A Pyridoxine Cyclic Phosphate and Its 6-Azoaryl Derivative Selectively Potentiate and Antagonize Activation of P2X₁ Receptors

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Analogues of the P2 receptor antagonists pyridoxal-5'-phosphate and the 6-azophenyl-2',4'disulfonate derivative (PPADS), in which the phosphate group was cyclized by esterification to a CH₂OH group at the 4-position, were synthesized. The cyclic pyridoxine- $\alpha^{4,5}$ -monophosphate, compound \hat{z} (MRS 2219), was found to be a selective potentiator of ATP-evoked responses at rat P2X₁ receptors with an EC₅₀ value of $5.9 \pm 1.8 \,\mu$ M, while the corresponding 6-azophenyl-2',5'-disulfonate derivative, compound 3 (MRS 2220), was a selective antagonist. The potency of compound **3** at the recombinant P2X₁ receptor (IC₅₀ 10.2 \pm 2.6 μ M) was lower than PPADS (IC₅₀ 98.5 \pm 5.5 nM) or iso-PPADS (IC₅₀ 42.5 \pm 17.5 nM), although unlike PPADS its effect was reversible with washout and surmountable. Compound 3 showed weak antagonistic activity at the rat P2X₃ receptor (IC₅₀ 58.3 \pm 0.1 μ M), while at recombinant rat P2X₂ and P2X₄ receptors no enhancing or antagonistic properties were evident. Compounds 2 and 3 were found to be inactive as either agonists or antagonists at the phospholipase C-coupled P2Y₁ receptor of turkey erythrocytes, at recombinant human $P2Y_2$ and $P2Y_4$ receptors, and at recombinant rat $P2Y_6$ receptors. Similarly, compounds 2 and 3 did not have measurable affinity at adenosine A_1 , A_{2A}, or A₃ receptors. The lack of an aldehyde group in these derivatives indicates that Schiff's base formation with the P2X₁ receptor is not necessarily required for recognition of pyridoxal phosphate derivatives. Thus, compounds 2 and 3 are relatively selective pharmacological probes of P2X₁ receptors, filling a long-standing need in the P2 receptor field, and are also important lead compounds for future studies.

Extracellular adenine, and likely uracil, nucleotides have a physiological role to play in the central and peripheral nervous systems through their activation of P2 receptors.^{1,2} Two families of P2 receptors have been defined:³ ligand-gated cation channels (P2X subtype), activated by adenine nucleotides, and G-protein-coupled receptors (P2Y subtype), most of which are activated by both adenine and uracil nucleotides.⁴ P2X₁₋₇ and P2Y_{1,2,4,6} designations have been unambiguously assigned to mammalian nucleotide receptors,^{5,6} although there is still uncertainty about the correspondence of these cloned sequences to the pharmacological phenotypes of native P2 receptors.

ATP acts as a cotransmitter with norepinephrine and other transmitters in sympathetic neurotransmission in mammals.⁷ In vas deferens, isolated blood vessels, intestine, kidney, and skin in a number of species, norepinephrine and ATP cause synergistic constriction via α_1 -adrenoceptors and P2X receptors (primarily the P2X₁ subtype), respectively. In rabbit coronary vessels, guinea pig taenia coli, rat aorta, and rat mesenteric artery, the predominant effect of those transmitters is relaxation via β -adrenoceptors and P2Y receptors. Besides sympathetic neurotransmission, P2 receptors also function in parasympathetic, sensory-motor, nonadrenergic noncholinergic (NANC) inhibitory, and somatic neuromuscular neurotransmission. For example, it appears that activation of the P2X₃ receptor subtype mediates nociception via the dorsal root ganglia; thus a selective antagonist may prove to be anti-nociceptive.^{8,9} The therapeutic potential and physiological role of P2 receptors in various central and peripheral biological systems has been reviewed.¹⁰

