

International Union of Pharmacology LVIII: Update on the P2Y G Protein-Coupled Nucleotide Receptors: From Molecular Mechanisms and Pathophysiology to Therapy

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Abstract—There have been many advances in our knowledge about different aspects of P2Y receptor signaling since the last review published by our International Union of Pharmacology subcommittee. More receptor subtypes have been cloned and characterized and most orphan receptors deorphanized, so that it is now possible to provide a basis for a future subdivision of P2Y receptor subtypes. More is known about the functional elements of the P2Y receptor molecules and the signaling pathways involved, including interactions with ion channels. There have been substantial developments in the design of selective agonists

and antagonists to some of the P2Y receptor subtypes. There are new findings about the mechanisms underlying nucleotide release and ectoenzymatic nucleotide breakdown. Interactions between P2Y receptors and receptors to other signaling molecules have been explored as well as P2Y-mediated control of gene transcription. The distribution and roles of P2Y receptor subtypes in many different cell types are better understood and P2Y receptor-related compounds are being explored for therapeutic purposes. These and other advances are discussed in the present review.

I. Brief Historical Background of Nucleotides and Their Receptors

The first description of the extracellular signaling by purines was by Drury and Szent-Györgyi (1929), and purinergic receptors were defined in 1976 (Burnstock, 1976). After an early hint (Spedding and Weetman, 1976), receptors for purines were subdivided into P1 (adenosine) and P2 (ATP and ADP) receptors (Burnstock, 1978), and later subdivision of P2 receptors into P2X and P2Y subtypes was made on the basis of pharmacology (Burnstock and Kennedy, 1985). It was recognized that some P2Y receptors responded to pyrimidines as well as purines (Seifert and Schultz, 1989). After cloning of P2 receptors and studies of transduction mechanisms in the early 1990s, the basis for subdivision into P2X and P2Y receptor families was confirmed and extended (Abbracchio and Burnstock, 1994) and seven subtypes of P2X receptors and eight subtypes of P2Y receptors are currently recognized (Ralevic and Burnstock, 1998; North, 2002; Burnstock, 2004).

II. Molecular Structure of P2Y Receptors

A. Nomenclature and Molecular History of P2Y Receptors

Regarding the currently used nomenclature, P2Y is used for functional mammalian receptor proteins and functional nonmammalian species. The lower case, p2y, is used for mammalian orphan receptors or functional nonmammalian receptors without a mammalian ortholog. The subscript number (1–n) following P2Y or p2y sequentially list proteins in their chronological order of cDNA cloning. The first P2 receptors were cloned in 1993 (Lustig et al., 1993; Webb et al., 1993). They corresponded to receptors previously characterized by pharmacological criteria: P2Y₁ (formerly P_{2Y}) and P2Y₂ (formerly P_{2U}). Since then several other subtypes were isolated by homology cloning and assigned a subscript

on the basis of cloning chronology (P2Y₄, P2Y₆, and P2Y₁₁). The long-awaited G_i-coupled ADP receptor (P2Y₁₂) of platelets was finally isolated by expression cloning (Hollopeter et al., 2001), and P2Y₁₃ and P2Y₁₄ receptors were characterized during a systematic study of orphan receptors (Chambers et al., 2000; Communi et al., 2001a). As of today, there are eight accepted human P2Y receptors: P2Y₁, P2Y₂, P2Y₄, P2Y₆, P2Y₁₁, P2Y₁₂, P2Y₁₃, and P2Y₁₄ (Abbracchio et al., 2003) (Table 1; Fig. 1) (see also section IV.). The missing numbers represent either nonmammalian orthologs or receptors having some sequence homology to P2Y receptors but for which there is no functional evidence of responsiveness to nucleotides. In particular p2y3 (Webb et al., 1996a) may be a chicken ortholog of P2Y₆ (Li et al., 1998), whereas p2y8 (Bogdanov et al., 1997) and tp2y (Boyer et al., 2000) could be the *Xenopus* and turkey orthologs of P2Y₄, respectively. p2y7 (Akbar et al., 1996) is a leukotriene B₄ receptor (Herold et al., 1997; Yokomizo et al., 1997); however, recently cross-reaction between agonists for some leukotriene receptors and some P2Y receptors has been found (see section IX.C.1), requiring further investigation. p2y5 (Webb et al., 1996b; Li et al., 1997) and p2y10 (Rao et al., 1999) must be considered as orphan receptors, although it has been reported (King and Townsend-Nicholson, 2000) that human p2y5 expressed in oocytes gives functional responses to ATP. p2y9 was reported to be a novel receptor for lysophosphatidic acid, distant from the Edg family (Noguchi et al., 2003). P2Y₁₅ was recently introduced to designate the orphan receptor GPR80/GPR99 on the basis that it would be a receptor for adenosine 5'-monophosphate (AMP²) (Inbe et al.,

² Abbreviations: AMP, adenosine 5'-monophosphate; TM, transmembrane domain; GPCR, G protein-coupled receptor; PTX, pertussis toxin; PLC, phospholipase C; PI3-K, phosphatidylinositol 3-kinase; IP₃, inositol triphosphate; UTP, uridine 5'-triphosphate; 2-MeSADP, 2-methylthio-ADP; MAP, mitogen-activated protein; ERK, extracellular signal-regulated protein kinase; PK, protein kinase; UDP, uridine 5'-diphosphate; CHO, Chinese hamster ovary; SCG, superior cervical

TABLE 1
Accepted human P2Y receptors

Receptor	Chromosome (Human)	Agonist (Human)	Phenotype of Knockout Mice
P2Y ₁	3q24–25	ADP	Inhibition of platelet aggregation Increased bleeding time Resistance to thromboembolism
P2Y ₂	11q13.5	ATP = UTP	Abolition of chloride secretory response to ATP/UTP in airways
P2Y ₄	Xq13	UTP	Abolition of chloride secretory response to ATP/UTP in jejunum and colon
P2Y ₆	11q13.5	UDP	No knockout mice are available
P2Y ₁₁	19p31	ATP	No murine P2Y ₁₁ gene
P2Y ₁₂	3q21–25	ADP	Inhibition of platelet aggregation Increased bleeding time Resistance to thromboembolism
P2Y ₁₃	3q24–25	ADP	Available, no phenotype yet
P2Y ₁₄	3q24–25	UDP-glucose	No knockout mice are available

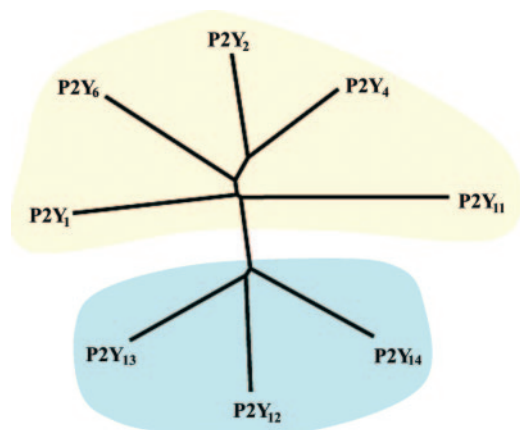


FIG. 1. A phylogenetic tree (dendrogram) showing the relationships among the current members of the P2Y receptor family. The P2Y receptors can be divided into two subgroups, shown with green and blue backgrounds. Sequences were aligned using CLUSTALX, and the tree was built using the TREEVIEW software. Reprinted from Abbracchio et al. (2003) with permission from Elsevier.

2004), but it is now firmly established that it is actually a receptor for α -ketoglutarate (He et al., 2004; Qi et al.,

ganglion; 2-MeSATP, 2-methylthio-ATP; PIP₂, phosphatidylinositol-4,5-bisphosphate; PI, phosphatidylinositol; GIRK, G protein-activated inward rectifier, Kir3; RGS, regulator of GPCR signaling; NHERF, Na⁺/H⁺ exchanger regulatory factor; CFTR, cystic fibrosis transmembrane conductance regulator; SAR, structure-activity relationship; ADP β S, adenosine 5'-O-(2-thiophosphate); MRS2255, 1,5-anhydro-2-(adenin-9-yl)-2,3-dideoxy-D-arabino-hexitol-6-triphosphate tetraammonium salt; β , γ -meATP, β , γ -methylene ATP; MRS2365, (1'S, 2'R, 3'S, 4'R, 5'S)-4-[(6-amino-2-methylthio-9H-purin-9-yl-1-diphosphoryloxymethyl)bicyclo[3.1.0]hexane-2,3-diol]; A3P5P, adenosine 3',5'-diphosphate; MRS2179, 2'-deoxy-N⁶-methyladenosine-3', 5'-bisphosphate; MRS2216, 2'-deoxy-2-chloro-N⁶-methyladenosine-3',5'-biphosphate; MRS2279, (N)-methanocarba-N⁶-methyl-2chloro-2'-deoxyadenosine-3', 5'-bisphosphate; MRS2500, 2-iodo-N⁶-methyl-(N)-methanocarba-20-deoxyadenosine-30,50-bisphosphate; MRS2584, (1'S, 4'R, 6'S, 7'S)-phosphoric acid mono-[6-(2-iodo-6-methylaminopurin-9-yl)-4-phosphonooxymethyl-2-oxabicyclo[2.2.1]hept-7-yl] ester; MRS-2298, 2-[2-(2-chloro-6-methylamino-purin-9-yl)methyl]propane-1,3-bisoxo(diammoniumphosphate); AR-C67085MX, 2-propylthio- β , γ -dichloromethylene-ATP; AR-C69931MX, N⁶-[2-(methylthioethyl)-2-(3,3,3-trifluoropropylthio)- β , γ -dichloromethyl ATP, cangrelor; AZD-6140, 3-[[[7-[2-3,4-difluoro-phenyl]-cyclopropylamino]-5-propylsulfanyl][1,2,3]triazolo[4,5-d]pyrimidine-3-yl]-5-(2-hydroxyethoxy)-cyclopentane-1,2-diol; INS49266, 6-phenylurea-2',3'-phenylacetaldehyde acetal ATP; INS50589, ((2S,3aR,4R,6R,6aR)-4-(6-(3-ethylureido)-9H-purin-9-yl)-tetrahydro-2-styrylfuro[3,4-d][1,3]dioxol-6-yl)methyl dihy-

2004; Suarez Gonzalez et al., 2004), as was also recently underlined in a report by the Subcommittee (Abbracchio et al., 2005).

B. Structural Aspects

In contrast with P2X receptors, P2Y receptor genes do not contain introns in the coding sequence, except for the P2Y₁₁ receptor. Site-directed mutagenesis of the P2Y₁ and P2Y₂ receptors has shown that some positively charged residues in transmembrane domains (TM) 3, 6, and 7 are crucial for receptor activation by nucleotides (Erb et al., 1995; Jiang et al., 1997b) (Fig. 2) (section V.).

drogen phosphate; MRS2395, 2,2-dimethyl-propionic acid 3-(2-chloro-6-methylaminopurin-9-yl)-2-(2,2-dimethyl-propionyloxymethyl)-propyl ester; PPADS, pyridoxalphosphate-6-azophenyl-2',4'-disulfonic acid; Ap₄A, diadenosine tetraphosphate; MRS2211, 3,5-diethyl-2-methyl-4-(trans-2-(4-nitrophenyl)vinyl-6-phenyl-1,4-dihydropyridine-3,5-dicarboxylate; ATP γ S, adenosine-5'-O-(3-thio)triphosphate; UDP β S, uridine β -thiodiphosphate; UTP γ S, uridine-5'-O-(2-thio)triphosphate; INS365, Up₄U, P¹,P⁴-di(uridine 5'-)tetraphosphate (tetrasodium salt); INS-37217, Up₄dC, P¹-(uridine 5'-)P⁴-(2'-deoxycytidine 5')tetraphosphate (tetrasodium salt); MRS2341, (1'S,2'R,3'S,4'R,5'S)-4-(2,4(H,3H)-dioxopyrimidin-1-yl)-1 (riphosphoryloxymethyl)bicyclo[3.1.0]-hexane-2,3-diol; AR-C126313, 5-(7-chloro-4H-1-thia-3-aza-benzof[4-yl]-3-methyl-6-thioxo-piperadin-2-one; AR-C118925, 5-((3,4-dihydro-2-oxo-5-2,8-dimethyl-dibenzo[2,3:6,7]cyclohept-4-enyl-4-thioxopyrimidin-1(2H)-yl)methyl)-N-(1H-tetrazol-5-yl)furan-2-carboxamide; MRS2578, 1,4-di-[[3-isothio-cyanato phenyl]-thioureido]butane; A3P5PS, adenosine 3'-phosphate 5'phosphosulfate; RT-PCR, reverse transcription-polymerase chain reaction; kb, kilobase(s); DAG, diacylglycerol; RGD, Arg-Gly-Asp; AA, arachidonic acid; NO, nitric oxide; IL, interleukin; MAPK, mitogen-activated protein kinase; JNK, c-Jun NH₂-terminal kinase; EGFR, epidermal growth factor receptor; SH3, Src homology 3; PDGF, platelet-derived growth factor; LPS, lipopolysaccharide; IBD, inflammatory bowel disease; Ap₅A, diadenosine pentaphosphate; Ap₆A, diadenosine hexaphosphate; h, human; CS-47, prasugrel (LY640315); Ap₃A, diadenosine triphosphate; GTP γ S, guanosine 5'-O-(3-thio)triphosphate; TX, thromboxane; VASP, vasodilator-stimulated phosphoprotein; MDCK, Madin-Darby kidney cell; ABC, ATP-binding cassette; MDR, multidrug resistance; VDAC-1, voltage-dependent anion channel-1; ACh, acetylcholine; NDPK, nucleoside diphosphokinase; E-NTPDase, ecto-nucleotide triphosphate diphosphohydrolase; E-NPP, ecto-nucleotide phosphophosphate/phosphodiesterase; NTP, nucleoside triphosphate; NDP, nucleoside diphosphate; FRET, fluorescence resonance energy transfer; CPA, cyclopentyl adenosine; CysLT, cysteinyl leukotriene receptor; I κ B, inhibitor of nuclear factor (NF)- κ B; PMA, phorbol 12-myristate 13-acetate; CREB, cAMP response element-binding trans-cription factor; NMJ, neuromuscular junction; AChR, acetylcholine receptor; AChE, acetylcholinesterase.

First subgroup												
hP2Y ₁	..	LQ	RFIFH	VN..	..	IPF	HVMK	TMN..	..	ATY	QVTR	GLA..
hP2Y ₂	..	LV	RFLFY	TN..	..	LPF	HVTR	TLY..	..	MAY	KVTR	PLA..
hP2Y ₄	..	FV	RFLFY	WN..	..	VVF	HITR	TIY..	..	VVY	KVTR	PLA..
hP2Y ₆	..	LV	RFLFY	AN..	..	LPF	HITK	TAY..	..	AAY	KGTR	PFA..
hP2Y ₁₁	..	LE	RFLFT	CN..	..	VPY	HIMR	VLN..	..	VGY	QVMR	GLM..

Second subgroup												
hP2Y ₁₂	..	VT	SVIFY	FT..	..	VVF	HFAR	IPY..	..	YVK	ESTL	WLT..
hP2Y ₁₃	..	FS	SVIFY	ET..	..	APF	HFAR	VPY..	..	IAK	ETTL	FLA..
hP2Y ₁₄	..	VS	AVLFY	VN..	..	VPY	HIAF	IPY..	..	YMK	EFTL	LLS..
GPR87	..	YT	SVLFY	AN..	..	LPY	HLCR	IPF..	..	YCK	EITL	FLS..
H963	..	VT	ACLTY	IN..	..	VPY	HIVR	IPY..	..	FAR	EATL	LLA..
GPR34	..	VV	CTLFY	MN..	..	VPY	HAFR	FIY..	..	KTN	ETML	VLS..

FIG. 2. Conserved residues among P2Y receptors are shown in a larger size font. The residues that have been mutated in the studies of Erb et al. (1995) and Jiang et al. (1997b) are underlined. Those residues that are crucial in the activation of those receptors are in bold. The corresponding sequences of orphan receptors structurally related to P2Y receptors of the second subgroup are also displayed.

They probably interact with the negative charges of the phosphate groups of nucleotides, since it is known that the receptor ligands are nucleotide species uncomplexed to magnesium or calcium. Actually, the eight P2Y receptors identified so far have a H-X-X-R/K motif in TM6. The P2Y₁, P2Y₂, P2Y₄, P2Y₆, and P2Y₁₁ receptors share a Y-Q/K-X-X-R motif in TM7, whereas another motif, K-E-X-X-L is found in P2Y₁₂, P2Y₁₃, and P2Y₁₄ receptors (Abbracchio et al., 2003) (see also Fig. 2 and sections IV. and V.). More recently, for P2Y₁₂, P2Y₁₃, and P2Y₁₄ receptors, one additional K residue in extracellular loop 2 has been suggested to be particularly important for nucleotide binding (Costanzi et al., 2004) (see also below; section V.).

C. Orphan Receptors Related to P2Y Receptors

The bioinformatic analysis of the human genome has revealed the existence of approximately 800 G protein-coupled receptors (GPCRs) (Fredriksson et al., 2003), of which 367 would be receptors for endogenous ligands, the remaining ones being olfactory and other chemosensory receptors (Vassilatis et al., 2003). Among the 367 "endoGPCRs," more than 150 remain orphans. Several of the currently available orphan GPCR sequences are structurally related to P2Y receptors (Lee et al., 2001; Wittenberger et al., 2001; Joost and Methner, 2002; Vassilatis et al., 2003): GPR87, H963, and GPR34. In particular, the sequence of those receptors contains the structural motifs in TM6 and TM7 described earlier (Fig. 2).

Human GPR87 has been shown to be highly expressed in placenta and thymus. The mouse ortholog has also been cloned and has been found to be expressed in brain and liver (Wittenberger et al., 2001).

Human H963 has been shown to be expressed in fibroblasts, human peripheral blood mononuclear cells, and T cells (S. Ferrario, D. Lecca, C. Volonté, and M. P. Abbracchio, unpublished data). Neither the mouse nor

the rat orthologs have been cloned; however, a recent study has suggested that GPR34 is not a nucleotide receptor (Sugo et al., 2006).

Human GPR34 is expressed in brain, heart, placenta, small intestine, pancreas, spleen, thymus, kidney, and skeletal muscle (Schöneberg et al., 1999). The mouse ortholog has also been described, and prominent expression has been found in liver and testis (Schöneberg et al., 1999).

Despite active research in several laboratories, the ligands of these receptors have not been identified. For example, despite the demonstration that GPR34 and ADP-like receptors (e.g., P2Y₁₂ and P2Y₁₃) have a common evolutionary origin (Schulz and Schöneberg, 2003), in the inositol phosphate assay, Cos-7 cells coexpressing GPR34 and Gα_{q14} did not show any response to ADP application (Schöneberg et al., 1999).

