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Evidence for Multiple P2Y Receptors in Trabecular Meshwork Cells

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ABSTRACT

The purpose of this study was to determine if functional purinergic P2 receptors are present in trabecular meshwork cells. The human trabecular cell line (HTM-3) and cultured bovine trabecular cells were used to assess the effects of P2 agonists on intracellular Ca^{2+} levels, extracellular signal-regulated kinase (ERK1/2) activation, and P2Y receptor expression. ATP, UTP, ADP and 2-methyl-thio-adenosine triphosphate (2-MeS-ATP) each produced a concentration-dependent increase in intracellular Ca^{2+} in bovine trabecular cells and the HTM-3 cell line. The addition of UDP did not produce any detectable rise in intracellular Ca^{2+} . Pretreatment with the P2Y₁ receptor antagonist, MRS-2179, blocked the ADP- AND 2-MeS-ATP-induced rise in intracellular Ca^{2+} . However, the ATP- or UTP-induced rise in intracellular Ca^{2+} was not inhibited by MRS-2179 pretreatment. The addition of ADP, 2-MeS-ATP, ATP or UTP were also found to activate the ERK1/2 signaling pathway. This activation of ERK1/2 was blocked by pretreatment with the MEK inhibitor, U-0126, or the PKC inhibitor, chelerythrine chloride, but not by MRS-2179. Analysis of mRNA from HTM-3 cells by RT-PCR revealed the expression of P2Y₁, P2Y₄, and P2Y₁₁ receptor subtypes. These data demonstrate that multiple P2Y receptors are present in trabecular cells. Our results are consistent with the idea that the mobilization of intracellular Ca^{2+} results from the activation of P2Y₁ and P2Y₄ receptors, while the activation of the ERK1/2 pathway results from the activation of P2Y₄ receptors alone. However, a role for the P2Y₁₁ receptors in mobilization of Ca^{2+} , or activation of the ERK1/2 pathway, cannot be discounted.

The existence of two families of purinergic receptors, P2X and P2Y that are activated by adenine or uracil nucleotides was proposed by Abbracchio and Burnstock (Abbracchio and Burnstock, 1994). To date, eight mammalian P2Y receptor subtypes and seven P2X receptor subtypes have been cloned and characterized pharmacologically. In general, mammalian P2Y receptors (i.e., P2Y_{1,2,4,6,11}) are coupled to the G_{q/11} family of G proteins and stimulate the formation of inositol trisphosphate and diacylglycerol with subsequent mobilization of Ca²⁺ from intracellular stores (von Kugelgen and Wetter, 2000). Additionally, P2Y₁₁ receptors have been shown to stimulate adenylyl cyclase. The P2Y₁₂₋₁₄ receptor subtypes are also G-protein coupled receptors, but have been found to primarily associate with Gi/o signaling systems (Abbracchio et al., 2003; Communi et al., 2001; Zhang et al., 2002). The P2X family of receptors function as ligand-gated cation channels (North, 2002).

The presence of adenine nucleotides in the humor of the eye has been known for some time (Greiner et al., 1991). Recent studies have also provided evidence that the activation of ocular P2 receptors can modulate IOP (Pintor et al., 2003). However, little is known about the expression and associated signaling events of P2 purinergic receptor subtypes in anterior segment tissues of the eye. The trabecular meshwork is a specialized region in the anterior chamber of the eye composed of connective tissue beams lined with smooth-muscle-like trabecular meshwork cells (Wiederholt et al., 2000). This meshwork forms the primary pathway for drainage of aqueous humor from the anterior chamber. Cells of the trabecular meshwork are thought to influence intraocular pressure (IOP) through their phagocytic actions (Tripathi and Tripathi, 1984), morphological changes altering intertrabecular space (Wiederholt et al., 2000), and influencing the extracellular matrix turnover (Shearer and Crosson, 2001; Yue, 1996). Consequently,

pharmacologic agents that target trabecular meshwork cells have the potential to regulate outflow resistance and IOP.

In these studies, we sought to determine if trabecular meshwork cells express receptors for adenine and uracil nucleotides, and begin to assess the signal transduction pathways coupled to these receptors. Our results show that trabecular meshwork cells express P2Y₁, P2Y₄, and P2Y₁₁ purinergic receptor subtypes. The activation of these receptors by P2 agonists leads to mobilization of intracellular Ca²⁺ and activation of the ERK1/2 MAP kinase pathway.

METHODS

Materials

Fetal bovine serum was obtained from HyClone Labs (Logan, UT) and DMEM was purchased from GIBCO-BRL (Grand Island, NY). Adenosine 5-triphosphate (ATP), 2-methyl-thio-triphosphate (2-MeS-ATP), adenosine 5-diphosphate (ADP), uridine 5-triphosphate (UTP), uridine 5-diphosphate (UDP), pyridoxal-phosphate-6-azophenyl-2,4-disulphonic acid (PPADS), MRS-2179, suramin, 8-sulfophenylthephylline (8-SPT), chelerythrine chloride and U-0126 were purchased from Sigma Chemical Company (St. Louis, MO). Fluo 3-AM was purchased from Molecular Probes (Eugene, OR).

Cell culture

Primary bovine trabecular cell cultures were established from trabecular meshwork explants by techniques described previously (Shearer and Crosson, 2001). Briefly, small strips of trabecular meshwork tissue were dissected from one or two eyes and homogenized by means of a Teflon hand-held homogenizer in DMEM containing 15% fetal bovine serum (FBS). The homogenized tissue was plated onto a 60 mm collagen-I-coated (Biocoat, Fort Washington, PA) cell culture plate and allowed to grow for 2 wk in DMEM containing 15% FBS. The resultant cells were harvested and plated onto polypropylene cell culture plates in DMEM containing 10% FBS. Second- or third-passaged cells were used in all studies. The transformed human trabecular meshwork cell line (HTM-3) was maintained on polypropylene cell culture plates and grown in DMEM containing 10% FBS (Pang et al., 1994). These cells were allowed to grow to approximately 80% confluence.

