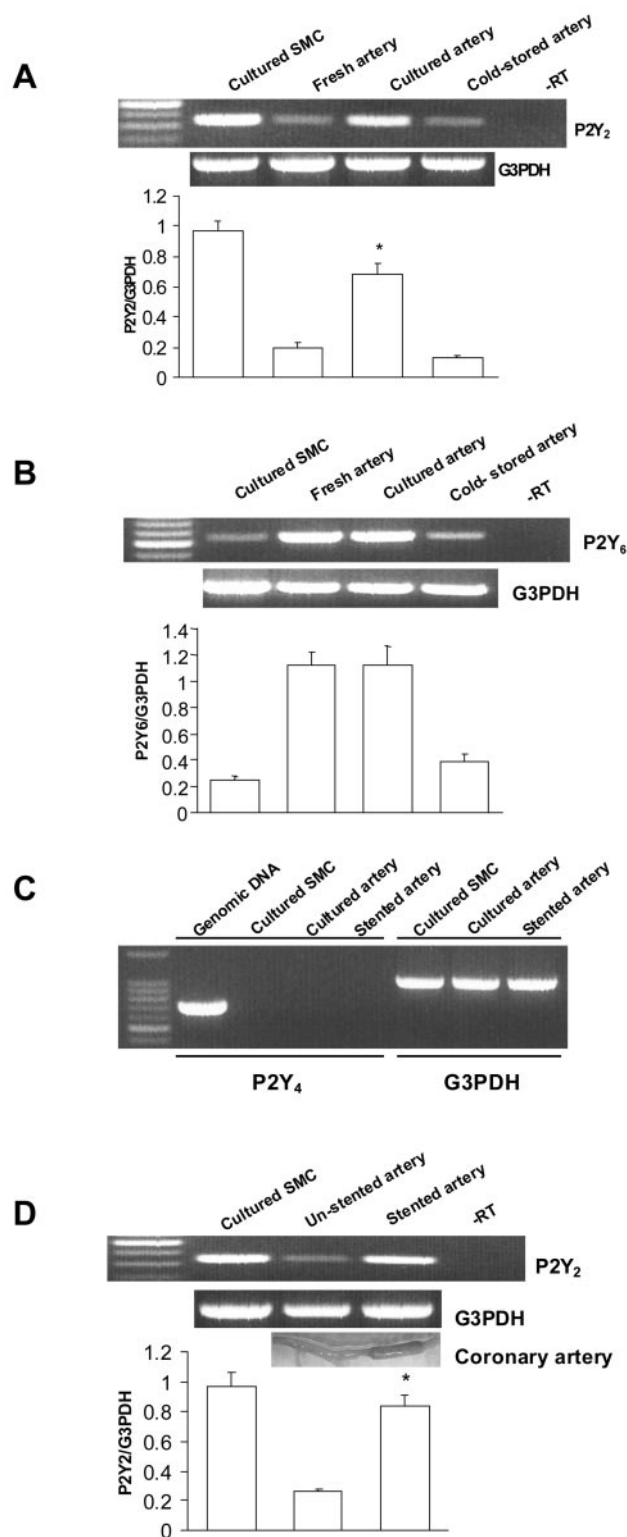


**Fig. 4.** Mitogenic effect of ATP, UTP, ITP, and UDP in cultured coronary artery smooth muscle cells. Cellular DNA (A) and protein synthesis (B) and cell number (C) were determined after stimulation of the CSMC with the indicated concentration of different nucleotides for 48 h (A and B) or 96 h (C). Protein expression levels of PCNA (D) were determined in CSMC stimulated with 100 μM nucleotide or PDGF (100 ng/ml) for 48 h followed by Western analysis. Blots were stripped and reprobed with anti-actin antibody (D). Data shown are the means  $\pm$  S.E.M. of results from four independent experiments performed in triplicate (A, B, and C) or duplicate (D). \*,  $p < 0.05$  and \*\*,  $p < 0.01$  relative to the respective control.