III. Second Messenger Systems and Ion Channels

A. Coupling to G Proteins and Intracellular Signaling Pathways

Coupling of the various P2Y receptors to specific G proteins was initially inferred from indirect evidence [measurement of intracellular levels of inositol phosphates, calcium, or cAMP and determination of pertussis toxin (PTX) sensitivity]. Direct evidence was recently obtained by measuring the effect of ADP on GTP hydrolysis in vesicles reconstituted with P2Y₁ and either Gα_{q11}β₁γ₂ or Gα₁₁β₁γ₂ (Waldo and Harden, 2004). Similar experiments demonstrated that P2Y₁₂ couples to Gα_{i2} more effectively than to Gα_{i1} and Gα_{i3} and not at all to Gα_o or Gα_q (Bodor et al., 2003). One given P2Y receptor can couple to functionally distinct G proteins. For instance, in HEL cells, activation of phospholipase C (PLC) by the P2Y₂ receptor is inhibited completely by a Gα₁₆ antisense oligonucleotide but also partially by PTX (Baltensperger and Porzig, 1997). Similarly, in gastric smooth muscle cells, it appears that P2Y₂ couples to PLC-β₁ via Gα_{q11} and to PLC-β₃ via Gα_{i3}β₁γ₂-derived βγ subunits (Murthy and Makhlof, 1998). The P2Y₂ receptor also has been shown to interact with α_v integrin to promote G_o-mediated chemotaxis in astrocytoma cells (Bagchi et al., 2005). The P2Y₁₂ receptor activates phosphatidylinositol 3-kinase (PI3-K) via Gα_i, but also RhoA and Rho kinase (Soulet et al., 2004). This action, which is insensitive to PTX, might be mediated by Gα_{12/13}, recently shown to play a critical role in platelet activation (Moers et al., 2003). Coupling of the same P2Y receptor to distinct G proteins and signaling pathways provides the possibility of agonist-specific signaling involving distinct active conformations of the receptor. For instance, activation of the P2Y₁₁ receptor by ATP leads to a rise in cAMP and in inositol trisphosphate (IP₃) and cytosolic calcium, whereas activation by uridine 5'-triphosphate (UTP) has been reported to produce calcium mobilization without IP₃ or cAMP increase (White et al., 2003). The P2Y₁₃ receptor can simultaneously

couple to G_{16} , G_i , and, at high concentrations of ADP, to G_s (like other G_i -coupled receptors such as the α_2 -adrenergic receptor): these three signaling pathways are characterized by different ratios of ADP to 2-methylthio-ADP (2-MeSADP) potency, suggesting the existence of ligand-specific conformations (Marteau et al., 2003). The activation of several P2Y receptors is commonly associated with the stimulation of several mitogen-activated protein (MAP) kinases, in particular extracellular signal-regulated protein kinase (ERK) 1/2. According to the cell context and the particular subtype, other classes of the MAP kinases, protein kinase (PK) C, calcium, and PI3-K are found to be involved to a variable extent (Soltoff et al., 1998; Huwiler et al., 2000; Communi et al., 2001a; Santiago-Pérez et al., 2001; Sellers et al., 2001).

B. P2Y Receptor Coupling to Ion Channels

1. Significance. In recent years, GPCRs in neurons and other excitable cells have been found to modulate the activity of voltage-gated ion channels in the cell membrane through certain actions of activated G proteins. Such actions are now well established in closing (or in certain cases in opening or potentiating) various classes of K^+ channels (Hille, 1994) and voltage-gated Ca^{2+} channels (Dolphin, 2003). Various voltage-gated Na^+ channels also have been observed to be modulated in certain cases via GPCR actions (Cantrell and Catterall, 2001; Maurice et al., 2001; Mantegazza et al., 2005), but this has not been reported for P2Y receptors. Although GPCR downstream signaling can lead to indirect effects on ion channels via activation of protein kinases, some GPCR regulation of several types of ion channel is more specific and more direct. This action has been investigated so far for only a very small fraction of the GPCR class. For P2Y receptors, specific couplings to certain K^+ and Ca^{2+} channels have been observed and analyzed. This coupling will be an important component of P2Y receptor signal transduction, but one that will usually be invisible in studies of second messenger or downstream pathways, since those channel interactions can occur in short timescales (down to ~ 100 ms) by a direct or quasi-direct pathway in the cell membrane.

Our consideration here of P2Y signaling through cell membrane channels is necessarily focused on cases in which the P2Y subtypes concerned have been identified. Thus, in a number of recent studies, the disturbing factors of enzymatic breakdown or interconversion of the nucleotides applied as agonists (Lazarowski et al., 2000) or the activation of adenosine receptors via such breakdown (Masino et al., 2002) have been experimentally minimized, allowing demonstration of ion channel responses upon activation of native P2Y receptors in brain neurons, with clear evidence for their identity (Wirkner et al., 2002; Khakh et al., 2003; Koizumi et al., 2003; Luthardt et al., 2003; Zhang et al., 2003b; Bowser and Khakh, 2004; Kawamura et al., 2004). That evidence shows that ATP (or UTP or their products ADP or UDP)

present at synapses, plus ATP diffusing from astrocytes, activates P2Y receptors on distinct subsets of brain neurons, regulating their activities by the coupling of those receptors to specific ion channels, as detailed below.

Although ion channel couplings of P2Y receptors are primarily of importance in neurons, they have in a few cases been detected also in various other tissues, e.g., in cardiac muscle cells (Vassort, 2001). As yet that category has been little explored.

2. Approaches to Analysis of the Channel Interactions of Molecularly Identified P2Y Receptors. Some studies of channel coupling by P2Y receptors have been made by heterologous expression in commonly transfected host cell lines such as CHO or HEK293, or in the *Xenopus* oocyte. However, usually both the P2Y receptor and the identified ion channel under study must be introduced into them, and the final relationship and protein environment of those components may be far from that in any native neuron, in which individual GPCR types can be located specifically with their effectors in microdomains (Delmas et al., 2004).

The problems there can be minimized if a suitable neuronal host cell can be found. A number of requirements for this exist (e.g., endogenous P2 receptors to be insignificant therein), and all of those conditions have been found to be met in the superior cervical ganglion (SCG) cell from the sympathetic nervous system of the young rat or mouse (Brown et al., 2000a). This cell type is well equipped with endogenous ion channels of the types found in neurons generally (Ikeda, 1996; Filippov et al., 1997). Its size readily allows nuclear injection of a receptor cDNA, a route that favors normal processing and trafficking of the protein. Transfection difficulties with neurons are avoided, and recordings of the channel couplings can be made in each receptor-expressing cell, as reviewed below. Because single cells are constantly perfused with medium and subsequently with the (purified) agonist and the assay period is extremely short, the method avoids potential problems well known in other P2 receptor activity studies, i.e., accumulation of nucleotides released from cell populations or their acute release by cell disturbance, as well as losses of the added agonist by metabolism.

3. Voltage-Activated Channels Regulated by P2Y Receptors. Among the channels with which the SCG cell membrane is well endowed are two types of voltage-gated channel that are important in receptor-based regulation of neuronal activity, the Ca^{2+} channel of the N-type and the M-current K^+ channel. The M-current K^+ channels are heteromers of subunits of the Kv7 family and are critical for setting the responsiveness of neurons to synaptic inputs (Selyanko et al., 2001, and references therein). Closing of the M-current K^+ and/or N-type Ca^{2+} channels by action of certain P2Y receptors has been shown to occur.

a. Ca^{2+} channels. For $G_{i/o}$ protein-coupled receptors in general, inhibition of the N-type Ca^{2+} current has

been shown to occur through $G\beta\gamma$ subunits, acting directly on the channel (reviewed by Dolphin, 2003). This $\beta\gamma$ action holds for the G_i -linked $P2Y_{12}$ receptor in the SCG system, as shown by the demonstration that closure of the N-type Ca^{2+} channel via the $P2Y_{12}$ receptor is fully sensitive to PTX and is totally abolished when $G\alpha$ -transducin, a $G\beta\gamma$ -scavenging protein, is coexpressed (Simon et al., 2002). For the $G_{i/o}$ -linked $P2Y_{13}$ receptor, inhibition of voltage-gated Ca^{2+} channels would again be expected. Indeed, evidence has been obtained in HEK293 cells (into which the N-type Ca^{2+} channel had been introduced by transfection) to suggest that a native $P2Y_{13}$ receptor is there and acts thus (Wirkner et al., 2004).

The action at N-type Ca^{2+} channels of activated $P2Y_1$ and $P2Y_2$ receptors (Table 1) is very similar to that of the endogenous M1 receptor ($G_{q/11}$ -linked) in the same cells, with all three receptors showing a PTX-sensitive and a PTX-insensitive component. α_q is presumably required for the latter component, as was established using knockout mice for the M1 inhibition of this channel (Haley et al., 2000), although those $P2Y$ receptors could conceivably also use α_{11} in some other cells. The PTX-sensitive component requires (at least for M1; Haley et al., 2000) the G_o protein. With $P2Y_1$, in contrast, $G_{i/o}$ does not act here (Table 2). However, with $P2Y_{1,2,6}$ receptors, both the PTX-insensitive and the PTX-sensitive N-type Ca^{2+} channel responses are abolished when $\beta\gamma$ subunits are sequestered by $G\alpha$ -transducin (Simon et al., 2002; Filippov et al., 2004). Hence, the α and the $\beta\gamma$ components of selected trimeric G protein(s) must operate together in this type of $P2Y$ receptor action, as summarized in Table 2.

Higher potency of the agonists for a given subtype than in second messenger assay systems is generally observed in the channel transduction, which follows a direct route within the cell membrane; e.g., for 2-MeSADP at the $P2Y_1$ receptor the EC_{50} value for Ca^{2+} channel closure is 0.57 ± 0.05 nM (Filippov et al., 2000).

(All experimental values quoted in section III. are at $20-22^\circ\text{C}$). Adenosine triphosphates (e.g., 2-MeSATP and ATP) when in pure form also are agonists in this $P2Y_1$ reaction, in conditions in which their agonistic diphosphates are excluded throughout; this has been controversial for $P2Y_1$ receptors in some other expression systems, but the potency and efficacy of the triphosphates vary strongly with the density of this receptor subtype (see discussion and references in Filippov et al., 2000). The N-type Ca^{2+} channel closure via the $P2Y_{12}$ receptor behaves likewise, with 2-MeSATP an agonist at $P2Y_{12}$ receptors, too, with exceptional potency (Simon et al., 2002) (see also section VI.F. for further discussion). This behavior, compared with the variation of behavior seen with these agonists at $P2Y_{12}$ receptors in different native situations (Barnard and Simon, 2001), again suggests a particularly strong dependence of the observed efficacy of these subtypes on receptor density in the membrane and on the directness of the transduction pathway involved.

Interestingly, the agonist selectivity of a $P2Y$ receptor can also be changed in the channel interactions from that observed for it in transductions downstream. Thus, whereas the transfected $P2Y_6$ receptor was reported to be UDP-selective and to have negligible action by UTP in its IP_3 formation (Nicholas et al., 1996), both those nucleotides are strong agonists in the closure of the N-type Ca^{2+} channel and likewise for the M-current K^+ channel response (Filippov et al., 1999). With impurity and metabolism of the UTP being excluded, the K_A of rat $P2Y_6$ for UTP in its channel activity is 20.1 ± 1.4 nM. A precedent for this phenomenon lies in the transduction dependence of relative agonist potencies described above for the $P2Y_{13}$ receptor (section III.A.); it is interesting for the $P2Y_6$ receptor because it indicates alternative binding site conformations for two native agonists, UTP and UDP.

b. The M-current K^+ channel. The M-current can be inhibited through the activation of a number of GPCRs

TABLE 2
Ion channel interactions of $P2Y$ receptors

Data derived from Filippov et al. (1998, 1999, 2000, 2003, 2004); Brown et al. (2000a); and Simon et al. (2002).

P2Y	Ca ²⁺ Channel Closure			G Protein ^c	K(M) Channel ^a		GIRK ^a	
	Whole Cell	Perf. ^a	PTX Block ^b		Closure	G Protein ^d	Activation	G Protein
			%					
1	Yes ^e		~50	$\alpha_{q/11} + \beta\gamma (G_q + G_o)$	Yes ^e	$\alpha_{q/11}$	Yes ^{e,f}	$\beta\gamma$
2	Yes		~60	$\alpha_{q/11} + \beta\gamma (G_q + G_o)$	Yes	$\alpha_{q/11}$	Yes ^f	$\beta\gamma$
4	No	Weak	~80 ^a	$\beta\gamma (G_o)^a$	Yes	$\alpha_{q/11}$	No	
6	Yes	Yes	~0 ^a	$\alpha_{q/11} + \beta\gamma (G_q)^a$	Yes	$\alpha_{q/11}$	No	
12	Yes		100	$\beta\gamma (G_{i/o})$	No		Yes ^f	$\beta\gamma$

K(M) channel, M-current K^+ channel; GIRK, G protein-activated inwardly rectifying K^+ channels.

^a Determined in perforated patch recording (Perf.), which avoids possible dialysis of some soluble cell components.

^b The percentage of the N-type Ca^{2+} current inhibition by $P2Y$ action, which is blocked by PTX pre-treatment.

^c The G protein subunits which are proposed to act at N-type Ca^{2+} channels; in parentheses are the parent heterotrimeric G proteins deduced to provide the $\beta\gamma$ subunits involved, this being noted only for the perforated patch state in which that is used; involvement of $\beta\gamma$ (where tested) was stated by showing total prevention of channel closure by coexpressing excess $G\alpha$ transducin.

^d The G protein subunit found to act at M-current K^+ channels.

^e 'Yes' denotes that the induced change occurs with agonist potencies similar to or greater than those known for other transductions of this receptor. 'No' denotes that the induced charge is essentially absent; 'Weak' denotes that the induced charge occurs but at greatly reduced agonist potency.

^f Highly sensitive to PTX pretreatment.

of the $G_{q/11}$ -linked class (Brown and Yu, 2000) but not of other classes. The G protein subunit involved in GPCR-mediated closing of this channel by M1 muscarinic receptors in rat or mouse SCG neurons is G_{α_q} , as shown both by anti- G_{α_q} antibody depletion and by G_{α_q} -gene knockout (Haley et al., 2000). For P2Y receptors, this parallel, together with the complete inability of the $G_{i/o}$ -linked P2Y₁₂ receptor to affect the M-current plus the PTX-insensitive closure of that channel by all four of the $G_{q/11}$ -linked subtypes examined (Table 2), indicates that this action can be ascribed to a G_{α_q} pathway also. Whether in other neurons $G_{\alpha_{11}}$ may also act thus is not yet excluded.

The pathway from G_{α_q} -GTP to the M-type channel Kv7 proteins has been deduced to be, in general, via PLC-mediated depletion of phosphatidylinositol-4,5-bisphosphate (PIP₂), removing its stabilizing interaction with those proteins and hence allowing channel closure (Ford et al., 2003; Zhang et al., 2003a; Suh et al., 2004; Winks et al., 2005). PIP₂ thus acts as a second messenger, but one that is diffusible only within the membrane and acting negatively. Restoration of the membrane PIP₂ occurs by a PI4-kinase and PI5-kinase pathway (Winks et al., 2005). The same PIP₂/PI4-kinase mechanism has been supported for the closure of the N-type Ca²⁺ channel by similar observations on G_q -linked muscarinic M1 receptor action and its recovery (Gamper et al., 2004). We presume that a similar mechanism is operative for P2Y receptors of the $G_{q/11}$ -linked class, but this has not as yet been specifically examined.

4. Activation or Inactivation of G Protein-Gated K⁺ Channels by P2Y Receptors. An entirely different class of K⁺ channels is specialized for interaction with GPCRs, i.e., the G protein-activated inward rectifiers (Kir3 or GIRK channels). Activations of $G_{i/o}$ -linked receptors, but not of those linked to $G_{q/11}$, generally open these channels (Fernandez-Fernandez et al., 2001; Stanfield et al., 2002). In line with this, in the rat SCG neurons 2-MeSADP, acting at an introduced rat P2Y₁₂ receptor, strongly activates a GIRK channel (EC₅₀: 0.099 ± 0.008 nM) (Simon et al., 2002).

Very exceptionally, some P2Y receptors, which usually link to $G_{q/11}$, have also been found to efficiently activate GIRK channels. This case occurs with P2Y₁ receptors (Simon et al., 2002), in which 2-MeSADP, for example, acts to open with high potency a GIRK channel in the SCG cell. Another instance has been seen in *Xenopus* oocytes, with the P2Y₂ receptor (activated by ATP or by UTP) opening a GIRK channel, when both that and the receptor are implanted therein (Mosbacher et al., 1998; Mark et al., 2000). In these cases the normally $G_{\alpha_{q/11}}$ -linked P2Y₁ or P2Y₂ receptors can display in neurons a channel-based transduction that is highly PTX-sensitive and dependent on G_{α_i} . This action is fast and direct, through $\beta\gamma$ subunits released in the cell membrane from $G_{\alpha_i}\beta\gamma$ trimers and binding at the K⁺ channel (Simon et al., 2002; Filippov et al., 2004), i.e., as

found with P2Y₁₂. This mechanism of the P2Y₁ receptor signal transduction via a K⁺ channel could be an indicator of an interaction of that receptor in neurons with a recruited GPCR-regulating component [e.g., a regulator of GPCR signaling (RGS) protein], since reconstituted P2Y₁ protein gives no coupling to G_{α_i} or G_{α_o} (Waldo and Harden, 2004) (see section VI.A.).

A second, independent, type of interaction with GIRK channels can occur with P2Y receptors, namely the closing of an open GIRK channel. This slow inactivation phase is common in $G_{q/11}$ -linked GPCRs, occurring in situ at GIRK channels already opened via endogenous $G_{i/o}$ -linked receptors. It occurs with P2Y₁ but not with P2Y₁₂ receptors (Simon et al., 2002). This inactivation is found also with P2Y₄ and P2Y₆ in the neuron (Filippov et al., 2004) and with P2Y₂ in oocyte expression (Mark et al., 2000). It is unaffected by PTX, with $G_{\alpha_{q/11}}$ being involved, as shown in the P2Y₁ case with evidence that included blockade by sequestering the released $\alpha_{q/11}$ subunits using the coexpressed RGS2 protein, which is specific for that action (Filippov et al., 2004). Again, when investigated for P2Y₁ receptors, depletion of PIP₂ from the membrane and the liberation of IP₃ in the cytosol can be seen to be correlated with the course of the GIRK current inactivation, when visualized by a sensor for them, PLC δ -PH, fluorescently tagged (Filippov et al., 2004). There appears to be a role for PIP₂ in the GIRK inactivation by P2Y receptors, related to that proposed for it (see above) in the M-current K⁺ channel inactivation, but this requires further specification.

5. Conclusions on the Interactions with Identified Ion Channels. Five P2Y subtypes have been compared so far (Table 2). Clear differences are seen between the P2Y subtypes, with only P2Y₁ and P2Y₂ receptors showing a common behavior in the three transductions. These considerable variations support the conclusion that the channel couplings seen are not a general phenomenon produced by overloading with an exogenous receptor. Likewise, for all of these P2Y receptors, their maximum inhibition of the N-type Ca²⁺ current is well below 100% of the total N-type Ca²⁺ current recorded and is less than that attainable by test activations of the native α_2 -adrenergic or M1 muscarinic receptors in the same cell (see references cited in Table 2). There is also evidence for the coupling of some native P2Y receptors to such ion channels in brain neurons (section III.B.1.) and also in autonomic neurons and related cells (Ennion et al., 2004; Lechner et al., 2004, and references cited therein).