Determination of intracellular calcium

Intracellular free Ca^{2+} was determined using a fluorometric imaging plate reader (FLIPR) system (Molecular Devices, Sunnyvale, CA). Cells for intracellular Ca^{2+} measurements were sub-cultured into 96-well clear-bottom black microplates (Corning Costar; Cambridge, MA). On the day of each experiment, cells were incubated with 4 $\mu\text{mol/L}$ fluo 3-AM (excitation at 488 nm, emission at 540 nm; Molecular Probes; Eugene, OR) in HEPES buffer (pH 7.4) containing 2.5 mmol/L probenecid for 1 hr at 37°C. Cells were then washed four-times, placed in the FLIPR and each well of the microplate were monitored at 1.5 sec intervals over 6 minutes. Six wells were averaged for each individual value and experiments were repeated at least three times. In selected experiments, the contribution of extracellular Ca^{2+} to these responses was investigated by omitting Ca^{2+} from the incubation buffer and adding 2 mmol/L EDTA. In cross-desensitization studies, agonist treatments were separated by 2 minutes. For antagonist studies, cells were treated for 10 minutes with individual antagonists prior to the addition of P2 agonists.

Extracellular signal-regulated kinase (ERK) assay

Cells were maintained in serum-free medium for 16 hr before the addition of any agent. Cells were then treated with P2 agonists for 10 minutes. In experiments evaluating the P2 antagonist, U-0126, or chelerythrine chloride, cells were pretreated for 30 min with individual agents prior to the addition of the agonist. At the end of the incubation periods, cells were rinsed with ice-cold PBS and lysed by the addition of 0.5 mL of lysis buffer (50 mmol/L β -glycerophosphate, 20 mmol/L EGTA, 15 mmol/L MgCl_2 , 1 mmol/L Na_3VO_4 , 1 mmol/L dithiothreitol (DTT), and 1 $\mu\text{g/mL}$ of a protease inhibitor cocktail (Roche Diagnostics, Indianapolis, IN). To determine the

level of ERK1/2 activation (phosphorylation), equivalent amounts of protein (15 μ g) were loaded onto 12% SDS-polyacrylamide gels, proteins separated according to molecular weight using standard SDS-PAGE protocols, and transferred to a nitrocellulose membrane. Total ERK levels (phosphorylated and non-phosphorylated forms) were determined by immunoblot techniques using polyclonal anti-ERK1/2 antibodies (New England Biolabs, Inc., Beverly, MA). Bands were visualized by the addition of anti-rabbit HRP-conjugated secondary antibodies and ECL reagents (Amersham Pharmacia Biotech, Piscataway, NJ). Blots were then stripped by incubation in “stripping buffer” (62.5 mmol/L Tris [pH 6.7], 100 mmol/L β -mercaptoethanol, 2% SDS) for 30 min at 50°C. The level of phosphorylated (activated) ERK1/2 was then determined by immunoblot analysis with polyclonal anti-phospho-ERK antibodies (New England Biolabs Inc., Beverly, MA) and visualized by the addition of anti-rabbit HRP-conjugated secondary antibodies and ECL reagents. Band densities were quantified by means of a BioRad Versa Doc Imaging System (Bio-Rad Laboratories, Hercules CA) and the level of phosphorylated ERK1/2 isoforms normalized for differences in loading, using band intensities from immunoblots of total ERK protein.

Reverse transcriptase-polymerase chain reaction

Total RNA was isolated from HTM-3 cells using a Trizol reagent RNA isolation kit (GIBCO BRL, Grand Island, NY, USA) according to the manufacturer’s instruction. Two micrograms of total RNA were reversely transcribed for cDNA synthesis using SuperScript™ RNase H-Reverse Transcriptase and oligo (dT)-12-18 primer (Invitrogen, Carlsbad, CA). Amplifications of targeted purinergic receptor cDNA were performed with specific primers that were designed based on GenBank nucleotide sequences. The PCR was allowed to proceed in a final volume of 20 μ L in a

programmable Master Cycler Gradient Thermocycle (Eppendorf, Mansfield, TX) with the following settings: 5 min at 95°C for initial denaturation followed by repeated cycles of denaturation at 95°C for 3 min, primer annealing for 1 min at 55°C, and extension at 72°C for 1 min 30 sec. After the final cycle, further extension was allowed to proceed for another 10 min at 72°C. The PCR products were resolved on a 1.0% ethidium bromide-stained agarose gel and then visualized under ultraviolet light transillumination. PCR product sizes were estimated from the migration of a DNA size marker run concurrently (1 kb plus DNA Ladder, Invitrogen Life Technologies, Carlsbad, CA). For each sample PCR was performed on RNA that had not been reversely transcribed to confirm that no genomic DNA was present in the samples. Positive reaction products were sequenced to confirm cDNA identity. Primers for each receptor were as follows: P2Y₁ (forward primer) TGTGGTGTACCCCTCAAGTCCC (reverse primer) ATCCGTAACAGCCCAGAATCAGCA P2Y₂ (forward primer) GAGCATCCTGACCTGGAGAG (reverse primer) AGTGCATCAGACACAGCCAG, P2Y₄ (forward primer) CCACCTGGCATTGTCAGACACC (reverse primer) GAGTGACCAGGCAGGGCACGC, P2Y₆ (forward primer) CGCTTCCTCTTCTATGCCAACC (reverse primer) CCATCCTGGCGGCACAGGCGGC, and P2Y₁₁ (forward primer) ACAGAGCGTATAGCCTGGTG (reverse primer) ACTGCGGCCATGTAGAGTAG.