Progress in the field of P2 receptors has been impeded by the lack of stable, selective, and bioavailable ligands, especially antagonists.¹¹ Synthetic polyanionic diazo derivatives of pyridoxal-5-phosphate (1a, Figure 1), PPADS (**1b**, pyridoxal- α^5 -phosphate-6-azophenyl-2',4'disulfonic acid), and iso-PPADS (1c, the 2,5-disulfonate isomer) were shown to be P2 receptor antagonists.¹² Pyridoxal-5'-phosphate, itself, is a weak antagonist of P2 receptors.¹³ In smooth muscle assays PPADS irreversibly antagonized P2X receptors in rabbit vas deferens,¹⁴ urinary bladder,¹⁵ isolated blood vessels,¹⁶ guinea pig isolated vas deferens,¹⁷ and perfused rat mesenteric arterial bed.¹⁸ PPADS is a relatively nonselective antagonist at P2 receptors, since it also acts at P2Y₁ receptors¹⁹ with a K_i value of approximately 1 μ M. PPADS does not antagonize the action of ATP agonists at P2X₄ and P2X₆ receptors and has an IC₅₀ of 45 µM at P2X₇ receptors.²⁰ Furthermore, PPADS has

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Figure 1. Structures of pyridoxal-5'-phosphate (**1a**) and its azo derivatives (**1b**, **1c**) and the cyclic pyridoxine- $\alpha^{4.5}$ -monophosphate derivatives (**2**, MRS 2219 and **3**, MRS 2220) described in the present study.

a low affinity at P2Y₂ and P2Y₆ receptors,²¹ at the P2Y_{AC} receptor in human platelets,^{21,22,40} and at the adenylate cyclase-coupled P2Y receptor in rat C6 glioma cells.²¹

Results

Analogues of the P2 receptor antagonists pyridoxal-5'-phosphate and 6-azophenyl-2',4'-disulfonate derivative (PPADS), in which the phosphate group was cyclized by esterification to a CH₂OH group at the 4-position, were synthesized (Figure 1) with the aim of developing more potent and selective antagonists for P2 receptor subtypes. Cyclic pyridoxine- $\alpha^{4,5}$ -monophosphate, **2** (MRS 2219), and cyclic pyridoxine- $\alpha^{4,5}$ -monophosphate-6-azophenyl-2',5'-disulfonic acid, 3 (MRS 2220), were prepared and characterized using NMR and highresolution mass spectroscopy, and purity of >98% was demonstrated using high-pressure liquid chromatography (HPLC). The possibility of contamination of **3** with the acyclic pyridoxine- α^5 -phosphate-6-azophenyl-2',5'disulfonic acid was ruled out using HPLC. The latter substance was prepared through sodium borohydride reduction of 1c, and no corresponding peak was observed in the HPLC trace of **3**, with a detection limit of 0.1%. Incubation of **3** in the buffer used for the P2X bioassay (see Materials and Methods) also failed to generate this acyclic compound.

The compounds were tested in a functional ion channel assay^{23,24} of ATP-induced current at recombinant rat P2X₁, P2X₂, P2X₃, and P2X₄ receptors, expressed in *Xenopus* oocytes, using the twin-electrode voltage-clamping technique. P1 and P2 receptors are present on the follicle cell layer of *Xenopus* oocytes,²⁵ but our experiments were carried out on defolliculated oocytes to avoid activation of endogenous receptors. In the control uninjected, defolliculated oocytes, treatment with ATP (<100 μ M) did not induce any current. Compounds **1a** and **1b** were previously reported to antagonize agonist-induced cation flux at recombinant P2X₁ receptors, with the IC₅₀ values of roughly 10 and 1 μ M, respectively.²⁶ Compounds **1a** was not evaluated in the present study; however we have found compounds



log [antagonist], M

Figure 2. Effects of compounds **1b** (PPADS, squares) and **1c** (iso-PPADS, triangles) on inward current induced by activation by 3 μ M ATP of recombinant rat P2X₁ receptors, expressed in *Xenopus* oocytes, using the twin-electrode voltage-clamping technique (pH 7.5, Ba²⁺ Ringer's solution). IC₅₀ values for compounds **1b** and **1c** (n = 4) were 98.5 ± 5.5 and 42.5 ± 17.5 nM, respectively. All data points were mean ± SEM of four observations. The apparent lack of error bars on some points is due to the size of symbols being greater than the size of error bars.