Some promiscuity between PTX-sensitive and PTX-insensitive G proteins is described here for these transductions (Table 2). Such cross-reaction is already known for P2Y₂ and P2Y₄ receptors in other signaling pathways (see sections III.A. VI.B, and VI.C.). We noted its occurrence here, however, also with the P2Y₁ receptor, in its coupling to a Ca²⁺ channel and to some but not all of its couplings to K⁺ channels. Indeed, the P2Y₁-linked acti-

vation of the GIRK K^+ channel described is almost entirely $G_{i/o}$ -mediated, but none of that component is seen in its PLC-dependent transductions; this finding underlines the independence of the channel-coupling pathways in P2Y receptor signaling. The coupling to that channel is not, however, necessarily linked to a relaxation of selectivity for $\alpha_{q/11}$ -containing heterotrimers, since the P2Y₆ receptor, in contrast with the others, strictly maintains that selectivity in all of the couplings (Table 2) as well as in other transductions. Again, in the independent inactivation reaction of the GIRK channel all four of these P2Y receptors generally associated with $G_{q/11}$ linkage were seen here to fully maintain that selectivity.

Such ion channel responses are usually recorded in whole-cell patch-clamping, but this may permit diffusible intracellular cofactors to dialyze out. Most of the analyses of P2Y receptor coupling in the SCG cell discussed above were made instead in the more-laborious perforated-patch mode (using amphotericin B to form small membrane pores), to avoid that possibility. In the case of the P2Y₆-mediated Ca^{2+} channel closure (Filipov et al., 1999), use of this configuration abolished a partial block by PTX (although for other subtypes it can remain). Hence a soluble component, possibly an RGS protein, may cooperate in the membrane-delimited reactions of the P2Y₆ receptor to direct its G protein coupling.

The P2Y₄ receptor has not so far been shown to occur in neurons, unlike P2Y₁ and P2Y₆ receptors; its mRNA is prominent in the brain ventricular system, cardiac and skeletal muscles, some smooth muscles, and some other peripheral sites but is undetectable in neurons (Webb et al., 1998). This may be why its coupling to the N-type Ca^{2+} channel of a neuron can be weak and readily lost (Table 2). Adaptation of such a P2Y subtype may evolve for a different signaling environment than that in neurons.

C. Other Potential Interactions with Ion Channels

Little is known of these as yet. One clue to some other interactions of the P2Y₁ receptor comes from the recent finding (Fam et al., 2005) that it can bind strongly to the Na^+/H^+ exchanger regulatory factor 2 (NHERF-2) through the extreme C-terminal motif DTSL, which is specific to P2Y₁ in this family. For comparison, binding of the related NHERF-1 to P2Y₁ receptors (Hall et al., 1998) is negligible (Fam et al., 2005). The three membrane-located NHERF subtypes either activate or inhibit various Na^+/H^+ exchangers, but these actions are indirect since it is now known that NHERFs are actually scaffolding proteins, which can localize various exchangers in membrane microdomains with selected receptors and signaling intermediates, e.g., G_{α_q} , Src, and certain isoenzymes of PLC and of PKC (Donowitz et al., 2005). For example, the NHERF-2 scaffold can link a cGMP-dependent protein kinase or a protein kinase A anchor-

ing protein and hence protein kinase A, to modulate a tethered Na^+/H^+ exchanger by specific phosphorylations thereon (Cha et al., 2005; Donowitz et al., 2005).

When the endogenous P2Y₁ receptor (as studied in C6 glioma cells) is linked through its tail to the NHERF-2 scaffold, the Ca^{2+} transients produced by its activation by 2-MeSADP become strongly prolonged (Fam et al., 2005). This will change the P2Y₁ selectivity for the various calcium-sensitive signaling cascades and for ion channel interactions. Another interaction of the P2Y₁ receptor is with the chloride channel of the cystic fibrosis transmembrane conductance regulator (CFTR); in renal epithelial cells, 2-MeSADP activation of native P2Y₁ receptors stimulates the chloride channel activity of the CFTR. This is again an indirect action arising from the NHERF-2 colocalization of this P2Y subtype and the CFTR; expression of a dominant negative NHERF-2 mutant blocks the CFTR regulation through P2Y₁, as does a blocker of the protein kinase A anchoring protein binding of PKA (Guerra et al., 2004). The evidence suggested that P2Y₁ receptor-mediated PKC activation leads to potentiation of PKA and its action on the associated CFTR channels.

Additionally, a highly unusual mode of GPCR interaction with an ion channel has been suggested for several P2Y receptors by Lee et al. (2003b). A novel, unidentified, voltage-gated channel of the *Xenopus* oocyte, T_{in} , was reported to be activated and modulated after expression of human P2Y_{1,2,6,11}, but not by P2Y₄ nor by other $G_{q/11}$ -linked GPCRs. It was deduced that this does involve a direct receptor binding to the channel. However, expression of G_{α} also activates this channel and the mechanism and physiological significance are at present unclear.

IV. Principles of P2Y Receptor Classification

As already outlined above, eight distinct mammalian P2Y receptors have been cloned and recognized so far: the P2Y_{1,2,4,6,11,12,13} and the recently reclassified P2Y₁₄ receptors (Abbracchio et al., 2003). The missing numbers in the P2Y₁₋₁₄ sequence represent GPCRs cloned from nonmammalian vertebrates or receptors for which a functional response to nucleotides has not yet been convincingly demonstrated.

Pharmacologically (Table 1) P2Y receptors can be broadly subdivided into 1) adenine nucleotide-preferring receptors mainly responding to ADP and ATP. This group includes human and rodent P2Y₁, P2Y₁₂, and P2Y₁₃, and human P2Y₁₁ (which has, however, been recently reported to also respond to UTP) (White et al., 2003); 2) uracil nucleotide-preferring receptors. This group includes human P2Y₄ and P2Y₆ responding to either UTP or UDP; 3) receptors of mixed selectivity (human and rodent P2Y₂, rodent P2Y₄ and, possibly, P2Y₁₁); and 4) receptors responding solely to the sugar nucleotides UDP-glucose and UDP-galactose (P2Y₁₄).

From a phylogenetic and structural (i.e., protein sequence) point of view, two distinct P2Y receptor subgroups characterized by a relatively high level of sequence divergence have been identified (Jacobson et al., 2002; Abbracchio et al., 2003). The first subgroup includes P2Y_{1,2,4,6,11} subtypes and the second subgroup encompasses the P2Y_{12,13,14} subtypes (see dendrogram in Fig. 1 reproduced from Abbracchio et al., 2003). Alignment of the deduced amino acid sequences of the cloned P2Y receptors has shown that the human members of this family are 21 to 48% identical (Table 3). The highest degree of sequence identity is found among the second subgroup of P2Y_{12,13,14}. Interestingly, despite clear phylogenetic relationships with the first subgroup, the P2Y₁₁ subtype seems to differ from all the others, for both sequence and pharmacological differences between species (e.g., canine versus human) and also based on its absence in the murine and rat genomes (Table 1). Thus, it might be hypothesized that this receptor differentiated from P2Y₁ and subsequently underwent many modifications and insertions that led to a dissimilar receptor, despite the common origin. Cotranscription and intergenic splicing of the P2Y₁₁ gene might be another evolutionary event accounting for its dissimilarity from the other P2Y receptors.

The two P2Y receptor subgroups highlighted above also differ in several other features. In particular, specific amino acid motifs in TM6 and TM7 have been previously proposed to be important for binding to extracellular nucleotides (Erb et al., 1995; Jiang et al., 1997b; Boarder and Webb, 2001; Jacobson et al., 2002). All human P2Y receptors share the typical TM6 H-X-X-R/K motif that might be important for agonist activity (Erb et al., 1995; Jiang et al., 1997b; Boarder and Webb, 2001; Jacobson et al., 2002) (Table 3; see also section V.). A Q/K-X-X-R defining motif in TM7 has also been proposed to participate in ligand binding for the P2Y_{1,2,4,6} and P2Y₁₁ receptors. In P2Y_{12,13,14} receptors, this motif

is substituted with K-E-X-X-L, which might affect ligand binding characteristics (Table 3). In humans, the genes of P2Y_{12,13,14} receptors cluster in the same region of chromosome 3, together with the gene encoding for the P2Y₁ receptor; in this region, three additional still unidentified "orphan" GPCRs structurally related to the known P2Y receptors can also be found (Table 1). The mapping of the known genes of the P2Y receptors and the structurally related orphans in the human genome is detailed in Simon and Barnard (2003). Interestingly, two of these orphan receptors (i.e., GPR87 and H963) also show full conservation of the defining motifs in TM6 and TM7 typically found in P2Y_{12,13,14}. Their functional characterization may eventually lead to inclusion in this P2Y receptor subgroup.

Finally, these two P2Y receptor subgroups also differ in their primary coupling to transductional G proteins. In particular, receptors in the first subgroup (i.e., P2Y_{1,2,4,6,11}) all principally use G_q/G₁₁ to activate the PLC β /IP₃ pathway and release intracellular calcium, whereas receptors in the second subgroup (i.e., P2Y_{12,13,14}) almost exclusively couple to members of the G_{i/o} family of G proteins (Table 3; see also section III. and individual receptor subsections and individual receptor summary tables that appear at the end of the article). Secondary couplings have been also reported, especially for receptors of the first subgroup in heterologous expression systems (Simon et al., 2002; Burnstock, 2003; King and Townsend-Nicholson, 2003; Köttgen et al., 2003; White et al., 2003). For receptors of the second group, P2Y₁₃ has been also reported to couple to G α_{16} and stimulate PLC in recombinant systems overexpressing this G protein (Communi et al., 2001a; Marteau et al., 2003), whereas activation of the native P2Y₁₄ receptor in astrocytes and microglia has been shown to increase intracellular calcium levels (Fumagalli et al., 2003; Bianco et al., 2005). Such "promiscuity" of G protein-coupling may depend on the indirect activation of

TABLE 3
Classification of P2Y receptors into two subsets

For P2Y_{1,2,4,6}, and ₁₁ receptors, amino acid residues reported in bold in TM6 and TM7 have been shown to be important for ligand binding in mutagenesis studies (Erb et al., 1995; Jiang et al., 1997b; Boarder and Webb, 2001; Jacobson et al., 2002; see also section V.). For P2Y_{12,13}, and ₁₄ receptors, direct evidence for the functional importance of the reported motifs in TM6 and TM7 is currently lacking. However, in a patient with a congenital bleeding disorder, a R-Q substitution in TM6 of the P2Y₁₂ receptor results in highly decreased receptor function (Cattaneo et al., 2003).

Receptor	Percentage of Identity								Proposed Amino Acid Motifs for Receptor Function		IUPHAR Receptor Code	Primary G Protein Coupling/ Second Messenger ^a
	P2Y ₁	P2Y ₂	P2Y ₄	P2Y ₆	P2Y ₁₁	P2Y ₁₂	P2Y ₁₃	P2Y ₁₄	TM6	TM7		
P2Y ₁	/	38	44	46	32	24	24	27	HXXK	QXXR	2.1.NUCT.01.000.00.00	G _{q/11} ; PLC β activation
P2Y ₂		/	41	41	29	25	26	26	HXXR	KXXR	2.1.NUCT.02.000.00.00	G _{q/11} ; PLC β activation
P2Y ₄			/	43	32	25	26	28	HXXR	KXXR	2.1.NUCT.03.000.00.00	G _{q/11} ; PLC β activation
P2Y ₆				/	34	24	24	23	HXXK	KXXR	2.1.NUCT.04.000.00.00	G _{q/11} ; PLC β activation
P2Y ₁₁					/	22	21	23	HXXR	QXXR	2.1.NUCT.05.000.00.00	G _{q/11} ; PLC β activation
P2Y ₁₂						/	48	47	HXXR	KEXXL	2.1.NUCT.06.000.00.00	G _{i/o} ; inhibition of adenylate cyclase
P2Y ₁₃							/	47	HXXR	KEXXL	2.1.NUCT.07.000.00.00	G _{i/o} ; inhibition of adenylate cyclase
P2Y ₁₄							/		HXXR	KEXXL	2.1.NUCT.08.000.00.00	G _{i/o} ; inhibition of adenylate cyclase

^a The main transductional mechanisms are reported; see text for more details.

additional G protein subtypes within protein complexes containing the P2Y receptor.

Thus, a division into two subgroups could be considered, based on 1) phylogenetic (i.e., sequence) similarity (Fig. 1; Table 3), the 2) presence of amino acid defining motifs proposed to be important for ligand binding (Table 3; see also section V.), and 3) selectivity of primary G protein coupling (Table 3). However, we prefer to wait to formally implement this subdivision until there is a more complete knowledge of these receptors, with some of the orphan P2Y-like receptors still waiting for deorphanization, the possibility of new receptors still to be discovered, and the place of the P2Y₁₁ receptor still to be clearly resolved.

V. Agonists and Antagonists

A. Chemical Structure of Agonist and Antagonist Ligands

Most of the P2Y receptor subtypes are still lacking potent and selective synthetic agonists and antagonists. However, considerable progress in exploring structure-activity relationships (SARs) has been made for P2Y₁ and P2Y₁₂ receptors and to a lesser extent for the P2Y₂ receptor. Radioligand binding studies have been successfully carried out at the P2Y₁ and P2Y₁₂ receptors, but so far not at any other P2Y receptor subtype. Here we describe the current state of molecular probes known

for the P2Y receptors, categorized by the chemical class of the endogenous agonists.

1. ADP-Preferring P2Y Receptors: P2Y₁, P2Y₁₂, and P2Y₁₃. ADP (**1** in Fig. 3) is the endogenous agonist at the P2Y₁, P2Y₁₂, and P2Y₁₃ receptors, and it interacts at these subtypes with generally greater affinity than does ATP (**3**) (Palmer et al., 1998; Boeynaems et al., 2003; Marteau et al., 2003). At P2Y₁ receptors, derivatives of ADP tend to be full agonists (the EC₅₀ of ADP is ~100 nM), whereas ATP appears to be a partial agonist. At P2Y₁₂ receptors, ADP derivatives activate (the EC₅₀ of ADP is ~100 nM) and 5'-triphosphate derivatives antagonize (Gachet and Hechler, 2005). At P2Y₁₃ receptors, both ADP and ATP might act as full agonists, with EC₅₀ values of ~100 nM. However, under some conditions, ATP can behave as a weak partial agonist, suggesting that, as described for the P2Y₁ receptor, the activity of ATP in recombinant systems may vary according to the level of expression of the P2Y₁₃ receptor (see also below).

Phosphate modifications among P2Y receptor ligands often serve to increase their stability toward ecto-nucleotidases. For example, the added stability of a terminal thiophosphate group resulted in its incorporation in some useful P2Y nucleotide agonists. One such analog, ADPβS (**2**), is a potent agonist of both P2Y₁ (EC₅₀ = 96 nM), P2Y₁₂ (EC₅₀ = 82 nM) (Jacobson et al., 2002), and

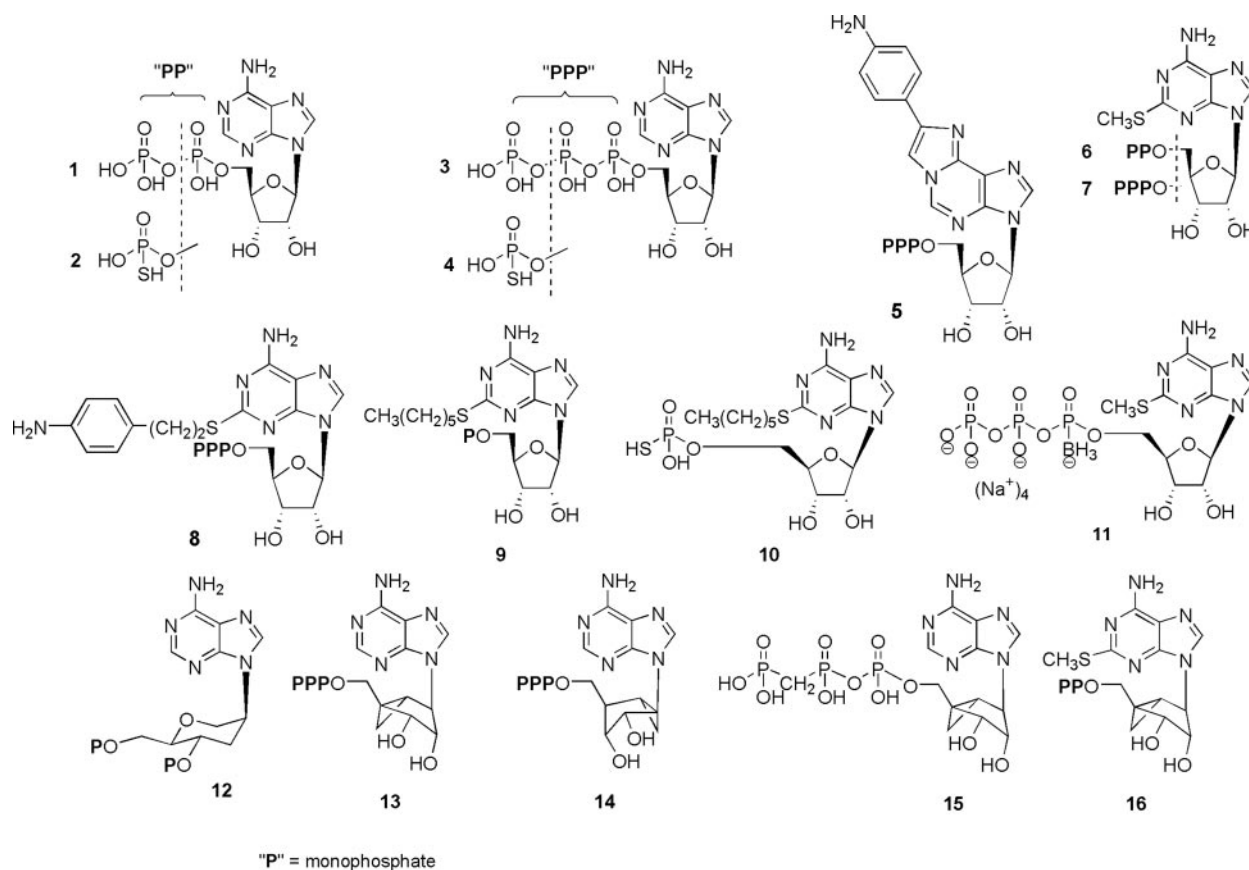


FIG. 3. Structures of adenine-derived nucleotide agonists of P2Y receptors.

P2Y₁₃ receptors (EC₅₀ = 42 nM) (Communi et al., 2001a). Although terminal thiophosphates are enzymatically more stable than the oxygen equivalents, they are subject to chemical oxidation reactions; thus, solutions of these compounds are prone to instability.