Statistical analysis

Data are presented as the mean, plus and minus the standard error (\pm SE), and were analyzed using analysis of variance followed by Duncan's multiple-range test for detecting differences, with $P < 0.05$ considered as significant. The dose-response curves were analyzed by nonlinear

regression analysis (GraphPad Software, Inc., San Diego, CA).

RESULTS

The effect of P2 agonists on Ca²⁺ mobilization. Exposure of HTM-3 or bovine trabecular cells to ATP, UTP or ADP (10⁻⁶ mol/L) produced a rapid increase in intracellular free Ca²⁺ concentration, peaking in 20-30 sec (Figure 1). The rise in intracellular Ca²⁺ was followed by a return to basal level in 60 to 80 seconds. At equivalent doses ATP and UTP produced similar increases in intracellular free Ca²⁺; however, the rise in Ca²⁺ measured following ADP addition was consistently less than that observed for ATP or UTP. In cells incubated in Ca²⁺-free buffer for 10 min, a rapid increase in intracellular free Ca²⁺ and subsequent decline in response to P2 agonists, was measured (data not shown). Figure 2 shows the concentration-response curve for peak rise in intracellular Ca²⁺ increase following the addition of various P2 agonists to BTM cells. The EC₅₀ and response maxima for P2Y agonists in bovine primary cell cultures and the HTM-3 cell line are listed in Table 1. Except for UDP, all agonists produced a dose-related increase in intracellular Ca²⁺; however, maximum response to ADP and 2-MeS-ATP was 35 to 45% lower than that observed for ATP and UTP.

To further characterize the P2 agonist-induced response in trabecular cells, the effects of P2Y₁-receptor antagonist, MRS-2179, and the nonselective P2 antagonists, suramin and PPADS, were evaluated. The increases in intracellular-free Ca²⁺ induced by 2-MeS-ATP and -ADP were blocked by the presence of MRS-2179 (10 μmol/L) (Figure 3). However, the ATP- and UTP-induced increase in Ca²⁺ mobilization was not altered by the presence of MRS-2179.

Pretreatment of cells with nonselective P2 antagonists, suramin and PPADS (10 μmol/L) each significantly inhibited the ADP- and 2-MeS-ATP-induced rise in intracellular Ca²⁺ by 60 to 80%,

but did not significantly alter the responses to ATP or UTP. Pretreatment with the adenosine receptor antagonist, 8-SPT (10 $\mu\text{mol/L}$), did not significantly alter the rise in intracellular Ca^{2+} induced by any of the P2 agonists (data not shown).

To evaluate cross-desensitization between ATP and UTP, agonists were administered sequentially within a 2 minute interval. The addition of ATP (10 $\mu\text{mol/L}$) did not alter the subsequent addition of UTP. However, the addition of UTP (10 $\mu\text{mol/L}$) reduced the response to ATP by 26% ($P < 0.05$).

Effects of P2 agonists on ERK1/2 activation. As shown in Figure 4, the addition of 10^{-7} mol/L ATP, UTP, ADP, and 2-MeS-ATP to bovine trabecular meshwork cells produced a significant increase in ERK1/2 phosphorylation. No increase in ERK1/2 activation was observed following the addition of UDP. In HTM-3 cells the addition of ATP, UTP, ADP and 2-MeS-ATP increased ERK1/2 activation by 233, 228, 247, and 190%, respectively. In both culture systems, the increase in ERK1/2 activation induced by each P2 agonist was not altered by pretreatment with MRS-2179, PPADS or suramin (data not shown).

To investigate the up-stream signaling events associated with P2 agonist-induced stimulation of ERK1/2, cells were pretreated with the MEK inhibitor, U-0126, or the PKC inhibitor, chelerythrine chloride. As shown in Figure 5, pretreatment of BTM cells with U-0126 (1.0 mol/L) blocked the ERK activation induced by ATP or UTP (10^{-7} mol/L). Pretreatment with the PKC inhibitor, chelerythrine chloride (20 $\mu\text{mol/L}$), also completely blocked the ATP- and UTP-induced ERK1/2 activation in these cells. In HTM-3 cells, pretreatment with U-0126 or

chelerythrine also completely blocked the ERK1/2 activation induced by ATP or UTP (10^{-7} mol/L).

Expression of P2Y receptor subtype mRNA in BTM and HTM-3 cells. To investigate the expression of P2Y-receptor subtypes, mRNA from human cell line (HTM-3) was analyzed by RT-PCR. As shown in Figure 6, mRNA for P2Y₁, P2Y₄, and P2Y₁₁ receptors was detected in HTM-3 cells. However, no message for P2Y₂ and P2Y₆ receptors could be detected in these cells.

DISCUSSION

Adenine nucleotides have been identified in the aqueous humor of animals (Greiner et al., 1991). However, our understanding of the role P2 receptors play in regulating anterior segment function has been limited. Although P2X receptors have been identified in the mammalian retina (Jabs et al., 2000; Wheeler-Schilling et al., 2001; Wheeler-Schilling et al., 2000), the expression of P2X receptors in anterior segment tissues have not been reported. In contrast, molecular and functional studies have provided evidence that P2Y receptors are expressed in a number of anterior segment tissues including corneal and ciliary epithelium, lens and conjunctiva (Collison and Duncan, 2001; Cowlen et al., 2003; Cullinane et al., 2001; Farahbakhsh and Cilluffo, 2002; Merriman-Smith et al., 1998). In this study, we investigated if trabecular meshwork cells express functional P2 receptors.