1b and **1c** to be considerably more potent than anticipated in inhibiting the inward current elicited by 3 μ M ATP in defolliculated oocytes expressing recombinant P2X₁ receptors, with IC₅₀ values of 98.5 ± 5.5 and 42.5 ± 17.5 nM, respectively (Figure 2). In oocytes expressing rat P2X₃ receptors, IC₅₀ values were 240 ± 38 nM (**1b**) and 83.5 ± 3.6 nM (**1c**).

The novel cyclic phosphate structurally related to compound 1c, i.e., compound 3, also antagonized activation of recombinant P2X₁ receptors by ATP (Figure 3B), while compound **2**, related to compound **1a**, selectively potentiated ATP-evoked responses at P2X₁ receptors effectively doubling the current (Figure 3A) with an EC₅₀ value of 5.9 \pm 1.8 μ M. Unlike PPADS, the effect of 1b at P2X₁ receptors was readily reversible upon washout and surmountable. Furthermore, compound **3** showed weak antagonistic activity at rat P2X₃ receptors (Figure 3B). The IC_{50} values for compound 3 at P2X₁ and P2X₃ receptors were 10.2 \pm 2.6 and 58.3 \pm 0.1 μ M, respectively, in the presence of 3 and 1 μ M ATP, respectively. Thus, compound 3 was 6-fold selective for P2X₁ vs P2X₃ receptors. Compounds **2** and **3** were both completely inactive as either potentiators or antagonists at P2X₂ or P2X₄ receptors (Figure 3). Therefore, although a less potent antagonist of P2X receptors than PPADS, the effect of **1b** was selective for the $P2X_1$ subtype.

Antagonism of phospholipase C (PLC) activity^{27,28} induced by activation of single subtypes of P2Y receptors was studied. At the P2Y₁ receptor of turkey erythrocytes, there was no significant effect on activation by 10 nM 2-MeSATP by compound **2** or **3** at concentrations up to 100 μ M (Figure 4). Furthermore, compounds **2** and **3** were found to be essentially inactive in stimulating, potentiating, or blocking PLC at recombinant human P2Y₂ and P2Y₄ receptors and at rat P2Y₆ receptors (Figure 4). Thus, compounds **2** and **3** were completely inactive as either agonists or antagonists of these four subtypes of P2Y receptors.

Affinity at adenosine (P1) receptors was examined in radioligand binding experiments at rat brain A_1 and A_{2A}



Figure 3. Effects of compounds **2** (A) and **3** (B) on inward current induced by activation by ATP, at the indicated concentrations, of recombinant rat P2X₁ (3 μ M), P2X₂ (10 μ M), P2X₃ (1 μ M), and P2X₄ (30 μ M) receptors, expressed in *Xenopus* oocytes, using the twin-electrode voltage-clamping technique. The agonist concentrations correspond approximately to the EC₇₀ values. IC₅₀ values for compounds **3** (n = 4) at P2X₁ and P2X₃ areceptors were 10.2 ± 2.6 and $58.3 \pm 0.1 \mu$ M, respectively. Slopes of the curves were 0.8 ± 0.1 and 0.9 ± 0.1 , respectively. All data points were mean \pm SEM of four observations. The apparent lack of error bars on some points is due to the size of symbols being greater than the size of error bars.

adenosine receptors and at recombinant human A_3 adenosine receptors expressed in HEK293 cells.²⁹ Neither compound **2** nor **3** at 100 μ M caused any significant displacement of [³H]R- N^6 -phenylisopropyladenosine at rat A_1 , [³H]-2-[4-[(2-carboxyethyl)phenyl]ethylamino]-5'-N-ethylcarbamoyladenosine at rat A_{2A} , or [¹²⁵I]N⁶-(4amino-3-iodobenzyl)-5'-N-methylcarbamoyladenosine at human A_3 receptors (data not shown). Thus, these two compounds display complete selectivity for the P2 over the P1 receptors.