Figure 3 shows the structures of adenine-derived nucleotide agonists of P2Y receptors. The structure of the nucleobase of adenine nucleotides has been extensively probed for effects at P2Y receptors. 8-Aza and 1-deaza modifications are generally tolerated. A fluorescent adenine-modified derivative of ATP (**5**) behaved as a potent P2Y₁ receptor agonist (Sharon et al., 2004). The 2-position of the adenine ring can accommodate a wide variety of substituents, with resultant activation of both P2Y₁ and P2Y₁₂ receptors. Long-chain and sterically bulky groups may be accommodated at the 2-position. In particular, 2-alkylthio ethers (Fischer et al., 1993; Brown et al., 2000b) appear to provide high potency at these subtypes when bonded to a variety of alkyl or alkylaryl groups. Notably, the smallest member of this class is 2-MeSADP [**6**, which is a potent agonist (EC₅₀) at P2Y₁ (6 nM), P2Y₁₂ (1 nM), and P2Y₁₃ (1 nM) receptors (Jacobson et al., 2002; Marteau et al., 2003); however, see also above and individual receptor subsection] and is highly selective in comparison with other P2Y receptor subtypes. For example, at the P2Y₂ receptor compound **6** is inactive at 100 μM. The corresponding triphosphate, 2-MeSATP (**7**), is less potent and selective as a P2Y receptor agonist, since it also activates P2X receptors (King, 1998). The sterically bulky *p*-aminophenylethylthio analog (2-[2-(4-aminophenyl)ethylthio]adenosine 5'-triphosphate) (**8**), potently activated the P2Y₁ receptor (EC₅₀ = 1 nM) (Fischer et al., 1993). The SAR of alkynyl substitutions at the 2-position of P2Y₁ receptor agonists has been explored (Cristalli et al., 2005).

Although AMP is inactive at the P2Y₁ receptor, addition of a 2-thioether substituent as a receptor "anchor" causes AMP analogs to activate P2Y₁ receptors. Among these derivatives, 2-hexylthioadenosine 5' monophosphate (**9**) was especially potent, with an EC₅₀ of 59 nM at the turkey P2Y₁ receptor (Boyer et al., 1996b). Certain 2-thioether derivatives of AMP derivatives also activate the P2Y₁₂ receptor in C6 glioma cells in the micromolar range. The α-thio modification of AMP analogs (e.g., **10**) increases potency at the P2Y₁ receptor (Fischer et al., 1999). Such monophosphate derivatives have also been reported to inhibit ecto-nucleotidases, which complicates their use as P2Y receptor agonists.

A BH₂ moiety may serve as a substitute for an ionized oxygen atom of the α-phosphate of ATP derivatives in promoting binding to the P2Y₁ receptor binding site. Thus, 5'-(1-boranotriphosphate) derivatives such as the 2-methylthio derivative (**11**) have been found to potently activate the P2Y₁ receptor (Nahum et al., 2002). Because the 1-boranotriphosphate moiety is chiral, it was possible to separate two stable isomers in this series. The

more potent isomer of **11** displayed an EC₅₀ of 2.6 nM at the rat P2Y₁ receptor.

The ribose moiety of nucleotide derivatives was also modified, resulting in enhanced potency at the P2Y₁ and P2Y₁₂ receptors. Simple carbocyclic (cyclopentyl) analogs of ATP were found to enhance antagonist affinity at the P2Y₁₂ receptor (see below) (van Giezen and Humphries, 2005). Similarly, at the P2Y₁ receptor, carbocyclic and even acyclic substitutions of ribose were studied. In general, carbocyclics and ring-constrained nucleotide analogs were able to maintain agonism at the P2Y₁ receptor, whereas acyclic derivatives proved to be exclusively antagonists (see below). A ring-expanded, yet nonglycosidic, dehydroanhydrohexitol analog MRS2255 (**12**) activated the P2Y₁ receptor with an EC₅₀ of 3.0 μM (Nandan et al., 2000).

Among the more successful examples of the use of carbocyclic or sterically constrained carbocyclic substitution of the ribose moiety for P2Y receptor interactions are the "methanocarba" analogs (Nandan et al., 2000; Kim et al., 2002). These analogs incorporate a conformationally fixed bicyclic ring system, consisting of fused cyclopentane and cyclopropane rings, in place of the ribose moiety. Depending on the position of fusion, the resulting nucleotides may adopt one of two conformations: (*N*), north, or (*S*), south. Correlation of ring geometry with the biological activities helped define the conformational requirements of the ribose moiety in P2Y receptor binding and led to pharmacological probes of unusual selectivity and affinity. For example, the two isomeric methanocarba equivalents of ATP indicated a strong preference (ratio of potency >100-fold) at the P2Y₁ receptor for the (*N*)-isomer **13** over the (*S*)-isomer **14** (Kim et al., 2002).

Combination of this ring system with other favorable modifications of ADP or ATP resulted in large qualitative differences from the native nucleotides in receptor activation. For example, whereas β,γ-methylene ATP (β,γ-meATP) is a weak partial agonist at the human P2Y₁ receptor, the corresponding (*N*)-methanocarba-β,γ-meATP (**15**) was a full agonist with an EC₅₀ of 158 nM (Ravi et al., 2002). MRS2365 (**16**), the most potent known agonist of the human P2Y₁ receptor, with an EC₅₀ of 0.4 nM, induces platelet shape change without aggregation (Chhatriwala et al., 2004). In addition, the high selectivity of **16** for the P2Y₁ receptor in comparison to its inactivity at P2Y₁₂ and P2Y₁₃ receptors was striking, in contrast to the relatively nonselective 2-MeSADP (**2**). Thus, the P2Y₁₂ and P2Y₁₃ receptors have very different conformational preferences within the ribose-binding region than does the P2Y₁ receptor. At P2Y₁₃ receptors, under optimal experimental conditions, ATP (**2**) and 2-MeSATP (**5**) are equipotent as agonists.

Figure 4 shows the structures of nucleotide-based antagonists of P2Y receptors. A successful approach to the development of potent and selective P2Y₁ receptor antagonists was made possible by the observation by Boyer

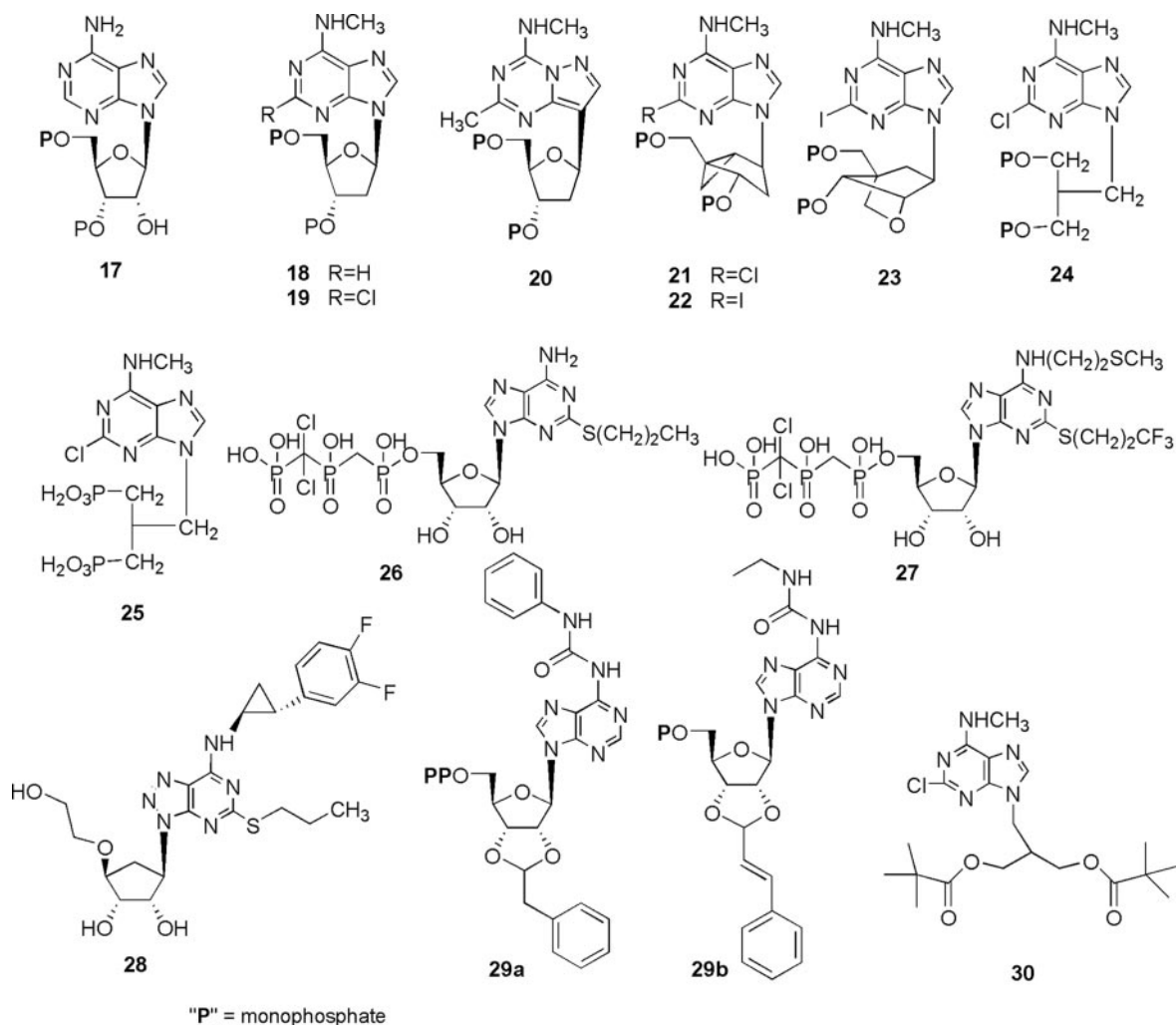


FIG. 4. Structures of nucleotide-based antagonists of P2Y receptors.

et al. (1996a) that naturally occurring adenosine bisphosphate derivatives such as A3P5P (**17**) act as partial agonists or antagonists of the receptor. Thus, the splitting and repositioning of the phosphate groups of the 5'-diphosphate group of ADP to separate ribose positions (5'- and either 3'- or 2'-) reduces efficacy at the P2Y₁ receptor. Removal of the 2'-hydroxyl group and addition of the potency-enhancing N⁶-methyl group resulted in MRS2179 (**18**), and the corresponding 2-chloro analog MRS2216 (**19**), which became full antagonists at the P2Y₁ receptor with IC₅₀ values of 300 and 100 nM, respectively (Nandanan et al., 1999; Brown et al., 2000b). The SAR of alkyl, thioether, and other substitutions at the 2-position of bisphosphate antagonists has been explored (Nandanan et al., 1999; Raboission et al., 2002a). Raboission et al. (2002b) synthesized a C-nucleotide bisphosphate derivative (**20**) that antagonized P2Y₁ receptors. In addition, the adenine N9 nitrogen is not essential in P2Y₁ receptor interaction, and similarly the N1 nitrogen was found to be unnecessary through the evaluation of 1-deaza analogs (Nandanan et al., 1999).

Upon introduction of the conformationally preferred (*N*)-methanocarbo ring system into this series of bisphosphate nucleotide antagonists, the P2Y₁ receptor affinity was further enhanced. Thus, MRS2279 (**21**), the (*N*)-methanocarbo equivalent of the riboside (**19**), and the corresponding 2-iodo derivative MRS2500 (**22**) demonstrated high affinity in competitive antagonism at the human, turkey, rat, and mouse P2Y₁ receptors (Nandanan et al., 2000; Boyer et al., 2002; Waldo et al., 2002; Cattaneo et al., 2004). The K_i value of MRS2500 was 0.78 nM, as determined in inhibition binding experiments at the human P2Y₁ receptor (Ohno et al., 2004), and the compound was highly specific for this subtype. MRS2279 or related antagonists were demonstrated to be inactive at P2Y_{2,4,6,11,12,13} and P2X_{2,3,4,7} receptors (Boyer et al., 2002; Marteau et al., 2003). Weak antagonism by MRS2279 of the rat P2X₁ receptor expressed in *Xenopus* oocytes was observed (Brown et al., 2000b). However, the potency of many of the known P2 receptor antagonists is magnified in this assay; thus, MRS2279 may still be considered highly selective for the P2Y₁ receptor. In platelet studies, for example, antagonism of

the P2X₁ receptor by this compound and related P2Y₁ receptor antagonists is not observed (Baurand et al., 2001). The K_i value of MRS2500 at the P2Y₁ receptors was found to be 0.79 nM, and it was consistently potent in inhibiting the ADP-induced aggregation of human and rat platelets. [³³P]MRS2179, [³H]MRS2279, and [³²P]MRS2500 were prepared and shown to be effective as radioligand probes in platelets and in other tissue (Baurand et al., 2001; Waldo et al., 2002; Houston et al., 2005). A novel ring system was incorporated in a carbocyclic locked nucleic acid derivative MRS2584 (**23**), which acted as a P2Y₁ receptor antagonist with a binding K_i of 23 nM (Ohno et al., 2004).

In addition to the approach of rigidifying the ribose moiety in a conformation that approximates the conformation preferred in receptor binding, the opposite approach, which used a flexible acyclic ribose equivalent, also produced bisphosphate antagonists of the P2Y₁ receptor of intermediate affinity. The acyclic nucleotide analog MRS2298 (**24**) represents a bisphosphate structure that was optimized for affinity at the P2Y₁ receptor, with an IC₅₀ in inhibition of PLC of 200 nM (Cattaneo et al., 2004). A related analog containing the metabolically stable phosphonate linkage, MRS2496 (**25**), displayed an IC₅₀ at the rat platelet P2Y₁ receptor of 0.68 μM (Xu et al., 2002a).

Extensive structure-activity studies of ATP derivatives as antagonists of the platelet P2Y₁₂ receptor resulted in high-affinity, selective antagonists of interest as antithrombotic agents. Several such 5'-triphosphate derivatives, including AR-C67085MX (**26**) and AR-C69931MX (**27**) (Ingall et al., 1999), were used in clinical trials, with the recognition that triphosphate derivatives would not be suitable for oral administration. In this series, it was also possible to substitute the unwieldy triphosphate group with uncharged moieties such as short alcohols or esters, thus proving that a highly anionic moiety is not needed for recognition by the P2Y₁₂ receptor. This discovery led to compounds such as the carbocyclic nucleoside derivative AZD6140 (**28**), which is an orally active P2Y₁₂ receptor antagonist of nanomolar affinity that inhibits platelet aggregation up to 8 h after administration (Springthorpe, 2003). The presence of the 3,4-difluorophenyl group limits the metabolism of compound **28**.

Analogs of ADP having neutral, hydrophobic substitutions at the ribose 2'- and 3'-hydroxyl groups and at the adenine NH₂ position were found to antagonize the P2Y₁₂ receptor. One such analog is INS49266 (**29a**), which displayed a K_B of 361 nM in the inhibition of ADP-induced platelet aggregation (Douglass et al., 2002). The agonist potencies of compound **29a** at P2Y₁ and P2Y₂ receptors were >10 and 14 μM, respectively. A related analog in this series, INS50589 (**29b**), has entered clinical trials as a platelet aggregation inhibitor with a rapid onset and offset mechanism of action that is intended for intravenous administration.

The acyclic template used in MRS2298 (**24**) has been adapted to antagonists of the P2Y₁₂ receptor (Xu et al., 2002a). Upon replacement of the two phosphate groups with hydrophobic esters, such as in the dipivaloate MRS2395 (**30**), the selectivity shifted entirely from the P2Y₁ receptor to the P2Y₁₂ receptor. MRS2395 displayed an IC₅₀ of 3.6 μM in the inhibition of ADP-induced aggregation of rat platelets by antagonizing the P2Y₁₂ receptor.

The action of the successful antithrombotic drug clopidogrel (**31** in Fig. 5) is also dependent on the P2Y₁₂ receptor present on platelets. Clopidogrel (**31**) produces a metabolite (**32**), which acts as an irreversible P2Y₁₂ receptor antagonist (Savi et al., 2000). CS-747 (**33**) (also known as prasugrel or LY640315) is acting with a similar mechanism (Sugidachi et al., 2001; Niitsu et al., 2005). CT50547 (**34**) is a P2Y₁₂ antagonist characterized by a non-nucleotide (and not highly charged) structure (Scarborough et al., 2001). Recently, a novel P2Y₁₂ receptor antagonist structure in the form of a pyrazolidine-3,5-dione derivative (**35**) was presented (Fretz et al., 2005). The search for more drug-like (nonphosphate-containing or uncharged), and structurally novel antagonists of the P2Y₁ receptor has so far had limited success. A dipeptide conjugate of adenosine, which bore carboxylate groups, was found to antagonize hP2Y₁ receptor responses with a K_B value of 4.0 μM (Sak et al., 2000). Metabolites of the hypolipidemic drug nafenopin may act as P2Y₁ receptor antagonists, including a coenzyme A conjugate that displayed a K_B value of 58 nM. A cyclic depsipeptide YM-254890 (**36**), a fermentation product of a *Chromobacterium* isolated from soil, inhibited the ADP-induced aggregation of platelets with an IC₅₀ of 31 nM and was inactive at P2Y₁₂ receptors (Uemura et al., 2006). The mechanism was found to be inhibition of Gα_q rather than direct interaction with P2Y₁ receptors.