The addition of a P2 agonist to human or bovine trabecular cells produced a rapid rise in intracellular Ca^{2+} . This increase in intracellular Ca^{2+} did not appear to result from the activation of ionotropic P2X receptors, as it was not blocked by the incubation of cells in Ca^{2+} -free media. Although adenosine receptors have been identified on trabecular meshwork cells (Shearer and Crosson, 2002), the inability of the adenosine antagonist, 8-SPT, to alter the response to adenine and uracil nucleotides demonstrates that the activation of adenosine receptors did not contribute to responses observed in these studies. Taken together, these data support the idea that the responses induced by P2 agonists in trabecular cells are mediated by P2Y receptors.

The difference in response maxima (see Table 1) and the selective blockade of P2 agonists responses by MRS-2179 (see Figure 3), indicate that the activation of at least two P2Y-receptor subtypes can mobilize intracellular Ca^{2+} in trabecular cells. The moderately selective P2Y₁ agonists, ADP and 2-MeS-ATP, exhibited response maxima that were 35 to 45% lower than values determined for ATP and UTP. The EC₅₀ values measured for ADP and 2-MeS-ATP are consistent with the EC₅₀ for P2Y₁ receptors measured in other mammalian tissues (Nicholas et al., 1996; Pacaud et al., 1996; Palmer et al., 1998; Ralevic and Burnstock, 1996; Simon et al., 1995). In addition, the ADP- or 2-MeS-ATP-induced increase in intracellular Ca^{2+} was blocked by the pretreatment of P2Y₁ antagonist, MRS-2179. These data support the idea that both ADP- and 2-MeS-ATP stimulate the release of intracellular Ca^{2+} in trabecular cells through activation of the P2Y₁ receptor. The expression of P2Y₁ receptors in the HTM-3 cell line was confirmed by RT-PCR. The precise signal transduction pathways employed by P2Y₁ receptors in these cells are yet to be fully explored. Our results are consistent with other studies showing that P2Y₁ receptors are coupled to an increase in intracellular Ca^{2+} through a G_{q/11}/IP3 signaling system.

The increase in intracellular Ca^{2+} following the addition of UTP provides evidence that bovine and human trabecular cells also express one or more of the conventional uracil-sensitive P2Y receptors (i.e., P2Y₂, P2Y₄ or P2Y₆). The lack of any measurable increase in intracellular Ca^{2+} following UDP administration, and the absence of any detectable mRNA in RT-PCR analysis demonstrates that this response is not due to the activation of P2Y₆ receptors in these cells. As RT-PCR analysis also failed to detect P2Y₂ receptors in the human HTM-3 cell line, and that the calculated EC₅₀ for UTP is similar to that reported for P2Y₄ receptors in other systems, the responses to UTP in these cells appears to result from P2Y₄ receptor activation. However, recent

studies have shown that UTP can mobilize intracellular Ca^{2+} by activating P2Y_{11} receptors (White et al., 2003). As RT-PCR analysis did detect P2Y_{11} receptor in HTM-3 cell line, we can not exclude the possibility that UTP mobilizes intracellular Ca^{2+} via the activation of P2Y_{11} receptors.

Studies have shown that P2Y_4 receptors derived from rat and human cells can be activated by both ATP and UTP (von Kugelgen and Wetter, 2000). However, in P2Y_4 receptors derived from human cells ATP potency is normally less than that measured for UTP (von Kugelgen and Wetter, 2000). In this study, the calculated EC_{50} , response maxima, and antagonist sensitivity for ATP were very similar to that determined for UTP, for both human- and bovine-derived cells. Cross-desensitization studies also demonstrated that ATP treatment did not alter subsequent responses to UTP. These studies provide evidence that ATP and UTP responses are mediated by separate receptors. As ATP responses were not blocked by P2Y_1 antagonist, MRS-2179, the ATP-induced increase in intracellular Ca^{2+} appears to be mediated by P2Y_{11} receptors expressed in these cells. However, the final determination will require the development of selective P2Y receptor antagonists.

Activation of the ERK1/2 pathway in trabecular cells has been shown to regulate matrix metalloproteinase secretion and cellular proliferation (Shearer and Crosson, 2001; Alexander and Acott, 2003; Jeon, 2003). In addition, the agents that alter IOP have been shown to regulate ERK1/2 activity in the trabecular meshwork (Shearer and Crosson, 2002). Hence, the activation of the ERK1/2 signaling pathway appears to play a central role in the regulation of trabecular function (Shearer and Crosson, 2002). The addition of ATP, UTP, ADP, or 2-MeS-ATP each

activated ERK1/2 in HTM-3 and bovine trabecular cells. The ATP- and UTP-induced activation of ERK1/2 was blocked by pretreatment with the MEK inhibitor, U-0126, or the PKC antagonist, chelerythrine chloride. These results are consistent with the idea that activation of P2Y receptors leads to mobilization of intracellular Ca^{2+} , activating calcium-sensitive PKC and eventually activating ERK1/2 in these cells. Unlike the Ca^{2+} mobilization studies, no significant difference in response magnitude was measured in these experiments. The activation of ERK1/2 by UTP in these studies indicates that uracil-sensitive P2Y₄ receptors are linked to ERK activation.

Additionally, the inability of MRS-2179 to inhibit ERK activation induced by ADP or 2-MeS-ATP argues that the P2Y₁ receptor is not linked to this pathway. However, we cannot exclude the possibility that ERK activation observed following the addition of nucleotides also results from the activation of P2Y₁₁ receptors.