Discussion

In the present study we describe a structural modification, a reduction of the aldehyde group and cyclization of the pyridoxal-5-phosphate moiety of pyridoxal phosphate-related P2 receptor antagonists, that apparently results in molecules that exhibit complete selectivity for P2X over P2Y receptors, and further selectivity for the P2X₁ subtype within the P2X class of signaling proteins. By virtue of this modification, the analogue corresponding to iso-PPADS, **3**, retains antagonist



Figure 4. Lack of either antagonism or potentiation (n = 3) by compounds **2** and **3** (100 μ M) of activation of phospholipase C activity induced at turkey erythrocyte P2Y₁ receptors (by 10 nM 2-MeSATP), recombinant human P2Y₂ receptors (by 100 nM UTP), recombinant human P2Y₄ receptors (by 100 nM UTP), and at recombinant rat P2Y₆ receptors (by 100 nM UDP).

properties at this subtype, while the analogue corresponding to pyridoxal-5-phosphate, **2**, becomes a potentiator of activation of this subtype. Compound **3** also weakly antagonized ATP action at recombinant P2X₃ receptors. Evans et al.²⁰ reported that PPADS blocks ATP action at recombinant P2X₃ receptors expressed in HEK293 cells with an IC₅₀ of 2 μ M.

PPADS also potentiated UTP responses in cell lines possessing P2U receptors and pyrimidinoceptors, but by an action considered to involve ectoATPase inhibition.³⁰ This action may explain the modest increase by compound **3** of uridine nucleotide activity at recombinant hP2Y₂ and hP2Y₄ receptors (see Figure 4).

While selective antagonists have been reported for $P2Y_{AC}$ and $P2Y_1$ receptors,^{22,29} this is the first example of a highly selective antagonist (compound **3**) for any P2X receptor subtype. Furthermore, although potentiation by various antagonists of the effects of P2X receptor agonists have been reported previously (e.g., suramin at P2X receptors in enteric neurons and PPADS at recombinant P2X₄ receptors),^{31,32} compound **2** displays unique selectivity as a potentiator of the P2X₁ subtype. Defolliculated oocytes contain no ectoATPases;³³ therefore the possibility that compound **2** could be acting as a potentiator through enzymatic inhibition can be ruled out.

The dramatic qualitative difference in activity between the structurally related compounds **2** and **3** may provide clues in the molecular modeling of the receptors³⁴ to distinguish native and activated conformations of this ion channel. It will be useful to study the effects of these compounds at mutant P2X receptors^{35,36} in order to form a hypothesis for the amino acids responsible for antagonist/potentiator binding.

Antagonism by PPADS at native P2X receptors in whole tissue bioassays is nonsurmountable and often reversed slowly,^{14–18} if at all, consistent with irreversible antagonism. However, blockade by iso-PPADS at P2X receptors on rat vagus nerve³⁷ and blockade by PPADS at recombinant P2X₂ receptors expressed in oocytes²³

are fully reversible within 1-2 h. The possibility of Schiff's base formation between the aldehyde group on the ligand and a P2X receptor to explain the slow reversibility was raised by Buell et al.³⁵ and Collo et al.³⁸ PPADS blockade was slowly reversible, but not fully reversible, at recombinant P2X_{1,2,5} receptors (expressed in HEK293 cells). PPADS blockade reversed rapidly at recombinant P2X₃ receptors (expressed in HEK293 cells), whereas PPADS only showed irreversible blockade at $P2X_{46}$ receptors at 100-fold higher concentrations. It was hypothesized that the aldehyde of PPADS might form a Schiff's base with lysine residues of P2X₁ and P2X₂ receptors. Scrutiny of the structures of P2X₁₋₆ receptors drew attention to a lysine on P2X_{1.2.5} receptors at a site equivalent to position 249 on P2X₄ receptors and position 251 on P2X₆ receptors. A threonine is present at the equivalent position on P2X₃ receptors. Substitution of glutamate with lysine (E249K) at P2X₄ receptors enhanced the antagonist potency of PPADS by 30-fold, although blockade was slowly reversible. The same substitution at the equivalent position (L251K) at P2X₆ receptors also enhanced PPADS sensitivity with slow reversibility. Substitution of lysine with glutamate (K246E) at P2X₂ receptors increased the rate of recovery of PPADS blockade. None of these procedures affected the blocking activity of the reversible antagonist suramin at these recombinant P2X receptors. At $P2X_1$ receptors, the conserved lysine would be unreactive with compounds 2 and 3, since the aldehyde group is absent. Thus, Schiff's base formation between the ligand and receptor is not necessarily required for recognition of pyridoxal phosphate derivatives at $P2X_1$ receptors. The absence of an aldehyde group in compound 3 may be responsible for the complete reversibility of its antagonism.