Figure 5 shows the structures of non-nucleotide antagonists of P2Y receptors. Known non-nucleotide antagonists of P2 receptors were also modified in an effort to achieve P2Y₁ receptor selectivity. Many of the classic antagonists of P2 receptors are highly negatively charged polycyclic compounds and tend to block certain P2Y as well as P2X responses. Although no antagonists are totally nonselective with respect to the entire P2 superfamily (such an antagonist would be a useful tool), these polyanions are the least selective and often have activities beyond the P2 receptors. For example, the nonselective P2X/P2Y antagonist Reactive blue 2 (**37**) and its derivatives (such as Acid Blue 80, Acid Blue 129, and Acid Violet 34) have been shown to block action at P2Y₁ receptors; however, high potency and selectivity have not been achieved (Brown and Brown, 2002; Jacobson et al., 2002). The polysulfonate suramin (**38**) and its derivatives, in addition to displaying trypanocidal drug properties, are relatively nonselective P2 antagonists with, in general, reversibility upon washout (King,

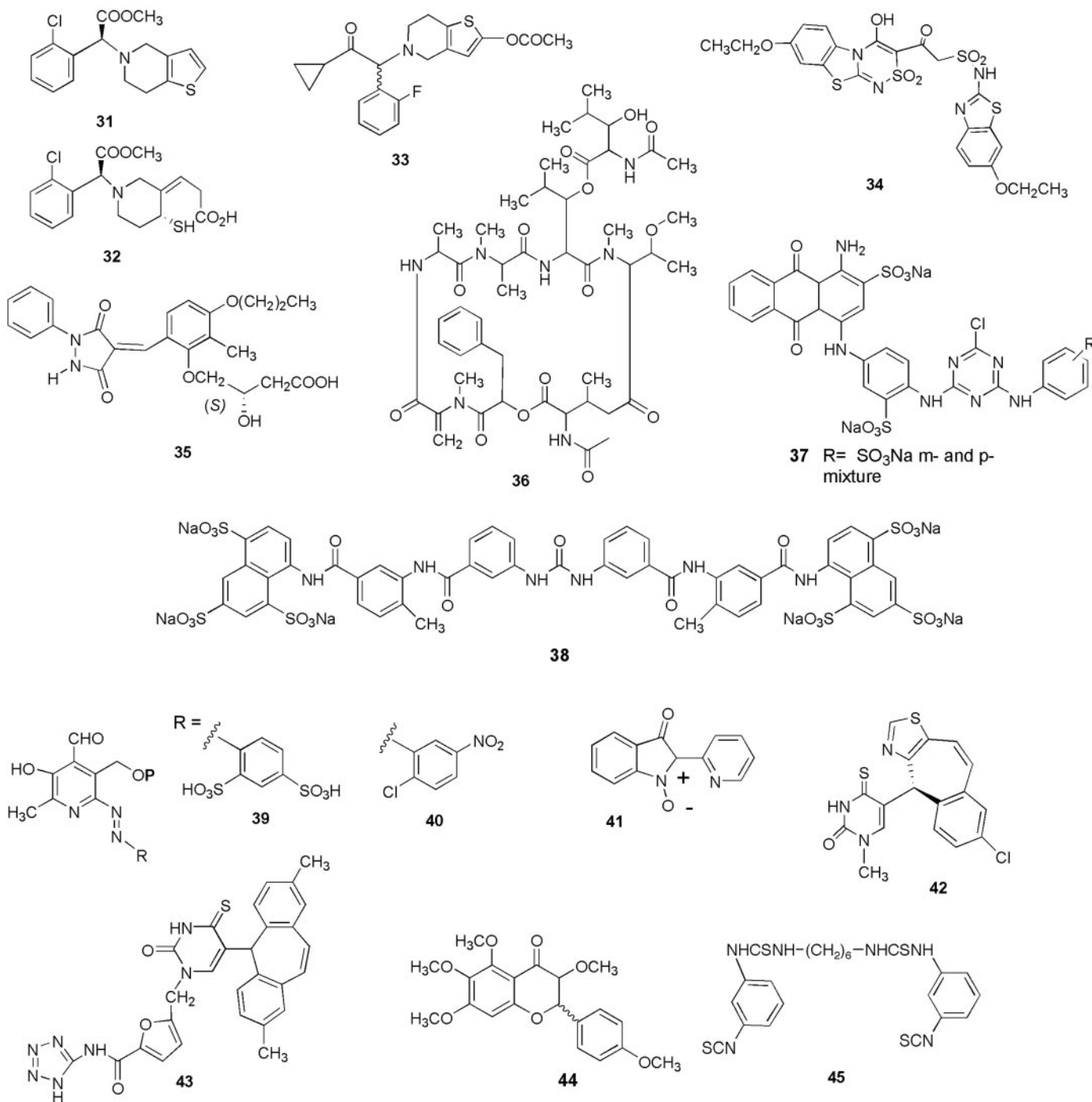


FIG. 5. Structures of non-nucleotide antagonists of P2Y receptors.

1998). Within the P2Y family, suramin has been characterized as an antagonist of P2Y₂ receptors (Wildman et al., 2003) and P2Y₁₁ (Communi et al., 1999b) receptors. Derivatives of the pyridoxal phosphate derivative PPADS (**39**) have been shown to antagonize P2Y₁ receptor effects in a competitive fashion, although at μM concentrations (Lambrecht et al., 2002). Extensive SAR manipulations within these families have not resulted in antagonists of nanomolar affinity.

A variety of structurally diverse, non-nucleotide antagonists of the P2Y₁ receptor have been demonstrated

to be noncompetitive inhibitors. For example, pyridyl isatogen tosylate (**41**), which has been explored as a possible allosteric modulator of the receptor (King et al., 1996), was found to be a P2Y₁-selective antagonist in recombinant P2Y receptor systems (Gao et al., 2004). It reduced the maximal effect of 2-MeSADP in stimulation of PLC, with an IC₅₀ of 0.14 μM , but had no effect on the binding of [³H]MRS2279.

Two nucleotide 5'-triphosphate derivatives (**26** and **27**) were shown to potently antagonize the P2Y₁₃ receptor (Kim et al., 2005), but in a noncompetitive manner.

The following compounds were found to antagonize action at the P2Y₁₃ receptor (IC₅₀): suramin (**38**) (2.3 μM), PPADS (**39**) (11.7 μM), Ap₄A (0.216 μM), and AR-C69931MX (**27**) (0.004 μM) (Marteau et al., 2003). Recently, a derivative of PPADS, MRS2211 (**40**), was shown to selectively antagonize the human P2Y₁₃ receptor (Kim et al., 2005). The antagonism of MRS 2211 of agonist-induced PLC was competitive with a pA₂ value of 6.3.

2. ATP-Preferring P2Y Receptor: P2Y₁₁. At P2Y₁₁ receptors, ATP is the preferred native ligand (Communi et al., 1999b), and ATP_γS (**4**) is a more potent agonist than ATP. Selective antagonists of the P2Y₁₁ receptors are under development (Ullmann et al., 2005). The P2Y₁₂ antagonist AR-C67085MX (**26**) acts as a potent agonist at the P2Y₁₁ receptor (Communi et al., 1999b).

3. UTP-Recognizing P2Y Receptors: P2Y₂ and P2Y₄.

The P2Y₂ receptor is activated nearly equipotently by UTP (**47** in Fig. 6) and ATP (**3**) but is not activated by the corresponding 5'-diphosphates, i.e., UDP (**46**) and ADP (**1**). The P2Y₄ receptor is primarily activated by uracil nucleotides, depending on species. In the rat, ATP is also a P2Y₄ agonist, but in humans it acts as a P2Y₄ antagonist. Uridine β-thiodiphosphate (UDPβS, **48**) and the γ-thiophosphate (UTP_γS, **49**) are selective agonists for P2Y₆ and P2Y₂/P2Y₄ receptors, respectively (Malmsjö et al., 2000). Numerous substitutions of the uracil ring of UTP have been reported to reduce potency at the P2Y₂

receptor (Müller, 2002). The 5-bromo derivative of UDP (**50**) retains potency at the P2Y₆ receptor.

The adenine dinucleotide Ap₄A is a potent agonist at the rat P2Y₄ receptor and is less potent than ATP at the P2Y₂ receptor. Other uracil dinucleotides, such as INS365 (Up₄U, **51**), also potently activate the P2Y₂ receptor (Shaver et al., 2005). The dependence of potency at various P2Y receptors on the number of bridging phosphate units in the dinucleotide series indicates an optimum at the tetraphosphate. Newer-generation P2Y₂ receptor agonists, such as INS37217 (Up₄dC, **52**), have been reported (Pendergast et al., 2001; Yerxa et al., 2002). INS37217 is less prone to enzymatic hydrolysis than more common dinucleotide agonists. P2Y₂ receptor agonists are of clinical interest for the treatment of pulmonary and ophthalmic diseases and possibly cancer.

Ribose substitution with the (*N*)-methanocarba ring system has been shown to preserve the potency of both adenine and uracil nucleotides at the P2Y₂ receptor and UTP (e.g., MRS2341, **53**) at the P2Y₄ receptor (Kim et al., 2002). However, inclusion of the same (*N*)-methanocarba ring system in the corresponding 5'-diphosphate prevented activation of the P2Y₆ receptor. Therefore, enzymatic cleavage of compound **53** to the diphosphate does not have complicating actions at the P2Y₆ subtype.

Figure 6 shows the structures of uracil-derived nucleotide agonists of P2Y receptors. Suramin (**38**) is a weak

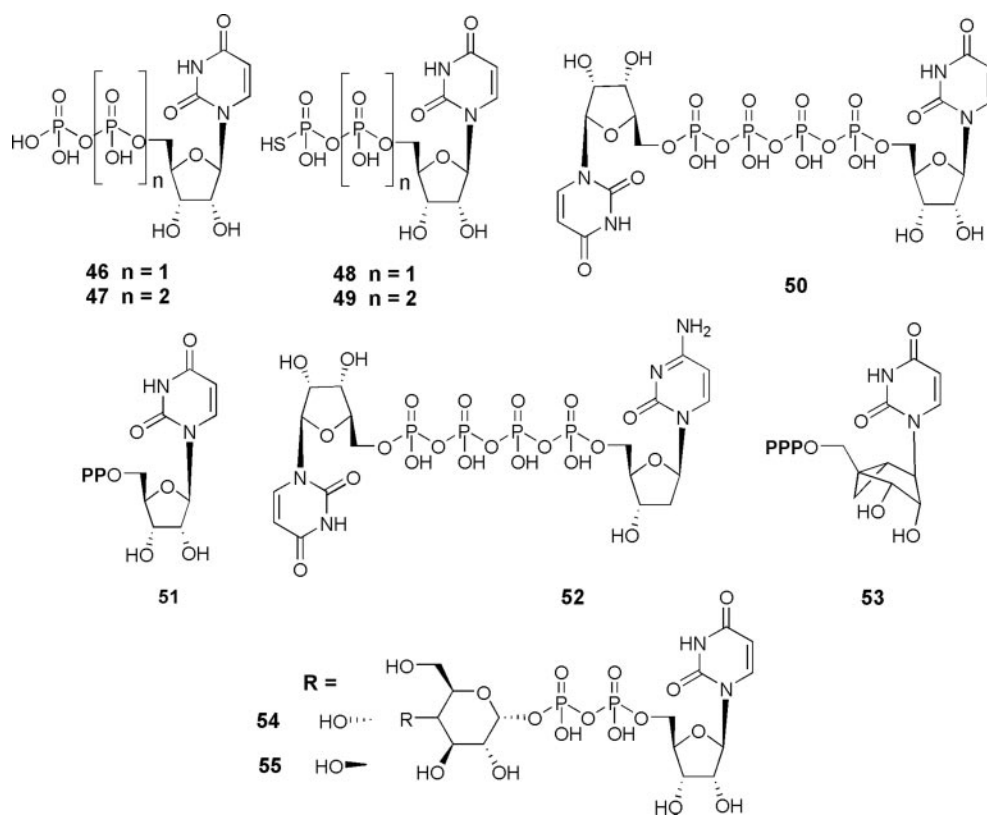


FIG. 6. Structures of uracil-derived nucleotide agonists of P2Y receptors.

antagonist at the P2Y₂ receptor with an IC₅₀ of 48 μM (Müller, 2002). A family of selective, heterocyclic antagonists of the P2Y₂ receptor containing a thiouracil moiety, including AR-C126313 (**42**) and the related aminotetrazole derivative AR-C118925 (**43**), has been reported (Meghani, 2002). Reactive Blue 2 (**37**) at a concentration of 100 μM effectively blocks rat P2Y₄ receptors but only partially blocks human P2Y₄ receptors. ATP antagonizes the human but not rat P2Y₄ receptor (Kennedy et al., 2000).

Flavonoids have been identified as a new lead for the design of P2Y₂ receptor antagonists (Kaulich et al., 2003). Tangeretin (**44**) is a potent, noncompetitive antagonist with an IC₅₀ of 12 μM.

4. UDP-Preferring P2Y Receptor: P2Y₆. UDP derivatives activate the P2Y₆ receptor more potently than the corresponding 5'-triphosphates (Malmjö et al., 2000; Müller, 2002); thus, UDP is a selective agonist at this subtype. The β-thiodiphosphate (**48**) was shown to be more potent than UDP in activation of the P2Y₆ receptor and more stable to degradation. INS48823 is a potent P2Y₆ agonist (Korcok et al., 2005).

Various diisothiocyanate derivatives were found to be potent insurmountable (and possibly irreversible by virtue of the reactive isothiocyanate groups) antagonists of human P2Y₆ as well as of other P2Y receptors (Mamedova et al., 2004). A 1,4-di-(phenylthioureido)butane derivative (MRS2578, **45**) selectively inhibited UDP-induced PLC activity through both human (IC₅₀ = 37 nM) and rat (IC₅₀ = 98 nM) P2Y₆ receptors expressed in 1321N1 human astrocytes and was inactive at human P2Y₁, P2Y₂, P2Y₄, and P2Y₁₁ receptors. Limitations of using these isothiocyanate derivatives as P2Y antagonists include the pharmacological irreversibility, relative instability of the compounds in aqueous medium, and hydrophobicity and consequent low aqueous solubility.

5. UDP-Sugar-Preferring P2Y Receptor: P2Y₁₄. The most recently cloned receptor, P2Y₁₄, responds to UDP-glucose (**54**) and has a sequence more similar to the P2Y₁₂ and P2Y₁₃ receptors than to the other P2Y subtypes (Abbracchio et al., 2003). The P2Y₁₄ receptor is also activated by UDP-galactose (**55**). It is the only known P2Y subtype to be activated by nucleotide sugars. The SAR at this subtype has not yet been explored. Antagonists of the P2Y₁₄ receptor are still unknown.

B. Molecular Modeling Studies

The two distinct subgroups of P2Y receptors were successfully modeled by homology modeling, with the high-resolution structure of bovine rhodopsin serving as a template (Moro and Jacobson, 2002). The putative TM binding site and other regions of the human P2Y₁ receptor have been extensively studied by means of mutagenesis (Fig. 2). To ascertain which residues of the P2Y₁ receptor are involved in ligand recognition, individual

residues of the TMs and extracellular loops were mutated to Ala and other amino acids (see also Table 4).

Recent computational models of all of the P2Y receptors were derived from a multiple-sequence alignment based on a combined manual and automatic approach, which takes into account not only the primary structure of the proteins but also the three-dimensional information deducible from the secondary and tertiary structures of the template (Costanzi et al., 2004). The receptors display the general motif of a single-polypeptide chain forming seven helical TMs, which are connected by three extracellular and three intracellular loops. The ends of the chain form an extracellular amino-terminal region and a cytoplasmic carboxyl-terminal region, as shown for the P2Y₁ and P2Y₁₂ receptors (Fig. 7). According to the models, at the cytoplasmic end of TM7 both the receptors fold at an angle of ~90° to form a helical segment that is homologous to H8 in rhodopsin and runs parallel to the plane of the cell membrane (Hoffmann et al., 1999).

Figure 7 shows the theoretical structures of the putative nucleotide binding sites of P2Y₁ (A) and P2Y₁₂ (B) receptors, based on mutagenesis and molecular modeling experiments as described in Costanzi et al. (2004). The large figures show the binding sites as viewed from the plane of the plasma membrane with docked nucleo-

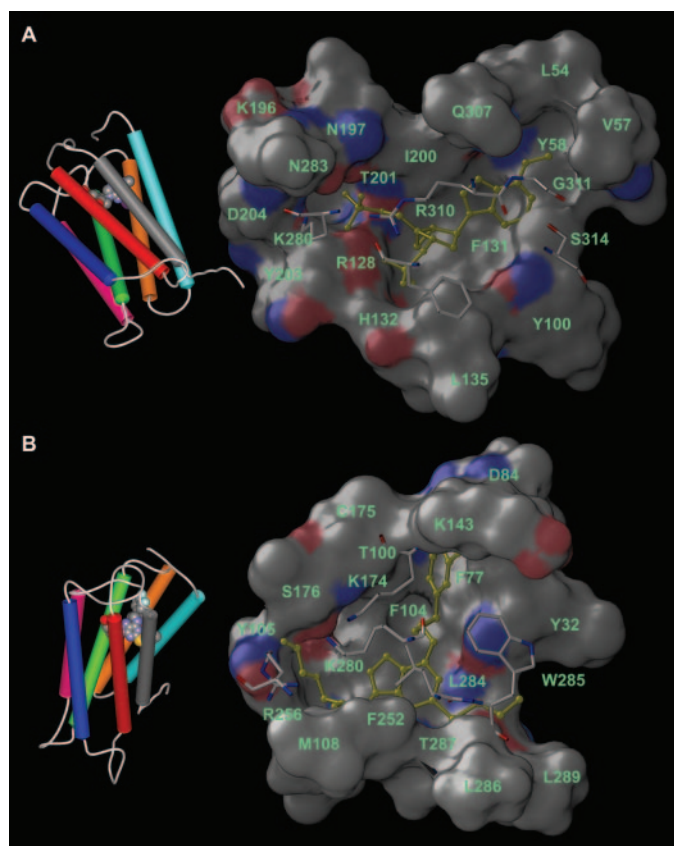


FIG. 7. Theoretical structures of the putative nucleotide binding sites of P2Y₁ (A) and P2Y₁₂ (B) receptors based on mutagenesis and molecular modeling experiments as described in Costanzi et al. (2004).

tide ligands (the antagonists MRS2500 for P2Y₁ and AZD6140 for P2Y₁₂). Key residues found to interact with the ligand in the human P2Y₁ and P2Y₁₂ receptors are indicated. To the left of each detailed structure is a smaller three-dimensional representation of the receptor including seven TMs (color coded: cyan, TM1; orange, TM2; green, TM3; magenta, TM4; blue, TM5; red, TM6; gray, TM7) and the connecting loops. The orientation of the entire receptor relative to the membrane is the same as for each detailed binding site model.

Table 4 summarizes the effects and structural role of specific amino acid residues of the human P2Y₁ receptor, deduced from site-directed mutagenesis, molecular modeling, and homology to other GPCRs. Ligand docking modeling was performed on the P2Y₁ and P2Y₁₂ receptor models. The results suggested that ADP binds to the P2Y₁ and P2Y₁₂ receptors on the exofacial side of the cavity delimited by TM1, TM2, TM3, TM6, and TM7 and capped with extracellular loop 2 (Fig. 7). Two different sets of three basic amino acids for each of the two subgroups are involved in coordination of the phosphate moiety of agonists (Costanzi et al., 2004). Molecular recognition in the P2Y₁ receptor of non-nucleotide antagonists, such as derivatives of PPADS (compound **39**), was also studied.

A cluster of positively charged amino acid side chains in TMs 3, 6, and 7 was proposed to form the counterions to the negatively charged 5'-di- or triphosphate moiety at the P2Y₁ receptor. Site-directed mutagenesis validated this prediction and further indicated several uncharged hydrophilic residues that may coordinate the nucleobase. Thus, the agonist 2-MeSADP (compound **6**) was inactive at R128(3.29)A and R310(7.39)A and S314(7.43)A mutant P2Y₁ receptors and had a markedly reduced potency at K280(6.55)A and Q307(7.36)A mutant P2Y₁ receptors.

In the P2Y₁₂ subgroup, the role of R3.29 in TM3 seemed to be fulfilled by a Lys residue in extracellular loop 2, whereas the residue R7.39 in TM7 seemed to be substituted by K7.35, located within the same TM but at a distance of four residues, i.e., one helical turn in the exofacial direction. Only R6.55 was common to the essential cationic residues of the two subclasses of P2Y receptors (Costanzi et al., 2004).

VI. P2Y Receptor Subtypes

The history of each currently recognized P2Y receptor subtype and available information on molecular structure, coupling to G proteins, transductional mechanisms, response to agonist/antagonist ligands, tissue distribution and function are reported in the following (additional information, including the official International Union of Pharmacology alphanumeric code, previous names, gene bank accession numbers, chromosomal localization, availability of radioligands, and

knockout or knockin animals, are also schematically reported in the individual tables for each subtype).