Original studies considered the trabecular meshwork a passive filter for aqueous humor, with changes in resistance to aqueous flow occurring indirectly through ciliary muscle contraction. Recent studies have demonstrated that trabecular cells are contractile in nature and are capable of modifying the extracellular matrix within the region, supporting the idea that these cells participate in the regulation of outflow resistance and IOP (Shearer and Crosson, 2001; Wiederholt et al., 2000; Yue, 1996). As the elevation in intracellular Ca^{2+} regulates trabecular cell contractility, and ERK has been shown to regulate matrix metalloproteinases, P2Y receptors may play an important role in regulation of trabecular function and aqueous outflow resistance. Recent studies have shown that P2 receptors modulate IOP (Pintor, 2003). Therefore, we speculate that pharmacological agents targeting trabecular P2Y receptors may prove to be efficacious agents for the treatment of glaucoma.

In summary, these data demonstrate that multiple P2Y receptors are present in human and bovine trabecular meshwork cells. Our results are consistent with the idea that the activation of P2Y₁, P2Y₄ and P2Y₁₁ receptors leads to the mobilization of intracellular Ca²⁺. The stimulation of the ERK1/2 signaling pathway appears to result from the activation of P2Y₄ receptors via a PKC-dependent system. However, a role for the P2Y₁₁ receptor in the activation of this pathway cannot be excluded.

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FOOTNOTES PAGE

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LEGENDS FOR FIGURES

Figure 1. Examples of P2 agonists-induced changes in trabecular intracellular Ca²⁺ levels.

Trabecular cells were loaded with 4 μmol/L fluo 3-AM in HEPES buffer (pH 7.4) for 1 hour at 37°C and fluorescence monitored (see methods). Wells were treated with 10⁻⁶ mol/L of indicated nucleotide at t = 12 seconds. Panel **A** show the responses from primary cultures of bovine trabecular cells. Panel **B** shows responses from HTM-3 cell line.

Figure 2. Concentration-response curves for P2 agonist-induced increase in intracellular

Ca²⁺. Bovine trabecular meshwork cells were treated with a single P2 agonist and fluorescence monitored (see methods). For each agonist concentration, the difference between the peak response and pre-stimulus baseline was calculated as the percent of control. Values represent the means ± SE from 5 experiments.

Figure 3. Effect of the P2Y₁ antagonist, MRS-2179, on the P2 agonist-induced increase in

intracellular Ca²⁺. Bovine trabecular meshwork cells were pretreated with MRS-2179 (10 μmol/L) for 10 minutes prior to the addition of P2 agonists. Asterisks denote significant difference (*P* < 0.05) between agonist stimulation alone and agonist stimulation following pretreatment with MRS-2179 (n = 4).

Figure 4. Effects of P2 agonists on ERK 1/2 activation.

Serum-deprived bovine trabecular cells were incubated for 10 minutes in the presence or absence (control) of individual P2 agonists (0.1 μmol/L). Panel **A** shows summary data from 5 experiments. Values are the means ± SE of

densitometry measurements from immunoblots cell lysates. Asterisks denote significant difference ($P < 0.05$) from control levels. Panel **B** shows representative immunoblots of phospho-ERK and total ERK from bovine trabecular cell lysates.

Figure 5. Inhibition of ERK1/2 activation by the MEK inhibitor U0126 and the PKC inhibitor chelerythrine. Representative immunoblots of phospho-ERK and total ERK from bovine trabecular cell lysates. Panel **A** shows responses from control cells, cells treated for 10 minutes with UTP (0.1 $\mu\text{mol/L}$) in the presence or absence of U0126 or chelerythrine for 30 minutes. Panel **B** shows responses from control cells, cells treated for 10 minutes with ATP (0.1 $\mu\text{mol/L}$) in the presence or absence of U0126 or chelerythrine for 30 minutes. Pathway inhibitors were added 30 minutes prior to the addition of ATP or UTP.

Figure 6. RT-PCR analysis of P2Y-receptor subtype expression in HTM-3 cells. For each panel, Lanes 1, 2 and 3 contain: RT product; non-RT product and genomic DNA, respectively.

TABLE PAGE

Table 1. Dose-response parameters for P2 agonist-induced mobilization of intracellular Ca^{2+} in trabecular meshwork cells.

Agonist	Bovine Trabecular Cells		HTM-3 Cell Line	
	EC ₅₀	Response	EC ₅₀	Response
	($\mu\text{mol/L}$)	Maximum (% Increase)	($\mu\text{mol/L}$)	Maximum (% Increase)
ATP	0.58	428	0.27	287
UTP	0.37	419	0.20	267
ADP	0.35	269	0.053	177
2-MeS-ATP	0.070	225	0.048	172
UDP	ND	ND	ND	ND

Parameters for UDP were not determined (ND) as this agonist did not produce any significant elevation in intracellular Ca^{2+} at the concentrations tested.