**Fig. 5.** Effect of pP2Y<sub>2</sub> anti-sense OGN on ATP- and UTP-induced increases in  $[Ca^{2+}]_i$  and proliferation in cultured CSMC. Subcultures of CSMC were pretreated with or without sense or anti-sense OGN (3 μM) for 48 h and then incubated with 100 μM ATP, UTP, or LPA for 3 min (A and B) or 48 h (C and D), and  $[Ca^{2+}]_i$  (A and B) and DNA (C) and protein synthesis (D) were determined as described under *Materials and Methods*. Data points in panel B represent the average peak increase in  $[Ca^{2+}]_i$  for 8 to 16 cells on each of five coverslips. Data points in panels C through D are the means  $\pm$  S.E.M. of results from four independent experiments in triplicate (C and D). \*,  $p < 0.05$  and \*\*,  $p < 0.01$  relative to the response of agonist alone.

cation to isolate an open reading frame of the porcine P2Y<sub>2</sub> receptor that has the highest homology to the human P2Y<sub>2</sub> receptor. Pharmacological characterization of the cloned porcine P2Y<sub>2</sub> receptor expressed in 1321N1 cells devoid of endogenous P2Y receptors clearly showed that UTP was a more potent agonist than ATP at increasing  $[Ca^{2+}]_i$  and phosphorylation of ERK1/2, whereas other endogenous nucleoside triphosphates were active and nucleosides except for UDP were inactive. In contrast, UTP and ATP are equipotent and



**Fig. 6.** Up-regulation of P2Y<sub>2</sub> receptor mRNA in organ-cultured and stented porcine coronary artery smooth muscle cells. Semiquantitative RT-PCR for detection of P2Y<sub>2</sub> receptor mRNA expression (340 bp) was performed with cultured CSMC or CSMC dispersed from freshly isolated, organ-cultured, and 5°C cold-stored porcine coronary arteries (A) or from in vivo stented and un-stented coronary artery segments (D), as described under *Materials and Methods*. P2Y<sub>6</sub> receptor mRNA expression (538 bp) was also determined in cultured CSMC or CSMC dispersed from freshly isolated, organ-cultured, and 5°C cold-stored porcine coronary arteries (B). The expression level of porcine P2Y<sub>4</sub> receptor mRNA (686 bp) was below the detection limit in cultured CSMC, intact coronary arteries, and

efficacious agonists of P2Y<sub>2</sub> receptors in human (Parr et al., 1994), mouse (Lustig et al., 1993), rat (Chen et al., 1996), and dog (Zamboni et al., 2000), receptors at which other nucleoside triphosphates are inactive (Dubyak, 2003). Differences in agonist potencies between species orthologs have been reported for other P2Y receptors. For example, the dog P2Y<sub>11</sub> receptor has 74% sequence identity with the human P2Y<sub>11</sub> receptor but different sensitivity for ATP versus ADP (Qi et al., 2001). ATP is a full agonist at the rat P2Y<sub>4</sub> receptor but is a competitive antagonist of the human P2Y<sub>4</sub> receptor (Kennedy et al., 2000). Although the sensitivity of the cloned porcine P2Y<sub>2</sub> receptor to the antagonists suramin, RB-2, and PPADS was similar to the cloned rat P2Y<sub>4</sub> receptor (Kennedy et al., 2000), these antagonists are relatively nonselective. Given the fact that the primary sequence of the cloned porcine P2Y receptor has the highest identity with the human P2Y<sub>2</sub> receptor (84%) and only about 40% homology with either the rat or human P2Y<sub>4</sub> receptor, we have termed it P2Y<sub>2</sub>.

The structural determinants of nucleotide selectivity (UTP versus ATP) at P2Y<sub>2</sub> receptors are unknown, although mutagenesis data suggest that positively charged amino acids in the 6th and 7th transmembrane domains are required for binding the negatively charged phosphate groups of nucleotides (Erb et al., 1995). The porcine P2Y<sub>2</sub> receptor contains positively charged amino acids in these domains, and their precise positioning has never been shown to be important for agonist activity. It seems apparent that only a few amino acid transitions can determine the nucleotide selectivity of the P2Y<sub>4</sub> and P2Y<sub>11</sub> receptors (Qi et al., 2001; Herold et al., 2003); therefore, we suggest that minor differences in the primary sequence underlie variations in the agonist potency profiles between P2Y<sub>2</sub> receptor species orthologs.

We also found that there was greater than a 2-fold difference between the EC<sub>50</sub> values of UTP and ATP for the induction of increases in ERK phosphorylation compared with increases in [Ca<sup>2+</sup>]<sub>i</sub>, although the reason for this difference is not apparent. A possible explanation is that Ca<sup>2+</sup> recording was done in a constant perfusion system that minimizes nucleotide degradation by ecto-ATPases (Kennedy et al., 2000), whereas ERK phosphorylation was determined over 5 min with monolayer cell cultures, where generation of the partial porcine P2Y<sub>2</sub> receptor agonist UDP could occur. It is also noted that ecto-ATPases can degrade ATP at different rates than UTP (Palmer et al., 1998; Kumari et al., 2003), and ADP is not an agonist of the porcine P2Y<sub>2</sub> receptor (Fig. 2C). The possibility that UDP can be converted by cell cultures into UTP (Nicholas et al., 1996) may also contribute to the relative agonist potencies.