A. P2Y₁

Human (Ayyanathan et al., 1996; Janssens et al., 1996; Léon et al., 1996; Schacter et al., 1996), rat (Tokuyama et al., 1995), mouse (Tokuyama et al., 1995), cow (Henderson et al., 1995), chick (Webb et al., 1993), turkey (Filtz et al., 1994), and *Xenopus* (Cheng et al., 2003) P2Y₁ receptors have been cloned and characterized. In most species, ADP is a more potent agonist than ATP and their 2-methylthio derivatives are more potent than the parent compounds. UTP, UDP, CTP, and GTP are inactive (Waldo et al., 2002; Waldo and Harden, 2004). At present, the most potent agonist known is the *N*-methanocarba analog of 2-MeSADP, MRS2365 (Chhatriwala et al., 2004). ATP is, in fact, a partial agonist at the P2Y₁ receptor (Palmer et al., 1998) and so at low levels of receptor expression will act as an antagonist (Léon et al., 1997; Hechler et al., 1998b). Extracellular acidification and alkalinization do not appear to modify the activity of ATP (King et al., 1996). The first antagonists to display selectivity for the P2Y₁ receptor were A3P5P and A3P5PS (Boyer et al., 1996a), but these have been superseded by the highly selective and more potent MRS2179 (Boyer et al., 1998; Camaioni et al., 1998) and MRS2279 (Boyer et al., 2002). More recent studies showed that modification of both MRS2179 (Mathieu et al., 2004) and MRS2279 (Kim et al., 2003a; Cattaneo et al., 2004) at the 2-position of the adenine moiety further increases antagonist potency. The increased potency and selectivity of MRS2279 has been used in binding studies showing that [³H]MRS2279 bound specifically to the human P2Y₁ receptor, with a *K_d* of 3.8 nM. The binding was displaced by 2-MeSADP > ADP = 2-MeSATP > ATP and by MRS2279 = MRS2179 > A3P5P (Waldo et al., 2002; Waldo and Harden, 2004).

Site-directed mutagenesis studies on the human P2Y₁ receptor have produced a binding pocket model in which amino acid residues in TM3, 6 and 7 are critical determinants in the binding of ATP and other nucleotide derivatives (Jiang et al., 1997b; Moro et al., 1998). Arginine 128 (TM3) and lysine 280 (TM6) are proposed to interact with the α and β phosphate groups of ATP, arginine 310 (TM7) also with the β phosphate, and threonine 222 (TM5) with the γ phosphate. Additionally, glutamine 307 (TM7) and serine 314 (TM7) may interact with the adenine ring. Alanine scanning mutagenesis studies revealed that four cysteine residues in the extracellular loops, which are conserved in P2Y receptors, are essential for proper trafficking of the human P2Y₁ receptor to the cell surface (Hoffmann et al., 1999). These studies also showed that several charged residues in the extracellular loops are essential for P2Y₁ activation.

TABLE 4

Analysis of the pharmacological effects and structural role of individual residues of the human P2Y₁ receptor, deduced from site-directed mutagenesis, molecular modeling, and homology to other GPCRs

Where applicable, the data indicate the effects of single amino acid replacement on the activation of PLC by mutant human P2Y₁ receptors transiently expressed in COS-7 cells and results of molecular modeling of the docked complex of 2-MeSADP or ATP within the wild-type hP2Y₁ receptor (see Moro and Jacobson, 2002; Costanzi et al., 2004; and references therein).

Site	Mutation (if Performed)	Position ^a	Putative Interactions, Pharmacological Role	Potency Loss (Ratio for 2-MeSADP) ^e
TM1				
Y58		1.39	H-bond donor to S314 ^{b,c,d}	N.A.
N69		1.50	H-bond donor to S317 ^{c,d}	N.A.
TM2				
S87		2.40	H-bond acceptor from Y324 ^{c,d}	N.A.
N92		2.45	H-bond acceptor from W176 ^{c,d}	N.A.
D97		2.50	NA ⁺ binding ^c	N.A.
Y110		2.63	Agonist effect ^e	N.A.
TM3				
R128	A	3.29	Counterion to α -phosphate ^{d,f}	>50,000
F131	A	3.32	Modulatory toward agonist, but not antagonist ^f	8
H132	A	3.33	Modulatory ^g	7
Y136	A	3.37	Modulatory ^g	6
S138		3.39	H-bond acceptor from N316 ^{c,d}	N.A.
H148		3.49	H-bond acceptor from R255 ^{c,d}	N.A.
TM4				
S172		4.46	H-bond acceptor from N92 ^{c,d}	N.A.
W176		4.50	H-bond donor to N92 ^{c,d}	N.A.
TM5				
T221	A	5.42	Small effect ^f	7
T222	A	5.43	Proximity to γ -phosphate ^f	5
F226	A	5.47	Binding of antagonists ^f	9
TM6				
R255		6.30	H-bond donor to H148 ^{c,d,h}	N.A.
Y273	A	6.48	Receptor activation ^{d,i}	>3,000
	F	6.48	Receptor activation rescued ^d	2
H277	A	6.52	Effect of agonist but not antagonist ^f	45
K280	A	6.55	Counterion to β -phosphate of ADP; action of PPADS ^{d,e,j}	810
TM7				
Y306	A	7.35	Modulatory toward agonist effects only ^d	5
	F	7.35	Modulatory ^{d,g}	6
Q307	A	7.36	H-bond acceptor from exocyclic NH of ADP ^{d,f}	210
R310	A	7.39	Counterion to α -phosphate ^{d,f}	>50,000
	K	7.39	Rescue of agonist (partial) and antagonist (full) effects ^f	190
S314	A	7.43	Activation, H-bond donor to adenine N1 or N3, ^{d,f} H-bond acceptor from Y58 ^b	>50,000
	T	7.43	Rescue of agonist effect ^f	5
N316		7.45	H-bond donor to S138 ^{c,d}	N.A.
S317	A	7.46	H-bond acceptor from N69; no pharmacological effect ^{d,f}	0.7
D320		7.49	Receptor activation ^{c,k}	N.A.
Y324		7.53	H-bond donor to S87 ^{c,d}	N.A.
Extracellular and cytoplasmic regions				
C42	A	N-terminal	Disulfide with C296 ^f	22,000
T47		N-terminal	Meta-binding site ^{c,f}	N.A.
C124	A	EL1	Disulfide with C202	>50,000
K125	A	EL1	No effect ^f	3
L157		IL2	G protein interaction ^{c,l}	N.A.
R195	A	EL2	No effect ^f	2
K196	A	EL2	No effect ^f	1
K198	A	EL2	Modulatory ^f	8
Y203	A	EL2	Agonist effects ^{d,m}	>3,000
	F	EL2	Rescue of agonist effects ^d	5
D204	A	EL2	Meta-binding site ^f	30
	N	EL2	Unable to rescue function ^f	270
	E	EL2	Unable to rescue function ^f	66
D208	A	EL2	No effect ^f	2
E209	A	EL2	Meta-binding site; H-bonding to ribose ^f	7800
	D	EL2	H-bonding rescued ^f	1
	Q	EL2	H-bonding rescued ^f	4
	R	EL2	H-bonding rescued ^f	2
R212	A	EL2	Small effect ^f	4
R285	A	EL3	Small effect ^f	4
R287	A	EL3	Meta-binding site; H-bonding to E209 ^f	6700
	K	EL3	Charge rescue (partial) of agonist effects ^f	34
	Q	EL3	Partial rescue ^f	240
	E	EL3	Disrupts E209 interaction ^f	>50,000
D289	A	EL3	No effect ^f	2
C296	A	EL3	Disulfide with C42 ^f	2300
D300	A	EL3	Agonist effects ^f	45
R301	A	EL3	No effect ^f	1
D329		C-terminal	Beginning of H8 ^{c,d,n}	N.A.

TABLE 4
Continued

Site	Mutation (if Performed)	Position ^a	Putative Interactions, Pharmacological Role	Potency Loss (Ratio for 2-MeSADP) ^a
R333		C-terminal	Required for Gq coupling ^e	N.A.
R334		C-terminal	Required for Gq coupling ^e	N.A.
K341		C-terminal	End of H8 ^{c,d,n}	N.A.

N.A. not applicable; EL extracellular loop; IL intracellular loop.

^a Residue identifier of format X.YZ refers to the TM X and residue YZ with respect to a highly conserved amino acid in each TM numbered 50 (Ballesteros et al., 2001; Costanzi et al., 2004); potency ratio shown is EC₅₀ at mutant receptor divided by EC₅₀ at the wild-type; the EC₅₀ of 2-MeSADP at the wild-type human P2Y₁ receptors was reported as 2.2 ± 0.5^f or 3 ± 1^g nM.

^b Applies only to ground state of receptor.

^c Not mutated, prediction from modeling only.

^d Costanzi et al. (2004).

^e Hoffmann et al. (2004).

^f Moro and Jacobson (2002) and references therein.

^g Modulatory toward effects of agonist and nucleotide antagonist.

^h Analogous to E(6.30) of the β₂ adrenergic receptor involved in an 'ionic lock' with R(3.50) but no electrostatic interaction with V(3.50) of the P2Y₁ receptor is possible; for the P2Y₁ receptor, a similarly placed interaction is predicted between R(6.30) and H(3.49).

ⁱ Mutation precludes only activation by agonist, but not binding of agonist or antagonist.

^j Guo et al. (2002).

^k Analogous to the N of the NPxxxY motif involved in activation of thyroid-stimulating hormone (Claeysen et al., 2002) and other receptors.

^l Moro et al. (1993).

^m Mutation precludes only activation and binding of agonist, but not binding of antagonist.

ⁿ An eighth helical region (cytoplasmic), by analogy to rhodopsin; a putative palmitoylation site (Cys) in this region is predicted for P2Y₄, P2Y₁₂, and P2Y₁₄, but not P2Y₁ receptors.

^o Ding et al. (2005).

A recent study on the purified and reconstituted human P2Y₁ receptor showed that it couples to Gα_q and Gα₁₁ but not to Gα_{i1}, Gα_{i2}, Gα_{i3}, or Gα_o (Waldo and Harden, 2004). This is consistent with a large body of studies showing that activation of this receptor evokes an increase in intracellular IP₃ levels and the release of intracellular Ca²⁺ stores, in a PTX-insensitive manner.

Northern blotting and RT-PCR revealed P2Y₁ receptor mRNA in most human tissues, including the brain, heart, placenta, lungs, liver, skeletal muscle, kidneys, pancreas, and various blood cells (Ayyanathan et al., 1996; Janssens et al., 1996; Léon et al., 1996). Two mRNA bands of 7.0 to 7.5 and 4.4 kb, whose expression pattern varied with the tissue examined, were seen (Ayyanathan et al., 1996; Léon et al., 1996). Quantitative RT-PCR indicated that expression was highest in the brain, prostate gland, and placenta and was also detected at varying levels in the pituitary gland, lymphocytes, spleen, heart, lung, liver, kidney, stomach, intestine, skeletal muscle, adipose tissue, and pancreas (Moore et al., 2001). In the study within the brain, P2Y₁ mRNA was highest in the nucleus accumbens, putamen, caudate nucleus, and striatum, with lower levels seen in the hippocampus, parahippocampal gyrus, globus pallidus, cingulate gyrus, and hypothalamus. Histochemical studies with a specific antibody also showed widespread distribution of the P2Y₁ receptor throughout the brain, with notable staining in the cerebral cortex, cerebellar cortex, hippocampus, caudate nucleus, putamen, globus pallidus, subthalamic nucleus, red nucleus, and mid-brain (Moore et al., 2000a). In addition, in post-mortem brain sections from persons with Alzheimer's disease, the P2Y₁-like immunoreactivity in the hippocampus and entorhinal cortex was localized to neurofibrillary tangles, neuritic plaques, and neuropil threads, which are characteristic Alzheimer's structures (Moore et al., 2000b). Western blots showed bands of 45, 90 and 180

kDa in the smooth muscle cells of the human left internal mammary artery and of 90 and 180 kDa in human umbilical vein endothelial cells (Wang et al., 2002).

A similar pattern of mRNA distribution is seen in rat tissues by Northern blotting (Tokuyama et al., 1995; Nakamura and Strittmatter, 1996). Mapping of the P2Y₁ mRNA by in situ hybridization has been performed across the chick brain (Webb et al., 1998), showing an abundant expression in many regions; at the cellular level this was located in many of the cell bodies of neurons, as well as in astrocytes. In rat brain, there is prominent P2Y₁-like immunoreactivity in neurons in Purkinje cells, the cerebellar cortex and hippocampus (Moran-Jiminez and Matute, 2000), ventral tegmentum (Krügel et al., 2001), midbrain, brainstem, and medulla (Fong et al., 2002). Glial cells in the brain also stain positive (Moran-Jiminez and Matute, 2000; Fong et al., 2002; Weick et al., 2003; Franke et al., 2004). The receptor is also expressed in sensory neurons (Nakamura and Strittmatter, 1996; Fong et al., 2002; Ruan and Burnstock, 2003; Gerevich et al., 2004), consistent with a potential role for P2Y₁ receptors in sensory reception. The presence of P2Y₁-like immunoreactivity (Fong et al., 2002) and mRNA (Buvinic et al., 2002; Kaiser and Buxton, 2002) in rat arterial endothelial cells is also consistent with the well-characterized vasodilatory activity of P2Y₁ receptors. P2Y₁-like immunoreactivity is also present in rat uterus epithelial cells (Slater et al., 2002), the pancreas (Coutinho-Silva et al., 2003), and glomeruli of rat kidney (Bailey et al., 2004). Northern blot analysis in chick embryos during the first 10 days of development showed that expression of P2Y₁ mRNA varied in a regulated manner in the limb buds, mesonephros, somites, brain, and facial primordia (Meyer et al., 1999). This suggests that the receptor may play a role in the development of each of these systems.

P2Y₁ receptor knockout mice have been generated by homologous recombination in two separate laboratories, and the reported phenotypes are identical (Fabre et al., 1999; Léon et al., 1999a) (see also Section.VII.J.). These mice are viable with no apparent abnormalities affecting their development, survival, and reproduction. Platelet counts and morphology are normal. In contrast, platelet shape change and aggregation to usual concentrations of ADP are completely abolished in these mice, whereas the ability of ADP to inhibit cAMP formation is maintained (Léon et al., 1999a). At higher concentrations of ADP (>10 μM), aggregation is observed without shape change, which is entirely due to P2Y₁₂ (Kauffenstein et al., 2001). Their bleeding time is mildly prolonged. These mice display resistance to systemic thromboembolism and to localized arterial thrombosis (Léon et al., 1999b, 2001; Lenain et al., 2003). Conversely, transgenic mice overexpressing the P2Y₁ receptor specifically in the megakaryocytic/platelet lineage have also been generated using the promoter of the tissue-specific platelet factor 4 gene (Hechler et al., 2003). This led to a phenotype of platelet hyper-reactivity *in vitro*. Moreover, overexpression of the P2Y₁ receptor enabled ADP to induce granule secretion, unlike in wild-type platelets, which suggests that the level of P2Y₁ expression is critical for this event and that the weak responses of normal platelets to ADP are due to a limited number of P2Y₁ receptors rather than to activation of a specific transduction pathway. In addition, transgenic mice display a shortened bleeding time and an increased sensitivity to *in vivo* platelet aggregation induced by infusion of a mixture of collagen and adrenaline (Hechler et al., 2003).

The P2Y₁ receptor is broadly expressed throughout the body. Thus, its gene knockout could have phenotypic consequences other than the sole hemostasis system. This is indeed the case with glucose homeostasis, since it has been found that the knockout mice display higher glucose levels as well as higher weight than wild-type mice (Léon et al., 2005).

For more information on the P2Y₁ receptor, see Table 5.

B. P2Y₂

P2Y₂ receptors, previously known as P_{2U}, have been cloned and pharmacologically characterized from human, rat, mouse, canine, and porcine cells or tissues (Lustig et al., 1993; Parr et al., 1994; Bowler et al., 1995; Rice et al., 1995; Chen et al., 1996; Zamboni et al., 2000; Shen et al., 2004). P2Y₂ receptors are fully activated by equivalent concentrations of ATP and UTP, whereas ADP and UDP are less effective agonists (Lustig et al., 1993; Parr et al., 1994; Lazarowski et al., 1995a). An exception is the porcine P2Y₂ receptor, which is relatively insensitive to ATP (Shen et al., 2004). UTP_γS has been shown to be a potent hydrolysis resistant agonist of P2Y₂ receptors (Lazarowski et al., 1995b). Suramin acts as a competitive antagonist of human and rat P2Y₂

receptors (Charlton et al., 1996; Lambrecht et al., 2002; Wildman et al., 2003). P2Y₂ receptors can directly couple to PLCβ₁ via G_{αq/11} protein to mediate the production of IP₃ and diacylglycerol (DAG), second messengers for calcium release from intracellular stores and PKC activation, respectively. Coupling of P2Y₂ receptors to other G protein subtypes has been reported (Baltensperger and Porzig, 1997; Murthy and Makhlof, 1998; Weisman et al., 1998; Bagchi et al., 2005). The P2Y₂ receptor is partially sensitive to PTX, a G_i/G_o protein inhibitor, and studies indicate that access to G_o protein is dependent upon association of the P2Y₂ receptor with α_vβ₃/β₅ integrins (Erb et al., 2001) and regulates nucleotide-induced chemotaxis (Bagchi et al., 2005). Expression of P2Y₂ receptor mRNA has been detected in human skeletal muscle, heart, brain, spleen, lymphocytes, macrophages, bone marrow, and lung, with lower expression levels detected in liver, stomach, and pancreas (Moore et al., 2001). Functional P2Y₂ receptors are expressed in epithelial, smooth muscle, and endothelial cells and in leukocytes, cardiomyocytes, osteoblasts, and cells derived from the peripheral and central nervous system, including Schwann cells, rat cortical neurons, oligodendrocytes, dorsal horn and cortical astrocytes, immortalized astrocytes, astrocytoma cells, and NG108-15 neuroblastoma × glioma hybrid cells (Bowler et al., 1995; Ho et al., 1995; Kirischuk et al., 1995; Rice et al., 1995; Berti-Mattera et al., 1996; Kim et al., 1996; Kunapuli and Daniel, 1998; Weisman et al., 1999; Pillois et al., 2002; Seye et al., 2002; Gendron et al., 2003; Kumari et al., 2003).

Site-directed mutagenesis of the P2Y₂ receptor has been used to demonstrate that replacement of positively charged amino acids in TM helices 6 and 7 with neutral amino acids decreases the potencies of ATP and UTP, suggesting that these domains play a role in binding the negatively charged moieties of nucleotide agonists (Erb et al., 1995) (see above). The P2Y₂ receptor undergoes agonist-induced desensitization in several cell types (Wilkinson et al., 1994; Garrad et al., 1998; Clarke et al., 1999; Otero et al., 2000; Velázquez et al., 2000; Santiago-Pérez et al., 2001), and mutagenesis studies indicate that deletion of structural motifs in the intracellular C-terminal domain of the P2Y₂ receptor that contain putative phosphorylation sites for GPCR kinase diminishes agonist-induced desensitization and internalization of the P2Y₂ receptor (Garrad et al., 1998). The P2Y₂ receptor also contains the consensus integrin-binding motif, Arg-Gly-Asp (RGD) in its first extracellular loop that facilitates P2Y₂ receptor colocalization with α_vβ₃/β₅ integrins when the P2Y₂ receptor is expressed in human 1321N1 astrocytoma cells that are devoid of endogenous G protein-coupled P2Y receptors (Erb et al., 2001). A mutant P2Y₂ receptor in which the RGD motif was replaced with Arg-Gly-Glu, a sequence that does not have high affinity for integrins, exhibited an EC₅₀ for nucleotide-induced calcium mobilization that was ~1000-fold

greater than that for the wild-type P2Y₂ receptor (Erb et al., 2001). The $\alpha_v\beta_3/\beta_5$ integrins are known to regulate angiogenesis and inflammatory responses including cell proliferation, migration, adhesion, and infiltration (Zhang et al., 2002b; Hutchings et al., 2003; Kannan, 2003; Li et al., 2003; Pidgeon et al., 2003), responses also mediated by P2Y₂ receptor activation (Wilden et al., 1998; Seye et al., 2002; Greig et al., 2003a,b; Schafer et al., 2003; Bagchi et al., 2005; Kaczmarek et al., 2005), suggesting that nucleotides may transactivate integrin signaling pathways by virtue of P2Y₂ receptor binding to integrins. The RGD sequence in the P2Y₂ receptor also has been shown to play an integrin-independent role in targeting of the receptor to the apical membrane of Madin-Darby canine kidney cells (Qi et al., 2005).