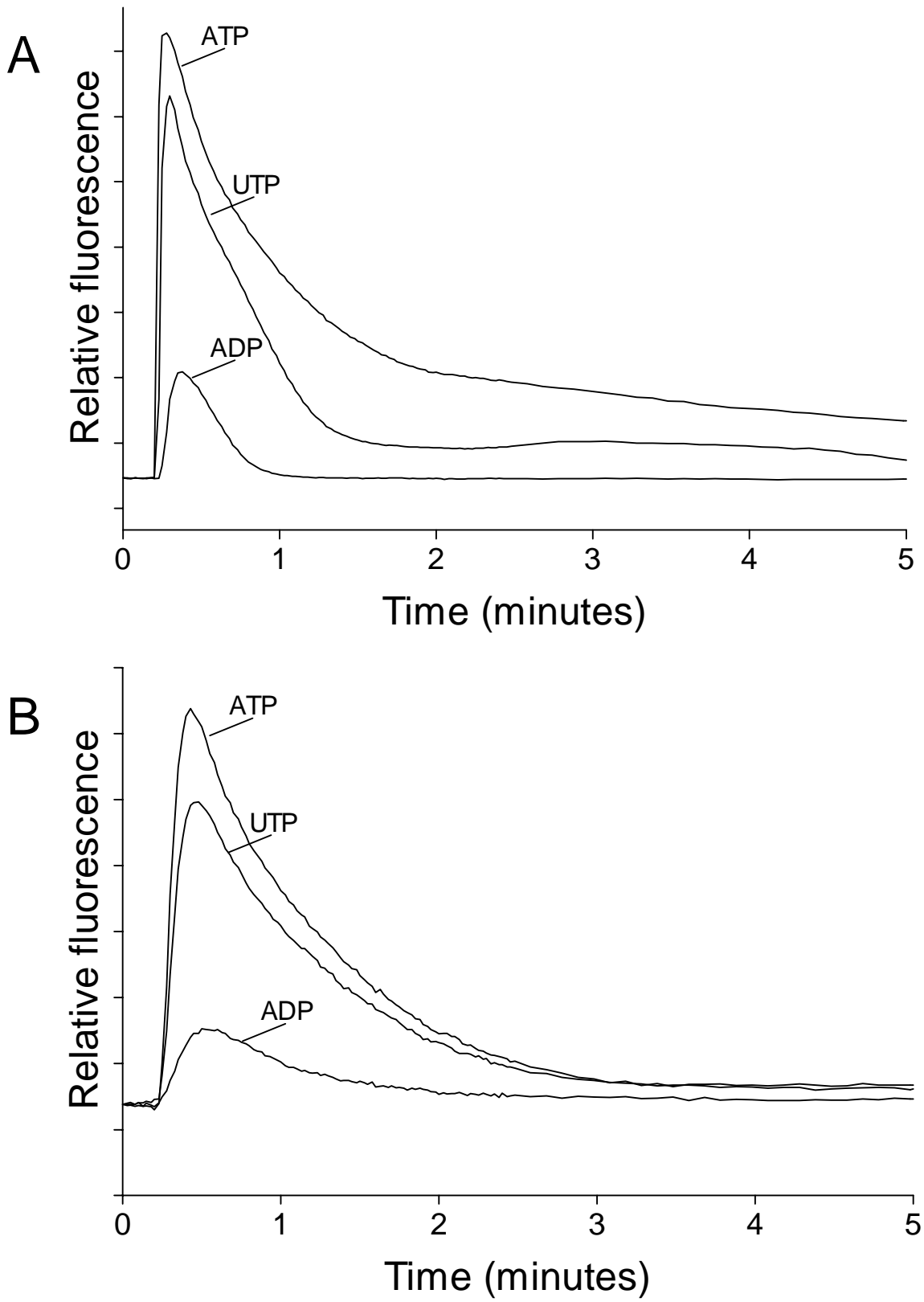


Figure 1

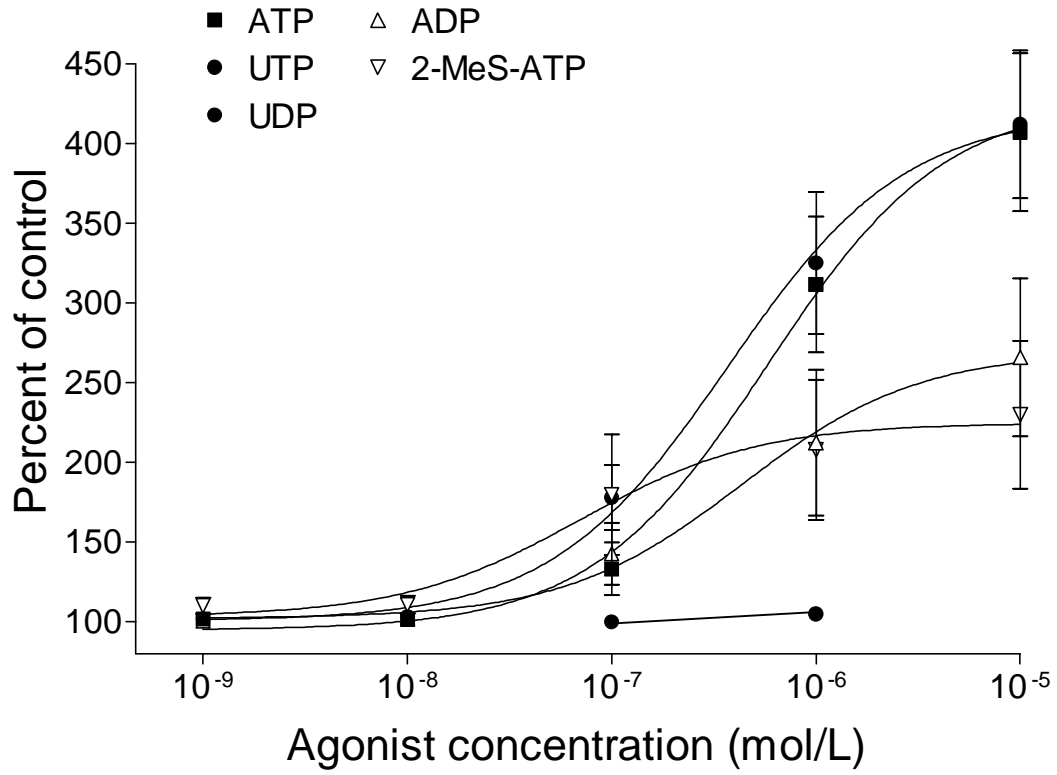


Figure 2