**Role of Porcine P2Y<sub>2</sub> Receptors in UTP- and ATP-Induced Proliferation of CSMC.** The mitogenic effect of extracellular nucleotides on VSMC has been known for years (Erlinge, 1998). However, a potent antiproliferative effect of

stented artery segments. The positive control for P2Y<sub>4</sub> receptor mRNA expression was 100 ng of genomic DNA isolated from cultured porcine CSMC (C). RT-PCR performed without reverse transcriptase is indicated as -RT. The piece of artery shown in D indicates the position of stented and unstented fragments of porcine coronary arteries. PCR products were electrophoresed on a 2% agarose ethidium bromide gel, and the relative densities of P2Y<sub>2</sub> (A) and P2Y<sub>6</sub> (B) receptor mRNAs were normalized to G3PDH (983 bp). Data shown are the means ± S.E.M. of results from five (A–C) or four pigs (D). \*, *p* < 0.05 relative to fresh arteries or unstented arteries.



UTP on VSMC also has been reported (White et al., 2000). In either case, the P2 receptor subtype(s) responsible for these effects on the proliferation of VSMC has not been determined. Earlier studies by Erlinge et al. (1993, 1995) showed that ATP or UTP increased DNA and protein synthesis in subcultured rat aortic VSMC. In the same cell culture model, however, Malam-Souley et al. (1993, 1996) were unable to detect increases in DNA synthesis after ATP/UTP stimulation, although ATP or UTP up-regulated the expression of mRNA to several cell cycle progression-related genes. Because P2X agonists were essentially inactive, it was concluded that a P<sub>2U</sub>-like receptor (now termed P2Y<sub>2</sub>) was responsible for the mitogenic effects of ATP/UTP. However, the role of a P2Y<sub>4</sub> receptor cannot be excluded, because the nucleotide agonist profile between rat P2Y<sub>2</sub> and P2Y<sub>4</sub> receptors is essentially indistinguishable (Wildman et al., 2003). Indeed, Harper et al. (1998) suggested that the P2Y<sub>4</sub> receptor mediated ATP/UTP-induced proliferation of rat aortic VSMC. In the present study, we found that ATP, UTP, or ITP, three agonists of the cloned porcine P2Y<sub>2</sub> receptor, increased DNA and protein synthesis and cell number in CSMC. The following evidence supports a role for the P2Y<sub>2</sub> receptor in proliferation of porcine CSMC: 1) UDP, the most potent agonist of P2Y<sub>6</sub> receptors, was inactive in stimulating cell proliferation; 2) antisense OGN for porcine P2Y<sub>2</sub> receptors nearly abolished UTP- or ATP-induced increases in [Ca<sup>2+</sup>]<sub>i</sub>, suggesting that the predominant UTP/ATP-sensitive P2Y receptor functionally expressed in subcultured porcine CSMC is the P2Y<sub>2</sub> subtype; 3) RT-PCR analysis showed that P2Y<sub>2</sub> receptor mRNA was highly expressed in CSMC, whereas P2Y<sub>6</sub> receptor mRNA levels were very low, and P2Y<sub>4</sub> receptor mRNA was undetectable; 4) UTP or ATP, but not UDP, significantly increased expression of PCNA, a cell proliferation marker; and importantly, 5) antisense knockdown of P2Y<sub>2</sub> receptors inhibited UTP- or ATP-induced cellular DNA and protein synthesis. Thus, we have provided several lines of compelling data demonstrating that P2Y<sub>2</sub> receptor plays a role in CSMC proliferation.

It is worth noting that porcine P2Y<sub>2</sub> receptor antisense OGN nearly abolished UTP-induced DNA and protein synthesis but only partially inhibited the mitogenic effect of ATP, whereas the P2Y<sub>2</sub> antisense OGN nearly eliminated UTP- or ATP-induced increases in [Ca<sup>2+</sup>]<sub>i</sub> in cultured CSMC. This inconsistency suggests that another receptor(s) for ATP, ADP, or adenosine might contribute to ATP-induced cell proliferation. The fact that ATP induced higher levels of DNA synthesis and PCNA expression than UTP yet triggered a partial Ca<sup>2+</sup> response strongly suggests that ATP or its hydrolysates can activate other receptors than P2Y<sub>2</sub> to mediate Ca<sup>2+</sup>-independent cell proliferation.