The pharmacological properties of compounds **2** and **3** at putative heterooligomers of the $P2X_1$ receptors, such as have already been demonstrated for $P2X_2/P2X_3$ subtypes,³⁹ should be examined.

Further structure activity studies are in progress to improve the affinity of compounds **2** and **3** at the P2X₁ receptor. Such a selective P2X₁ receptor antagonist derived from **3** may have potential utility in controlling receptor-mediated contraction of visceral and vascular smooth muscle (e.g., vascular hypertension and instability of the urinary bladder detrusor muscle). A selective enhancer of P2X₁ receptor activity derived from **2** may have potential utility in enhancing contractions at P2X₁ receptor-controlled muscle sphincter tone (e.g., treatment of urinary incontinence) and sympathetic vascular tone. In conclusion, these novel derivatives are useful pharmacological probes, filling a long-standing need in the P2 receptor field, and are also important lead compounds for future studies.

Materials and Methods

Synthesis. Pyridoxine and the reagents for azo coupling reactions were purchased from Aldrich (St. Louis, MO). Aniline-2,5-disulfonic acid was obtained from K & K Laboratories, Inc. (Hollywood, CA).

Proton nuclear magnetic resonance spectroscopy was performed on a Varian GEMINI-300 spectrometer, and spectra were taken in D_2O . The chemical shifts are expressed in ppm relative to the HOD peak at 4.78 ppm.

³¹P NMR spectra were recorded without proton decoupling mode, at room temperature using Varian XL-300 spectrometer (121.419 MHz); orthophosphoric acid (85%) was used as an external standard. High-resolution FAB (fast atom bombardment) mass spectra were determined with a JEOL SX102 spectrometer, and electron spray mass spectra were obtained using a Hewlett-Packard 1100 LC-ESPRAY system. Determination of purity were performed with a Hewlett-Packard 1090 HPLC system using an OD-5-60 C18 analytical column (250 $mm \times 4.6 mm$, Separation Methods Technologies, Inc., Newark, DE) in two different linear gradient solvent systems. One solvent system (A) was 0.1 M triethylammonium acetate buffer: $CH_3CN = 95:5$ to 40:60 for 20 min with flow rate 1 mL/min. The other (B) was 5 mM tetrabutylammonium phosphate buffer: $CH_3CN =$ 80:20 to 40:60, in 20 min with flow rate 1 mL/min. Peaks were detected by UV absorption using a diode array detector.

Cyclic Pyridoxine- $\alpha^{4,5}$ -monophosphate (2). To a suspension of 0.1 g (0.59 mmol) of pyridoxine in 2 mL of anhydrous benzene was added 1 mL of trimethylsilyl polyphosphate under nitrogen atmosphere. The mixture was stirred at 40 °C for 2 days and poured into 10 mL of anhydrous ether. The white precipitate was collected by filtration, washed with anhydrous ether, and dissolved in 2 mL of water. The water solution was stirred at 40 °C for 30 min to remove the trimethylsilyl group and passed through Amberlite CG-50 resin (H⁺ form, weakly acidic) with the elution of water (flow rate 0.5 mL/min). The pure fractions were combined and lyophilized. The solid residue was washed with a minimum of water, and 0.05 g of white powdered product was obtained (yield 37%). ¹H NMR (D₂O): δ 2.62 (3H, s, CH₃), 5.14 (2H, d, J = 15.6 Hz, CH₂O), 5.34 $(2H, d, J = 15.6 \text{ Hz}, CH_2O), 8.08 (1H, s, H-6).$ ³¹P NMR (D₂O): 4.62 (pen, J = 15.9 Hz). MS: negative FAB, 230 (M - H), positive FAB, 232 (M + H); negative API-ES, 230 (M - H). HRMS (FAB-): calcd 230.0218, found 230.0216. HPLC retention time: 4.0 min using solvent system A, 5.4 min using solvent system B (purity >98%).