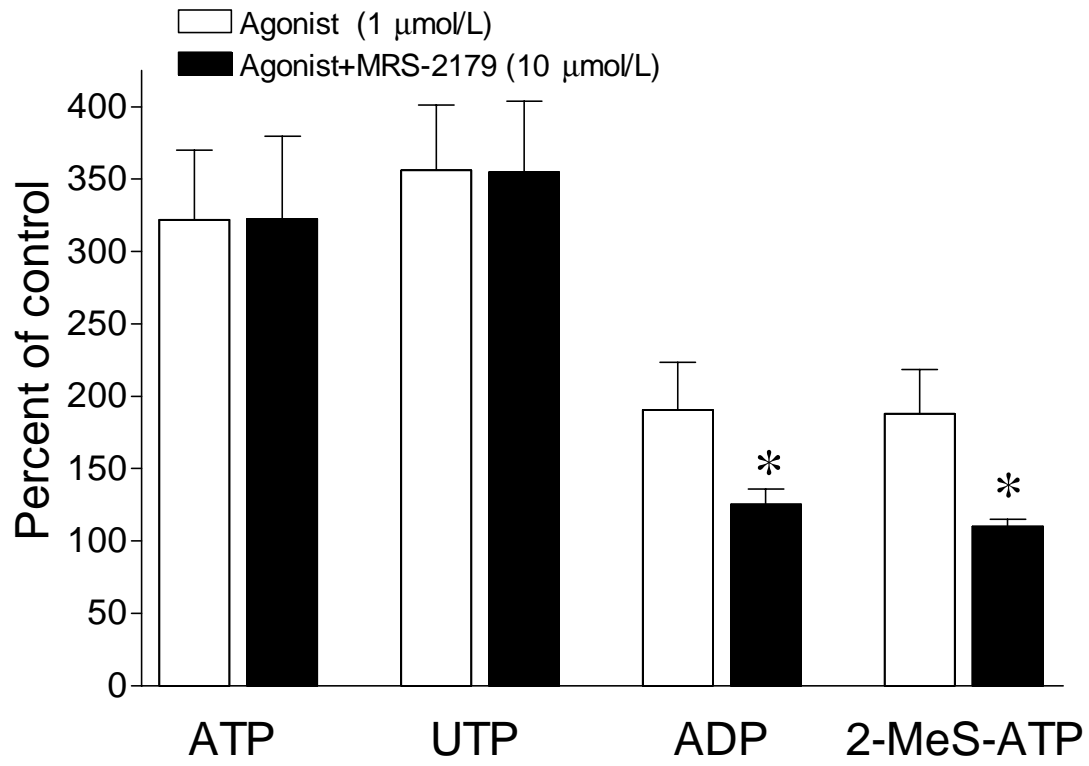


Figure 3

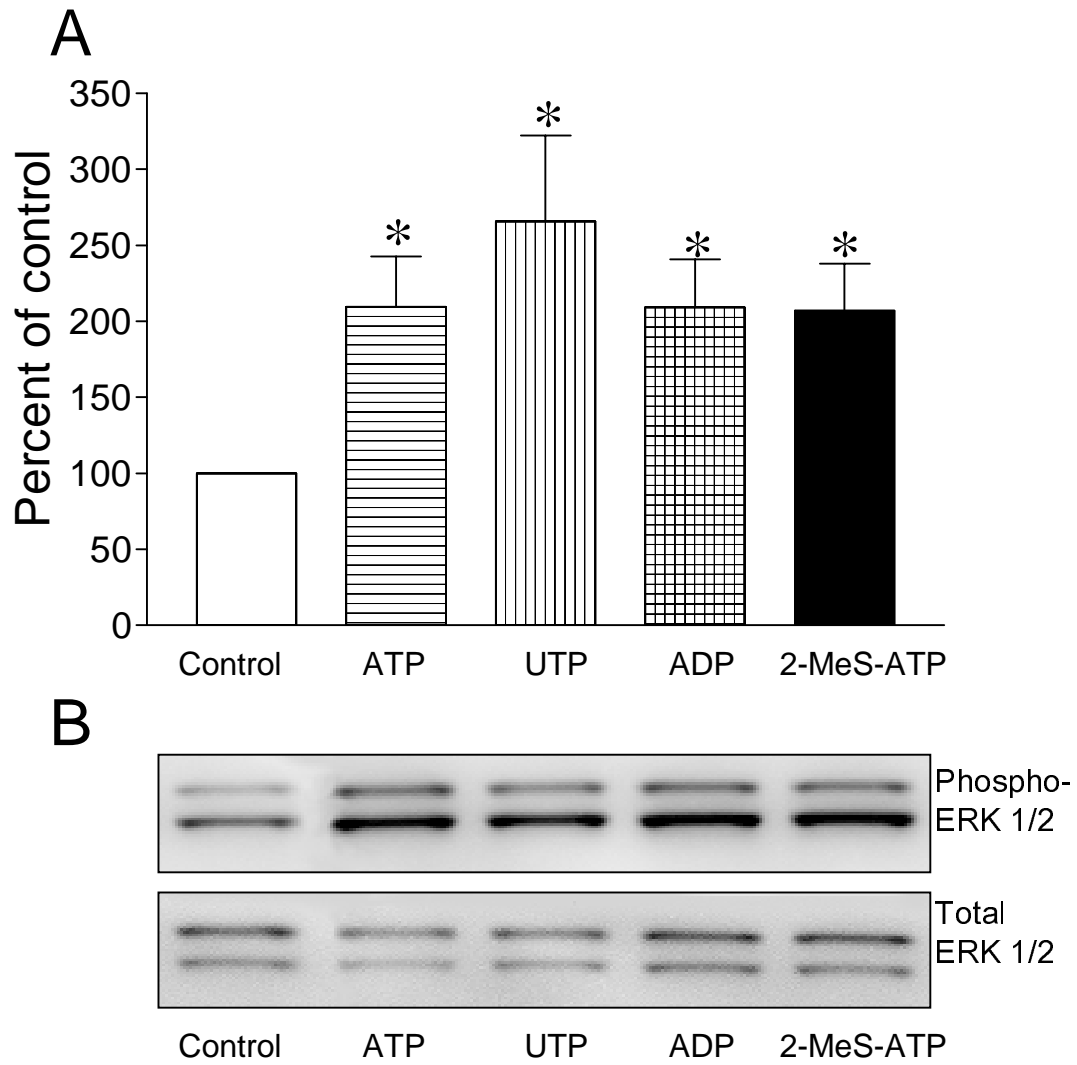


Figure 4