**Selective Up-Regulation of P2Y<sub>2</sub> Receptors in CSMC of Organ-Cultured and Stented Arteries.** Our previous studies have shown that a functional UTP-sensitive P2Y receptor was up-regulated in porcine coronary smooth muscles after several days in organ culture (Hill et al., 2001; Hill and Sturek, 2002). However, the receptor subtype that was up-regulated was not determined. In the present study, we found that P2Y<sub>2</sub> receptor mRNA levels were dramatically increased in CSMC from organ-cultured arteries compared with freshly harvested arteries, whereas P2Y<sub>6</sub> receptor mRNA levels were unchanged, and P2Y<sub>4</sub> receptor mRNA was undetectable. These data suggest that the P2Y<sub>2</sub> receptor is

selectively up-regulated in organ culture, an accepted *ex vivo* model for studying vascular biology. Our findings are consistent with a previous study showing that a short-term culture of rat salivary gland epithelial cells caused up-regulation of P2Y<sub>2</sub> receptors (Turner et al., 1997). The limited expression of P2Y<sub>2</sub> receptor mRNA in control CSMC is consistent with the inability of UTP or ATP to induce increases in [Ca<sup>2+</sup>]<sub>i</sub> in freshly isolated CSMC from porcine coronary arteries (Hill et al., 2001; Hill and Sturek, 2002). Moreover, the cloned porcine P2Y<sub>2</sub> receptor expressed in 1321N1 cells demonstrated similar relative potencies of UTP and ATP as the endogenous receptor in CSMC (Hill and Sturek, 2002), and antagonist sensitivities (i.e., suramin, RB-2, and PPADS) were essentially the same for the endogenous and cloned receptors. Furthermore, we found that UTP desensitized ATP-induced Ca<sup>2+</sup> responses in organ-cultured CSMC (unpublished data), suggesting a common receptor target, presumably P2Y<sub>2</sub>.

The high expression level of P2Y<sub>6</sub> receptor mRNA in CSMC from normal porcine coronary arteries was unexpected, because there was no detectable Ca<sup>2+</sup> response to UDP stimulation in these cells. However, the data are consistent with a previous finding with human coronary artery rings, in which very high levels of P2Y<sub>6</sub> receptor mRNA were detected, yet UDP, the most potent and efficacious agonist of this receptor subtype, failed to induce contraction of the arteries (Malmsjö et al., 2000b). These results suggest that either the P2Y<sub>6</sub> receptor is not expressed at the protein level or perhaps is expressed but fails to couple with Gq protein.

To extend our findings to a more clinically relevant model of arterial injury (for review, see Johnson et al., 1999), we compared P2Y receptor expression in CSMC of stented and unstented segments of coronary arteries. We found that P2Y<sub>2</sub> receptor mRNA levels were significantly increased in the stented CSMC compared with the unstented cells. This result confirms our recent findings with collared rabbit carotid arteries, in which selective up-regulation of P2Y<sub>2</sub> receptors in smooth muscle cells was associated with the development of neointimal hyperplasia (Seye et al., 2002). Collectively, these studies suggest that up-regulation of P2Y<sub>2</sub> receptors after vascular injury may be a general phenomenon in VSMC, which might be important in the development of vascular diseases such as atherosclerosis and angioplasty-induced restenosis and the failure of bypass vein grafts.

In summary, we report the cloning of the porcine P2Y<sub>2</sub> receptor, which is the first species ortholog of P2Y<sub>2</sub> that exhibits different agonist potencies for UTP and ATP. In addition, we demonstrate that the activation of the P2Y<sub>2</sub> receptor is fully responsible for the UTP-induced proliferation of porcine CSMC but only partially involved in the mitogenic effect of ATP, thereby highlighting an additional mechanism for ATP-induced cell proliferation. Last, we show that selective up-regulation of P2Y<sub>2</sub> receptors in porcine CSMC occurs in both *ex vivo* models of vessel organ cultures and in coronary arteries after stent angioplasty *in vivo*.

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**Address correspondence to:** Dr. Michael Sturek, Professor and Chair, Department of Cellular and Integrative Physiology, Indiana University, School of Medicine, 635 Barnhill Drive, MS 309, Indianapolis, IN 46202-5120. E-mail: msturek@iupui.edu