P2Y₂ receptor activation increases the synthesis and/or release of arachidonic acid (AA), prostaglandins and nitric oxide (NO) (Lustig et al., 1992; Pearson et al., 1992a,b; Xing et al., 1999; Xu et al., 2002b, 2003; Welch et al., 2003). In primary murine astrocytes, P2Y₂ receptors mediate the activation of calcium-dependent and calcium-independent PKCs and ERK1/2 that can activate cytosolic phospholipase A₂, leading to production of AA (Gendron et al., 2003; Xu et al., 2003), the precursor of eicosanoids, prostaglandins, and leukotrienes (Balsinde et al., 2002). Activation of P2Y₂ receptors in isolated UTP- or ATP-perfused rat hearts induces pronounced vasodilatation (Godecke et al., 1996), consistent with the role of P2Y₂ receptors in relaxation of smooth muscle through the endothelium-dependent release of NO and prostacyclin (Lustig et al., 1992; Pearson et al., 1992a,b). P2Y₂ receptor expression in smooth muscle cells is up-regulated by agents that mediate inflammation, including interleukin (IL)-1 β , interferon- γ , and tumor necrosis factor- α (Hou et al., 1999, 2000) and P2Y₂ receptor up-regulation has been shown to promote nucleotide-induced activation of PKC, cyclooxygenase, and MAPK (Koshiba et al., 1997; Turner et al., 1998; Seye et al., 2002). In addition to ERK1/2, P2Y₂ receptor activation can induce the phosphorylation of the stress-activated kinases JNK and p38 (Gendron et al., 2003). P2Y₂ receptor activation also induces p38- and ERK1/2-dependent up-regulation of genes that regulate cell survival in human astrocytoma cells (i.e., Bcl-2 and Bcl-xl) and genes that regulate neurite outgrowth in PC-12 cells (Chorna et al., 2004). Human neutrophil P2Y₂ receptors have been shown to regulate neutrophil degranulation induced by fibrinogen, independent of AA metabolites (Meshki et al., 2004), and P2Y₂ receptors have been suggested to play a role in the wound healing process (Burrell et al., 2003; Greig et al., 2003a,b).

Studies have indicated that P2Y₂ receptor-mediated MAPK activation in rat-1 fibroblasts and PC12 cells is dependent upon transactivation of the epidermal growth factor receptor (EGFR) via a Src/Pyk2-dependent pathway (Soltoff, 1998; Soltoff et al., 1998). In contrast, embryonic fibroblasts derived from Src^{-/-}, Pyk2^{-/-}, or

Src^{-/-}Pyk2^{-/-} mice have been used to demonstrate that Src and Pyk2 were essential for GPCR-mediated transactivation of the EGFR but not for GPCR-mediated MAPK activation (Andreev et al., 2001). G $\beta\gamma$ subunits have been shown to regulate Src-mediated transactivation of growth factor receptors (Luttrell et al., 1997), which may represent a common pathway whereby GPCRs stimulate cell proliferation. Other studies have identified two SH3-binding domains (i.e., PXXP motifs in which P is proline and X is any amino acid) in the intracellular carboxyl-terminal tail of the human P2Y₂ receptor that are necessary for transactivation of epidermal or platelet-derived growth factor (PDGF) receptors by ATP or UTP (Liu et al., 2004). Deletion of these SH3-binding domains inhibited nucleotide-induced P2Y₂ receptor colocalization with EGFR and UTP-induced EGFR transactivation (i.e., phosphorylation) but did not suppress ERK1/2 activation when the mutant receptors were expressed in 1321N1 astrocytoma cells (Liu et al., 2004), most likely because of the ability of the P2Y₂ receptor to also activate Src and ERK1/2 via PLC and integrin signaling pathways (Erb et al., 2001). Furthermore, Src coimmunoprecipitated with the P2Y₂ receptor in UTP-treated cells expressing the wild-type P2Y₂ receptor, but not the SH3-binding domain deletion mutant (Liu et al., 2004), strongly suggesting that activation of the P2Y₂ receptor promotes Src binding.

Activation of P2Y₂ receptors causes proliferation and/or migration of human epidermal keratinocytes, lung epithelial tumor cells, glioma cells, and smooth muscle cells (Wilden et al., 1998; Tu et al., 2000; Seye et al., 2002; Schafer et al., 2003; Greig et al., 2003b). P2Y₂ receptor activation also can induce cell cycle progression in smooth muscle cells from G₁ to S and M phases (Malam-Souley et al., 1996; Miyagi et al., 1996). HeLa cell proliferation in response to P2Y₂ receptor activation is associated with the PI3-K- and ERK1/2-dependent expression of the early response protein c-fos (Muscella et al., 2003). Consistent with a role for P2Y₂ receptors in cell proliferation, P2Y₂ receptor mRNA expression is down-regulated during cell differentiation (Martin et al., 1997). The P2Y₂ receptor is up-regulated in thymocytes as an immediate early gene response (Koshiba et al., 1997) and in injured and stressed tissues (Turner et al., 1997; Seye et al., 2002), and nucleotides induce ERK-dependent astrocyte proliferation under a variety of conditions including stretch- or stab-induced injury (Neary et al., 1994, 1996, 1999, 2003; Franke et al., 1999). Up-regulation of P2Y₂ receptors occurs in arteries after balloon angioplasty (Seye et al., 1997), in rat submandibular gland cells after short-term culture or upon duct ligation in vivo (Turner et al., 1997, 1998), and in a mouse model of Sjögren's syndrome (Schrader et al., 2004). Placement of a vascular collar around rabbit carotid arteries increases expression of P2Y₂ receptors in smooth muscle and endothelium that is associated with neointimal hyperplasia, monocyte infiltration, and

smooth muscle cell expression of the proliferative protein osteopontin, responses that were enhanced by application of the P2Y₂ receptor agonist UTP (Seye et al., 2002). In addition, P2Y₂ receptor up-regulation in endothelial cells increases the binding of monocytes to endothelial cells due to P2Y₂ receptor-mediated increases in the endothelial expression of vascular cell adhesion molecule-1 (Seye et al., 2003), a process that promotes vascular inflammation in atherosclerosis. Up-regulation of vascular cell adhesion molecule-1 was shown to be dependent on P2Y₂ receptor-mediated transactivation of vascular endothelial growth factor receptor-2 (KDR/Flk-1), a response that was inhibited by deletion of the SH3 binding motifs from the P2Y₂ receptor, demonstrating a mechanism whereby P2Y₂ receptors can cause inflammatory responses (Seye et al., 2004).

P2Y₂ receptor activation increases Cl⁻ secretion and inhibits Na⁺ absorption in epithelial cells, which has potential relevance for the treatment of cystic fibrosis, a disease that is caused by genetic defects in the gene for the CFTR, a major epithelial anion channel (Clarke and Boucher, 1992; Parr et al., 1994; Clarke et al., 2000; Kellerman et al., 2002). The synthetic P2Y₂ receptor agonist dCp₄U (INS37217) has been used to promote chloride and water secretion in tracheal epithelium and increase ciliary beat frequency and mucin release in human airway epithelium (Yerxa et al., 2002) and to stimulate subretinal fluid reabsorption in a rabbit model of retinal detachment (Meyer et al., 2002).

A P2Y₂ receptor knockout mouse that is defective in nucleotide-stimulated ion secretion in airway epithelial cells has been produced, confirming a role for the P2Y₂ receptor in the regulation of epithelial transmembrane ion transport (Cressman et al., 1999). In addition, P2Y₂ receptors have been shown to inhibit bone formation by osteoblasts (Hoebertz et al., 2002), and N-type calcium currents in neurons (Filippov et al., 1997; Brown et al., 2000a). P2Y₂ receptors also can induce α -secretase-dependent amyloid precursor protein processing in astrocytoma cells, suggesting a neuroprotective role (Camden et al., 2005). Collectively, studies with the P2Y₂ receptor and its signaling pathways have elucidated potential pharmacological targets in atherosclerosis, inflammation, cystic fibrosis, eye disease, osteoporosis, cancer, and neurodegenerative disorders.

For more information on the P2Y₂ receptor, see Table 6.

C. P2Y₄

Human (Communi et al., 1995; Nguyen et al., 1995; Stam et al., 1996), rat (Bogdanov et al., 1998b; Webb et al., 1998), and mouse (Lazarowski et al., 2001b; Suarez-Huerta et al., 2001) P2Y₄ receptors have been cloned and characterized (Communi et al., 2005b). UTP is the most potent activator of the recombinant human P2Y₄ receptor (Nicholas et al., 1996). GTP and ITP are approximately 10 times less potent than UTP

(Communi et al., 1996a). ATP behaves as a competitive antagonist (Kennedy et al., 2000). Up₄U (INS365) (Pendergast et al., 2001) and dCp₄U (INS37217) (Yerxa et al., 2002) are agonists of the human P2Y₄ receptor, whereas Ap₄A is inactive. On the contrary, the recombinant rat and mouse P2Y₄ receptors are activated equipotently by ATP and UTP and also with a lower potency by ITP, GTP and CTP (Bogdanov et al., 1998b; Webb et al., 1998). Pharmacological discrimination between rodent P2Y₄ and P2Y₂ receptors is thus difficult. No selective antagonist is available. Extracellular acidification enhanced the potency of ATP and UTP at rat P2Y₄, but not at rat P2Y₂ (Wildman et al., 2003). Zn²⁺ inhibited the ATP response at the rat P2Y₄ receptor but had no effect on rat P2Y₂ (Wildman et al., 2003). A study of chimeric human/rat P2Y₄ receptors showed that the structural determinants of agonism versus antagonism by ATP are located in the N-terminal domain and the second extracellular loop. Mutational analysis revealed that three residues in the second extracellular loop contribute to impart agonist property to ATP: Asn-177, Ile-183, and Leu-190 (Herold et al., 2004). Although their sequence is only distantly related to mammalian P2Y₄, *Xenopus* p2y8 (Bogdanov et al., 1997) and a turkey p2y (Boyer et al., 2000) could represent orthologs. They are activated with similar potencies by ATP, UTP, GTP, ITP, and CTP. The tp2y receptor is coupled to the stimulation of PLC and inhibition of adenylyl cyclase, indicating a dual coupling to G_{q/11} and G_{i/o}.

In the absence of reconstitution data, there is no hard molecular evidence of P2Y₄ receptor interaction with specific G proteins. The IP₃ response to UTP of the recombinant human P2Y₄ receptor is partially inhibited by PTX, at early times after agonist addition but not later (Communi et al., 1996a). This finding suggests that the P2Y₄ receptor couples mainly to a G_{q/11} protein and accessorially to a G_{i/o} protein. In sympathetic neurons injected with P2Y₄ cDNA, UTP inhibited the N-type Ca²⁺ current. This inhibition was relieved by PTX and by expression of the $\beta\gamma$ subunits of transducin (Filippov et al., 2003). These observations are consistent with the involvement of $\beta\gamma$ subunits released from G_{i/o} proteins in the function of the P2Y₄ receptor. In the same model, UTP inhibited the M-type K⁺ current in a PTX-insensitive way.

Among human organs, P2Y₄ mRNA was detected in placenta by Northern blotting (Communi et al., 1995) and was most abundant in the intestine, according to TaqMan quantitative RT-PCR (Moore et al., 2001). RT-PCR revealed the presence of P2Y₄ message in human umbilical vein endothelial cells, peripheral blood leukocytes (Jin et al., 1998b), fetal cardiomyocytes (Bogdanov et al., 1998a), and various cell lines derived from the human lung (Communi et al., 1999a). In the airway submucosal cell line 6CFSMEo-, an IP₃ response to UTP but not to ATP suggests a functional

expression of the P2Y₄ receptor (Communi et al., 1999a). In the rat, the expression of P2Y₄ message in heart and brain was much higher in neonates than in adults (Webb et al., 1998; Cheung et al., 2003). In situ hybridization on the adult rat brain neuronal P2Y₄ receptor mRNA was not detectable, it being present in the ventricular/choroid plexus system and strongly in the pineal gland, as well as apparently in astrocytes (Webb et al., 1998). In mouse, RT-PCR revealed that P2Y₄ message was the most abundant in stomach, liver, and intestine (Suarez-Huerta et al., 2001). Several histochemical studies have been performed using commercially available anti-rat P2Y₄ polyclonal rabbit antibodies, the specificity of which remains questionable (as is illustrated by Tung et al., 2004). Those studies suggest expression of the P2Y₄ receptor in rat dorsal root and trigeminal ganglia neurons (Ruan and Burnstock, 2003) and in the proximal convoluted tubular epithelium of the rat kidney (Turner et al., 2003). A combination of RT-PCR, Western blotting, immunohistochemistry, and pharmacology suggests that the P2Y₄ receptor is expressed and plays a role in the inner ear of rat, guinea pig, and gerbil (organ of Corti, stria vascularis marginal cells, and vestibular dark cells) (Teixeira et al., 2000; Sage and Marcus, 2002; Parker et al., 2003). These data are consistent with the hypothesis that nucleotides could play a role in the adaptation of the cochlea to noise exposure (Housley et al., 2002). Noise would increase the release of ATP and UTP in the scala media, leading to the inhibition of K⁺ release at the apical surface of the stria vascularis, presumably via activation of P2Y₄ receptors (Marcus and Scofield, 2001). In conclusion, a major site of P2Y₄ expression appears to be the intestine. Indeed RT-PCR has revealed that P2Y₄ message was particularly abundant in the human and murine intestine. This is consistent with the loss of the chloride secretory response to apical UTP in the jejunal epithelium of P2Y₄-null mice (Robaye et al., 2003).

P2Y₄-null mice have apparently normal behavior, growth, and reproduction (Robaye et al., 2003). The proportion of genotypes is consistent with X-linked Mendelian transmission. The chloride secretory response of the jejunal epithelium to apical UTP and ATP, measured in Ussing chambers, is abolished in P2Y₄-null mice. At the basolateral side, both P2Y₄ and P2Y₂ receptors are involved (Ghanem et al., 2005). In the colon, the epithelial secretion of chloride in response to UTP is mediated exclusively by the P2Y₄ subtype (Ghanem et al., 2005), whereas both P2Y₂ and P2Y₄ are involved in the UTP-induced secretion of potassium (Matos et al., 2005). Thus, in the intestine, the epithelial response to nucleotides is mediated mainly by the P2Y₄ receptor, but with a contribution of the P2Y₂ subtype, whereas in airways it involves mainly the P2Y₂ receptor (Cressman et al., 1999).

For more information on the P2Y₄ receptor, see Table 7.

D. P2Y₆

The mouse (Lazarowski et al., 2001b), rat (Chang et al., 1995; Nicholas et al., 1996), and human (Communi et al., 1996b) P2Y₆ receptors are UDP receptors. The p2y3 receptor is the avian ortholog of the mammalian P2Y₆ receptor and also displays selectivity for UDP (Webb et al., 1996a; Li et al., 1998). At the human P2Y₆ receptor, the rank order of potency of various nucleotides is as follows: UDP > UTP > ADP > 2-MeSATP ≫ ATP (Communi et al., 1996b). Adenine dinucleotides have little effect on P2Y₆, whereas diuridine triphosphate is a selective agonist of P2Y₆ (Pendergast et al., 2001). No selective competitive antagonist is available. Some aryl diisothiocyanate derivatives behave as potent insurmountable antagonists and exhibit selectivity for P2Y₆ compared with P2Y₁, P2Y₂, P2Y₄, and P2Y₁₁ receptors (Mamedova et al., 2004).

In the absence of reconstitution data, there is no hard molecular evidence of P2Y₆ interaction with specific G proteins. The IP₃ response to UDP of the recombinant P2Y₆ receptor is insensitive to PTX inhibition, suggesting a coupling to G_{q/11} (Chang et al., 1995; Robaye et al., 1997). A P2Y₆-mediated increase in cAMP has been reported, but it is probably an indirect effect mediated by prostaglandins, since it was at least partially inhibited by indomethacin (Köttgen et al., 2003). A unique feature of the P2Y₆ receptor compared with other P2Y subtypes is its slow desensitization and internalization (Robaye et al., 1997; Brinson and Harden, 2001). This can be explained by the short C-terminal sequence of P2Y₆ that contains a single threonine and misses the Ser333 and Ser334 that play a key role in the UTP-dependent phosphorylation, desensitization, and internalization of P2Y₄ (Brinson and Harden, 2001).

Northern blotting has revealed a rather wide tissue distribution of P2Y₆ mRNA. In particular, the P2Y₆ transcript has been found in human spleen, thymus, placenta, intestine, and blood leukocytes (Communi et al., 1996b) and in rat lung, spleen, stomach, intestine, and aorta (Chang et al., 1995). Consistent with those initial observations, the expression and potential role of the P2Y₆ receptor has been documented in placenta (Somers et al., 1999), vascular smooth muscle, epithelia, and immune cells. P2Y₆ message is present in smooth muscle cells cultured from the rat aorta (Chang et al., 1995; Hou et al., 2002). UDP acts as a growth factor for these cells (Hou et al., 2002). In P2X₁-deficient mice, the vasoconstrictor effect of ATP on mesenteric arteries is abolished (Vial and Evans, 2002), but a contractile effect of UTP and UDP is maintained and probably involves the P2Y₆ receptor, although the pharmacological profile of that response, characterized by the equipotency of UTP and UDP, is atypical. UDP and UDPβS induced the contraction of other vessels such as rat (Malmsjö et al.,

2003a) and human (Malmsjö et al., 2003b) cerebral arteries. The expression of P2Y₆ receptors on the basolateral side of rat colonic epithelial cells was demonstrated by immunohistochemistry and is involved in a sustained stimulatory effect of UDP on chloride secretion (Köttgen et al., 2003). In mouse gallbladder, apical UDP stimulates secretion of chloride, an effect that is maintained in P2Y₂-null mice and is likely to involve the P2Y₆ receptor (Cressman et al., 1999; Lazarowski et al., 2001b). Expression of the P2Y₆ receptor has also been demonstrated by RT-PCR and immunohistochemistry in human nasal epithelial cells (Kim et al., 2004), in which apical UDP stimulates chloride secretion (Lazarowski et al., 1997c). In human monocytic THP-1 cells, UDP stimulates the production of IL-8 (Warny et al., 2001). Furthermore, endogenous UDP released from those cells contributes to the IL-8 secretion induced by lipopolysaccharide (LPS), which is indeed partially decreased by P2Y₆ antisense oligonucleotides. UDP also acts on murine and human dendritic cells, where it increases cytosolic calcium, induces chemotaxis, and stimulates the release of chemokine CXCL8 (Marriott et al., 1999; Idzko et al., 2004). Northern blots revealed the expression of P2Y₆ mRNA in activated but not resting human CD4⁺ and CD8⁺ T cells (Somers et al., 1998). This expression was associated with a UDP-induced rise of [Ca²⁺]_i in the activated T cells only. In situ hybridization showed the presence of P2Y₆ mRNA in T cells infiltrating the lesions of patients with inflammatory bowel disease (IBD).