Cyclic Pyridoxine- $\alpha^{4,5}$ -monophosphate-6-azophenyl-2',5'-disulfonic Acid (3). To a solution of 0.055 g (0.216 mmol) of aniline-2,5-disulfonic acid in 2 mL of water and 0.22 mL of 1 N HCl was added 0.015 g (0.216 mmol) of solid sodium nitrite at 0 °C. This solution was stirred for 5 min, and the pH was adjusted to ~ 10 with 1 N NaOH. To the mixture was added dropwise a solution of 0.05 g (0.216 mmol) of 2, previously dissolved in aqueous NaOH (pH \sim 10). The pH was adjusted to \sim 9, and the yellow color changed to red. After 30 min of stirring 0 °C, the mixture was purified by ion-exchange column chromatography using Amberlite CG-50 resin (H⁺ form, weakly acidic) with the elution of water (flow rate 0.5 mL/min). The red fraction showing a single peak in HPLC was collected and lyophilized to give 0.08 g of the disodium salt form of the desired compound (yield 69%). ¹H NMR (D₂O): δ 2.56 (3H, s, CH₃), 5.24 (2H, d, J = 15.6 Hz, CH₂O), 5.70 (2H, d, J = 15.6 Hz, CH₂O), 7.89 (1H, bs, phenyl), 8.08-8.14 (2H, m, phenyl). ³¹P NMR (D₂O): 5.56 (pen, J =15.9 Hz). HRMS (FAB-): calcd 493.9729, found

493.9721. HPLC: 7.7 min using solvent system A, 12.4 min using solvent system B (purity >98%).

Pyridoxine-α⁵-**phosphate**-6-**azophenyl**-2',5'-**disulfonic Acid (HPLC Standard).** To a stirred solution of 0.027 g of **1c** (0.05 mmol, Tocris Cookson Inc., St. Louis, MO) in 2 mL of H₂O was added 0.004 g of solid NaBH₄ (0.1 mmol) at room temperature. The mixture was stirred for 30 min and passed through Amberlite CG-50 resin (H⁺ form, weakly acidic) with water elution (flow rate 0.5 mL/min). The pure fractions of the major component, indicated by single peak in HPLC, were collected and lyophilized (0.015 g, 58%). ¹H NMR (D₂O): δ 2.70 (3H, s, CH₃), 5.21 (2H, s, CH₂OH), 5.61 (2H, d, J = 6.8 Hz, CH₂OP), 7.88–8.23 (3H, m, phenyl). ³¹P NMR (D₂O) 0.05 (t, J = 6.8 Hz). HPLC: 5.9 min using solvent system A. HRMS (FAB–): calcd 511.9835, found 511.9828.

Pharmacology: Antagonist Activity at Recombinant P2X Receptors. *Xenopus* oocytes were harvested and prepared as previously described.²³ Defolliculated oocytes were injected cytosolically with rat P2X₁, P2X₂, P2X₃, or P2X₄ receptor cRNA (40 nL, 1 μ g/mL), incubated for 24 h at 18 °C in Barth's solution, and kept for up to 12 days at 4 °C until used in electrophysiological experiments.