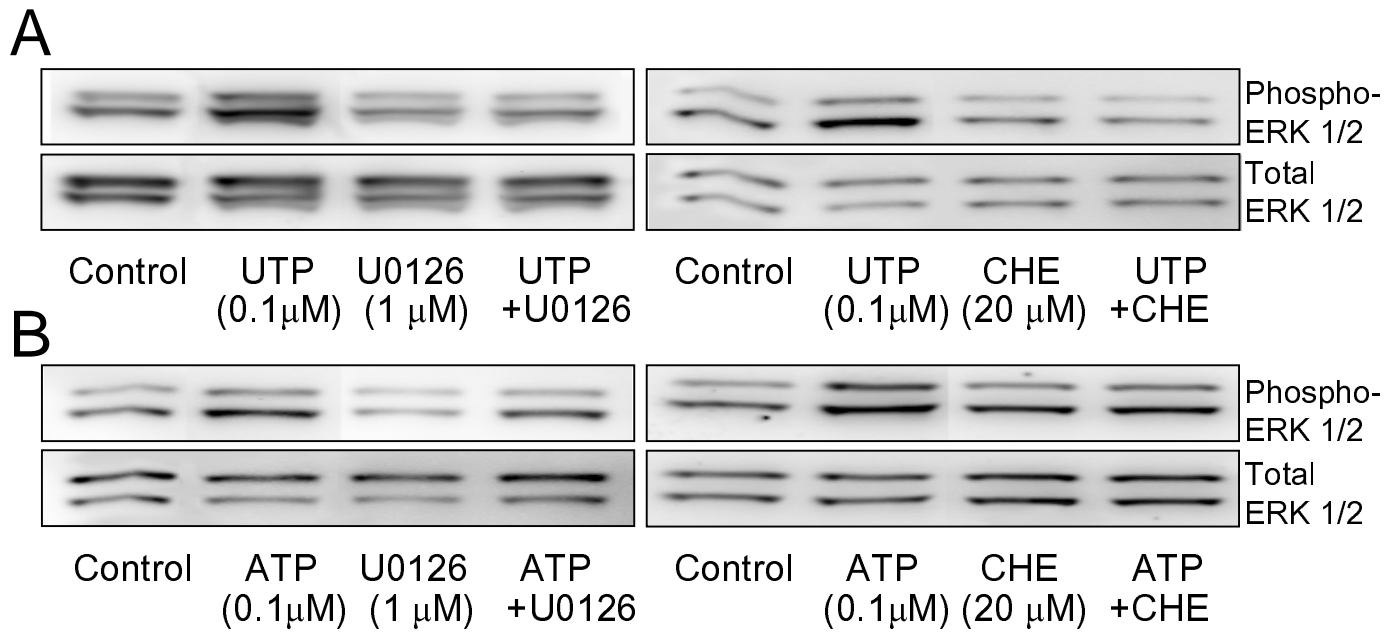


Figure 5

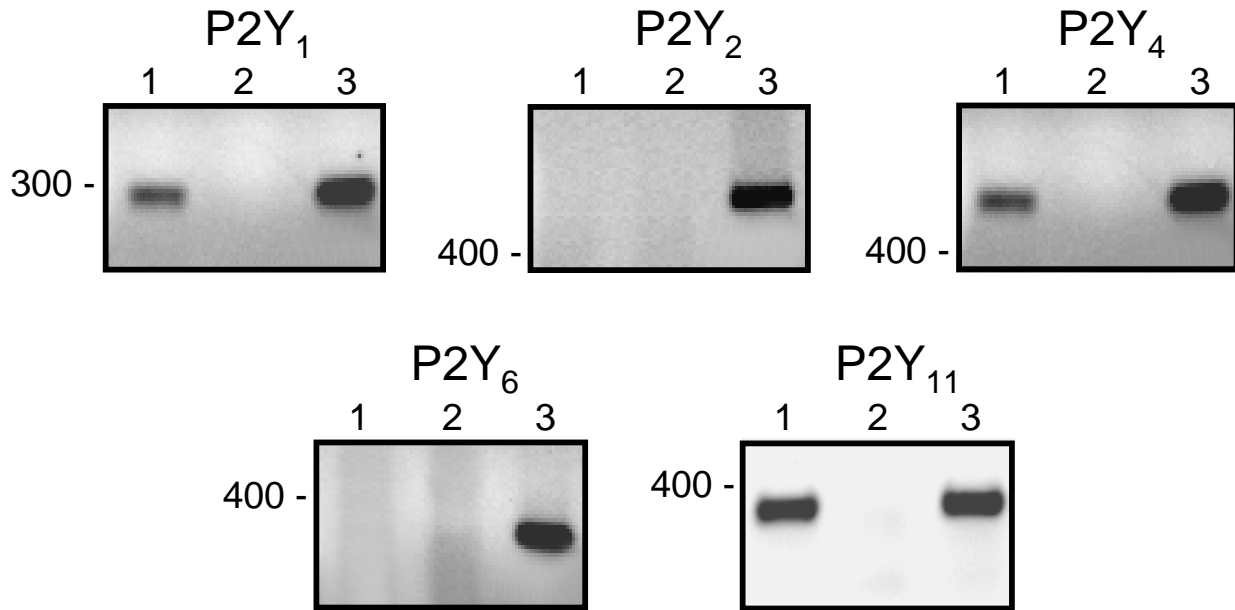


Figure 5