ATP-activated membrane currents ($V_{\rm h} = -90$ mV) were recorded from cRNA-injected oocytes using the twin-electrode voltage-clamp technique (Axoclamp 2B amplifier). Voltage recording (1–2 M Ω tip resistance) and current-recording microelectrodes (5 M Ω tip resistance) were filled with 3.0 M KCl. Oocytes were held in an electrophysiological chamber and superfused with Ringer's solution (5 mM/min, at 18 °C) containing (mM) NaCl, 110; KCl, 2.5; HEPES, 5; BaCl₂, 1.8, adjusted to pH 7.5.

ATP (at the EC₇₀ values in μ M for respective subtypes: P2X₁ 3, P2X₂ 10, P2X₃ 1, and P2X₄ 30) was superfused over oocytes for 60–120 s and then washed out for a period of 20 min. For inhibition curves, data were normalized to the current evoked by ATP at pH 7.5. Test substances were added for 20 min prior to ATP exposure; all compounds were tested for reversibility of their effects. The concentration required to inhibit the ATP-response by 50% (IC₅₀) was taken from Hill plots constructed using the formula log($I/I_{max} - I$), where *I* is the current evoked by ATP in the presence of an antagonist. Data are presented as Mean \pm SEM (n =4) for data from different batches of oocytes.

Phospholipase C Assay at P2Y Receptors. P2Y₁ receptor-promoted stimulation of inositol phosphate formation by 2-MeSATP (10 nM) was measured in turkey erythrocyte membranes as previously described.^{25,26} The values were averaged from three to eight independent determinations. Briefly, 1 mL of washed turkey erythrocytes was incubated in inositolfree medium (DMEM; Gibco) with 0.5 mCi of 2-[³H]myoinositol (20 Ci/mmol; American Radiolabelled Chemicals Inc., St. Louis, MO) for 18-24 h in a humidified atmosphere of 95% air 5% CO₂ at 37 °C. Erythrocyte ghosts were prepared by rapid lysis in hypotonic buffer (5 mM sodium phosphate, pH 7.4, 5 mM MgCl₂, 1 mM EGTA) as described.²⁶ PLC activity was measured in 25 μ L of [³H]inositol-labeled ghosts (~175 μ g of protein, 200–500000 cpm/assay) in a medium containing 424 μM

CaCl₂, 0.91 mM MgSO₄, 2 mM EGTA, 115 mM KCl, 5 mM KH₂PO₄, and 10 mM Hepes, pH 7.0. Assays (200 μ L final volume) contained 1 μ M GTP γ S and the indicated concentrations of nucleotide analogues. Ghosts were incubated at 30 °C for 5 min, and total [³H]inositol phosphates were quantitated by anion exchange chromatography as previously described.^{25,26}

Stimulation of inositol phosphate formation in 1321N1 human astrocytoma cells stably expressing recombinant human $P2Y_2$ receptors (activated by 100 nM UTP), recombinant human $P2Y_4$ receptors (activated by 100 nM UTP), and recombinant rat $P2Y_6$ receptors (activated by 100 nM UDP) was measured in a similar fashion.

Abbreviations. ATP, adenosine 5'-triphosphate; HEK, human embryonic kidney; Hepes, *N*-[2-hydroxyethyl]piperazine-*N*-[3-propanesulfonic acid]; HPLC, highpressure liquid chromatography; HRMS, high-resolution mass spectroscopy; *K*_i, equilibrium inhibition constant; iso-PPADS, pyridoxal- α^5 -phosphate-6-azophenyl-2',5'disulfonic acid; MeATP, adenosine-5'-methylenetriphosphate, (α,β) or (β,γ) isomers; 2-MeSATP, 2-methylthioadenosine-5'-triphosphate; MRS 2219, cyclic pyridoxine- $\alpha^{4,5}$ -monophosphate; MRS 2220, cyclic pyridoxine- $\alpha^{4,5}$ -monophosphate-6-azophenyl-2',5'-disulfonic acid; MS, mass spectrum; PLC, phospholipase C; PPADS, pyridoxal- α^5 -phosphate-6-azophenyl-2',4'-disulfonic acid; SAR, structure–activity relationship; TLC, thin-layer chromatography.

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