For more information on the P2Y₆ receptor, see Table 8.

E. P2Y₁₁

Among P2Y receptors, the human P2Y₁₁ has a unique profile: 1) it is the only P2Y receptor gene that contains an intron in the coding sequence; 2) the potency of its natural agonist ATP is relatively low; and 3) it is dually coupled to PLC and adenylyl cyclase stimulation. There does not seem to be a rodent ortholog of P2Y₁₁, confirming that this receptor subtype is quite different from the other members of the family. At the recombinant human P2Y₁₁ receptor, the rank order of potency with which nucleotides increase either cAMP or IP₃ is AR-C67085MX ≥ ATPγS ≈ BzATP > dATP > ATP > ADP (Communi et al., 1997, 1999b; Qi et al., 2001a). UTP, GTP, CTP, TTP, ITP, and dinucleotides (Ap₄A, Ap₅A, and Ap₆A) were completely inactive (although UTP has been claimed to be an agonist in some studies; see below and White et al., 2003). The EC₅₀ of ATP is in the 5 to 100 μM range, whereas in the same expression systems (i.e., 1321N1 or CHO cells), the EC₅₀ characterizing the activation of the other P2Y subtypes by their respective ligands is in the 10 to 500 nM range. Suramin behaves as a competitive antagonist of the human (h) P2Y₁₁ receptor with a K_i close to 1 μM (Communi et al., 1999b). The dog P2Y₁₁ receptor, the sequence of which shares

only 70% amino acid identity with hP2Y₁₁, has a completely different profile characterized by a greater potency of diphosphates compared with triphosphates (Zamboni et al., 2001). Indeed, the rank order of agonist potency is as follows: 2-MeSADP > ADPβS > ADP > ATP. A mutational analysis revealed that the change of Arg-265, located at the junction between TM6 and the third extracellular loop in the hP2Y₁₁, to Gln in cP2Y₁₁ is at least partially responsible for the diphosphate selectivity of the canine receptor (Qi et al., 2001b). Although the existence of a functional P2Y₁₁ receptor in *Xenopus* embryos was recently reported (C. Drew, unpublished data), an ortholog gene could not be detected in the murine genome, and there is no evidence of a functional P2Y₁₁ receptor in rat or mouse.

The hP2Y₁₁ gene differs from other P2Y genes by the presence in the coding sequence of a 1.9-kb intron that separates an exon encoding the first six amino acid residues from a second exon encoding the remaining part of the protein (Communi et al., 2001b). The hP2Y₁₁ gene is adjacent to the gene encoding the human ortholog of Ssf1, a nuclear protein playing an important role in *Saccharomyces cerevisiae* mating. Chimeric mRNA resulting from the cotranscription and intergenic splicing of the two genes is ubiquitously present in human organs. However, the fusion protein could only be detected after recombinant overexpression. Only half a dozen cases of intergenic splicing have been described in mammalian cells, and they are likely to represent an evolutionary tool to create new function via fusion of preexisting protein domains (Finta et al., 2002).

Activation of recombinant hP2Y₁₁ or cP2Y₁₁ receptors leads to an increase of both cAMP and IP₃, presumably via the dual activation of G_s and G_{q/11} (Communi et al., 1997, 1999b). Indeed, the use of various pharmacological tools (inhibition of PLC or prostaglandin synthesis, chelation of intracellular calcium, and down-regulation of PKC) has demonstrated that the cAMP increase is not merely an indirect consequence of rises in IP₃, [Ca²⁺]_i, and PKC activity (Suh et al., 2000; Qi et al., 2001a). However, PKC activation plays some role and amplifies the stimulation of adenylyl cyclase. In addition, it has been claimed that UTP acts via the hP2Y₁₁ receptor to induce an IP₃-independent Ca²⁺ mobilization that is sensitive to PTX inhibition, whereas the effect of ATP is not (White et al., 2003). This finding illustrates the fact that different agonists can recruit distinct signaling pathways via the same receptor.

Northern blotting has revealed a significant expression of P2Y₁₁ mRNA in human spleen and to a lesser extent intestine and liver (Communi et al., 1997, 2001b). According to a quantitative RT-PCR study, P2Y₁₁ message is most abundant in human brain and pituitary (Moore et al., 2001). It is also present in B lymphocytes from patients with chronic lymphocytic leukemia (Conigrave et al., 2001) as well as in human HL-60 (Communi et al., 1997, 2000) and NB4 (van der Weyden et al.,

2000b) promyelocytic leukemia cells. In the HL-60 cell line, P2Y₁₁ message is up-regulated by all the agents that induce granulocytic differentiation, such as retinoic acid and granulocyte colony-stimulating factor (Comuni et al., 2000). ATP has been shown to induce the differentiation of HL-60 cells into neutrophil-like cells (Jiang et al., 1997a). This action was associated with a rise in cAMP and the rank order potency of various nucleotides is consistent with that of recombinant P2Y₁₁ (Conigrave et al., 1998). P2Y₁₁ mRNA is also expressed in human monocyte-derived dendritic cells, in which ATP induces a semimaturational state characterized by increased surface expression of costimulatory molecules, inhibition of the production of proinflammatory cytokines such as IL-12, stimulation of IL-10 production, and modification of the repertoire of chemokine and chemokine receptor expression, resulting in a change of migratory behavior (Wilkin et al., 2001, 2002; La Sala et al., 2002; Schnurr et al., 2003). Semimature dendritic cells can orient CD4⁺ T lymphocytes toward a Th2 rather than a Th1 response or induce tolerance.

For more information on the P2Y₁₁ receptor, see Table 9.

F. P2Y₁₂

The human (Hollopeter et al., 2001; Savi et al., 2001; Zhang et al., 2001), rat (Hollopeter et al., 2001), and mouse (Foster et al., 2001) P2Y₁₂ receptors have been identified and characterized. ADP is the natural agonist of this receptor, whereas conflicting results were reported concerning the effects of ATP and its triphosphate analogs. For diphosphates, the rank order of agonist potency in all cases reported is 2-MeSADP \gg ADP > ADP β S. Concerning ATP and its analogs, they were found to be agonists either in native P2Y₁₂-expressing cells (Simon et al., 2001; Unterberger et al., 2002) or in some heterologously transfected cells (Takasaki et al., 2001; Zhang et al., 2001; Simon et al., 2002). However, in platelets there have been reports over a long period that ATP and a wide range of its triphosphate analogs behave as antagonists of ADP-induced adenylyl cyclase inhibition (reviewed by Gachet, 2001). This has recently been confirmed: ATP and its triphosphate analogs are antagonists of the P2Y₁₂ receptor both in human and mouse platelets, provided care is taken to remove contaminants and to prevent enzymatic production of ADP or 2-MeSADP (Kauffenstein et al., 2004). There are currently, therefore, two alternative interpretations extant of the action of adenosine triphosphates on the P2Y₁₂ receptor (either native or recombinant). One (Kauffenstein et al., 2004) is that, both in the platelet and in nucleated (nonplatelet) cells, these agents are antagonists: an apparent agonist action of triphosphates would then be due in all cases to a secondary introduction of diphosphate, as just noted. The second (Barnard and Simon, 2001; Simon et al., 2002) is that the triphosphates are intrinsically agonistic at the

P2Y₁₂ receptor generally, but that they behave as apparent antagonists in the platelet, due to the much lower receptor density there (or possibly to an absence in the platelet of some modifying component involved in nucleated cells). The latter interpretation has been made also (see section VI.A.) for a parallel difference in adenosine triphosphate behavior of the P2Y₁ receptor in the platelet and in most other cell types tested.

Behavior of ATP and ATP γ S rather similar to that reported for these agents at P2Y₁₂ in the platelet can be obtained in a system reconstituted with purified P2Y₁₂ receptor protein (see section III.A.), in which enzymatic degradation or interconversion of the nucleotides is not possible (Bodor et al., 2003). There, also, one of the same explanations just given for the platelet case could hold, e.g., the effective density of functional P2Y₁₂ receptors in the vesicle surface may be too low for those agents to show intrinsic agonism. The general dichotomy discussed here need not apply to all adenosine triphosphates: ATP γ S may behave at P2Y₁₂ receptors in general as an antagonist, that action being shown clearly in transfected 1321 N1 cells as well as in the platelet (Kauffenstein et al., 2004) (ATP γ S has not yet been tested with full precautions elsewhere).

A clear agonist action of adenosine triphosphates of P2Y₁₂ receptors other than in platelets was measured for adenylate cyclase inhibition in the cases noted above (Simon et al., 2001, 2002; Unterberger et al., 2002) with all of the precautions used by Kauffenstein et al. (2004) and with a demonstration of negligible loss of ATP occurring during the assay. Furthermore, in the direct transduction by the P2Y₁₂ receptor in neurons through the N-type Ca²⁺ channel (see section III.B.3.), in which a fast application of pure 2-MeSATP is made with constant perfusion flow, in contact with only a low cell density and with an assay time of 50 ms, 2-MeSATP (EC₅₀ 0.042 nM) is a definite agonist, 2.5 times more potent than 2-MeSADP (Simon et al., 2002). Clearly this effect cannot be due to an artifactual introduction of the diphosphate. In conclusion, the situation concerning the agonist selectivity of the P2Y₁₂ receptor is controversial, may vary with the cell type, and is at present left open here.

The P2Y₁₂ receptor is mostly expressed in the megakaryocyte/platelet lineage in which it is the molecular target of the active metabolite of the antiplatelet drug clopidogrel (Savi and Herbert, 2005; see also below). This metabolite covalently binds cysteine residues of the extracellular loops resulting in inhibition of ligand binding (Savi et al., 2001). Ticlopidine and clopidogrel are efficient antithrombotic drugs of the thienopyridine family of compounds. A third antithrombotic thienopyridine, CS-747 or prasugrel, is currently under clinical evaluation (Niitsu et al., 2005). Potent direct competitive P2Y₁₂ antagonists also exist, including the AR-C69931MX compound named cangrelor as well as other AR-C compounds, which are all ATP analogs (Ingall et

al., 1999). Of these, AZD6140 is a nonphosphorylated and orally active compound currently under clinical evaluation (Peters and Robbie, 2004).

In addition to the platelet lineage, the P2Y₁₂ receptor has also been shown to be expressed in subregions of the brain (Hollopeter et al., 2001), but its function there is not yet known. Glial cells (Fumagalli et al., 2003; Sasaki et al., 2003; Bianco et al., 2005), brain capillary endothelial cells (Simon et al., 2001), smooth muscle cells (Wihlborg et al., 2004), and chromaffin cells (Ennion et al., 2004) express P2Y₁₂ receptors. The precise role in these locations is still under study.

The platelet P2Y₁₂ receptor is coupled to G_{α₁₂} (see also section VII.J.), as was first been shown by photolabeling with radiolabeled GTP (Ohlmann et al., 1995) and confirmed in G_{α₁₂}-deficient mouse platelets (Jantzen et al., 2001) as well as in reconstituted systems (Bodor et al., 2003). In the latter case, G_{α₁₂} was found to be the preferred G_α subunit, whereas G_{α₁₁} and G_{α₁₃} were poorly effective. No coupling with G_z was observed.

P2Y₁₂ knockout mice have been generated (Foster et al., 2001; André et al., 2003; see also section VII.J.), which display the phenotype of clopidogrel-treated animals, i.e., prolonged bleeding time, inhibition of platelet aggregation to ADP, and resistance to arterial thrombosis in various models (Conley and Delaney, 2003). In humans, molecular defects of this receptor exist, which result in hemorrhagic syndromes. Four families of patients have been described so far with essentially the same phenotype. Among these, three have a defect in receptor expression (Cattaneo et al., 1992, 1997; Nurden et al., 1995), whereas in one family, a mutant form of the receptor is expressed with defective function (for a recent review, see Cattaneo, 2005; Cattaneo et al., 2003).

For more information on the P2Y₁₂ receptor, see Table 10.

G. P2Y₁₃

The human (Communi et al., 2001a; Zhang et al., 2002a), mouse (Zhang et al., 2002a), and rat (Fumagalli et al., 2004) P2Y₁₃ receptors have been identified and characterized (see also Communi et al., 2005a). ADP and Ap₃A are naturally occurring agonists of the P2Y₁₃ receptor. IDP is also a potent agonist of the murine P2Y₁₃ receptor, but is 10-fold less potent than ADP on the human one (Zhang et al., 2002a). Ap₄A, Ap₅A, and Ap₆A are inactive. When contaminating ADP was enzymatically removed and testing was performed over a short period, ATP behaved as a weak partial agonist (Marteau et al., 2003). As described for the P2Y₁ receptor, the activity of ATP may vary according to the level of expression of the P2Y₁₃ receptor in different recombinant systems. The relative potencies of ADP and 2-MeSADP differed according to assays used. 2-MeSADP was more potent than ADP in competing with [³³P]2-MeSADP on intact 1321N1 cells expressing hP2Y₁₃ and in stimulating binding of GTPγ[³⁵S] to membranes of the same

cells, whereas ADP was more potent than 2-MeSADP on the rat P2Y₁₃ (Fumagalli et al., 2004). In CHO-K1 cells expressing hP2Y₁₃, ADP and 2-MeSADP produced an equipotent inhibition of cAMP accumulation. These discrepancies suggest that the P2Y₁₃ receptor might exist in multiple active conformations characterized by differences in affinity for 2-MeSADP versus ADP, kinetics, and preference for G proteins. The antiplatelet and antithrombotic action of clopidogrel is mediated by an active metabolite. That metabolite has been shown to inhibit the binding of [³³P]2-MeSADP to hP2Y₁₂ with an IC₅₀ of 100 nM (Savi et al., 2001), but it had no effect on hP2Y₁₃ up to 2 μM (Marteau et al., 2003). Cangrelor (AR-C69931MX) is currently in development as an antiplatelet and antithrombotic agent. It is an ATP derivative that inhibits platelet aggregation by ADP at nanomolar concentrations (Ingall et al., 1999). It was previously believed to be a selective antagonist of the hP2Y₁₂ receptor (IC₅₀ = 2.4 nM) (Takasaki et al., 2001), but in the same range of nanomolar concentrations, it is also an antagonist of human and rat P2Y₁₃ receptors (Marteau et al., 2003; Fumagalli et al., 2004). Two other P2Y₁₂ antagonists, Ap₄A and 2-MeSAMP, are also antagonists of the P2Y₁₃ receptor (Marteau et al., 2003).

The effects of ADP mediated by the recombinant P2Y₁₃ receptor were all inhibited by PTX: increased binding of GTPγ[³⁵S], inhibition of cAMP formation, ERK1/2 phosphorylation, and accumulation of IP₃ in cells coexpressing G_{α₁₆} (Communi et al., 2001a; Marteau et al., 2003). This finding suggests that the P2Y₁₃ receptor is primarily coupled to a G_{i/o} protein. The only exception was the increased cAMP formation observed at high ADP concentrations, and that presumably results from promiscuous coupling to G_s, a phenomenon observed with other recombinant G_{i/o}-coupled receptors, such as the α₂-adrenergic receptor (Communi et al., 2001a). More direct evidence for the coupling to G_{i/o} derives from the measurement of [Ca²⁺]_i increases in HEK cells coexpressing various chimeric G proteins (Zhang et al., 2002a). A significant stimulation by ADP was obtained in cells expressing either G_{α_{q/i}} or G_{α_{q/13}}, that is, G_{α_q} in which the five C-terminal residues have been replaced by the corresponding sequence in either G_{α_{11/2}} or G_{α₁₃}, respectively.

P2Y₁₃ mRNA was amplified by RT-PCR in several human organs. Signals were the most intense in spleen and brain (Communi et al., 2001a). In dot blot analysis, the spleen gave the most intense positive signal, followed by placenta, liver, bone marrow, lung, and various brain regions (Zhang et al., 2002a). Quantitative RT-PCR revealed a significant expression in human monocytes, T cells, and dendritic cells derived from blood monocytes or bone marrow, but not in human platelets (Zhang et al., 2002a; Wang et al., 2004a). Northern blots were positive for murine spleen, brain, liver, and heart. In the rat, again the RT-PCR signal was the most intense in the spleen (Fumagalli et al., 2004).

P2Y₁₃-null mice have been generated recently (A. Ben Addi and B. Robaye, unpublished data). No phenotype has been characterized so far.

For more information on the P2Y₁₃ receptor, see Table 11.

H. P2Y₁₄

From a phylogenetic and structural point of view, the P2Y₁₄ receptor (previously known as GPR105 or UDP-glucose receptor) lies with the P2Y₁₂ and P2Y₁₃ receptors in the second main branch of the P2Y receptor family and is 47% identical to these receptors. The gene for this receptor has been found in human chromosome 3q24–3q25 where a cluster of other related GPCRs, consisting of P2Y₁-P2Y₁₂, and P2Y₁₃ receptors and the orphan receptors GPR87, GPR91, and H963 have been found (Abbracchio et al., 2003).

The P2Y₁₄ receptor is activated by UDP-glucose as well as UDP-galactose, UDP-glucuronic acid, and UDP-N-acetylglucosamine but not by uridine or adenine nucleotides (Chambers et al., 2000; Harden, 2004). Of these endogenous ligands, to date only UDP-glucose has been shown to be released extracellularly by a variety of cell lines (Lazarowski et al., 2003b). At present, no selective antagonists are available, although it has to be underlined that the currently available P2 receptor antagonists have not been tested on this receptor. No radioligand binding assay is available for quantification of P2Y₁₄ receptor binding sites.

The complete sequences of the rat (VTR-15-20; Charlton et al., 1997; Freeman et al., 2001) and mouse orthologs (Freeman et al., 2001) have also been described. The rat and mouse orthologs show 80 and 83% amino acid identity, respectively, with human P2Y₁₄ and show similar agonist pharmacology (Freeman et al., 2001).

The P2Y₁₄ receptor couples to the G_{i/o} family of G proteins. In particular, data obtained on the recombinant receptor in HEK-293 cells show coupling to G α subunits of the G_{i/o} family (G α ₁₆, G α _{qo5}, and G α _{qi5}), but not to G_s family or to endogenous G_{q/11} proteins (Moore et al., 2003). Stimulation of native P2Y₁₄ receptors in primary rat cortical astrocytes as well as in murine N9 and rat primary microglial cells results in transient intracellular calcium increases (Fumagalli et al., 2003; Bianco et al., 2005), although the mechanisms at the basis of this transductional effect are not known yet.

P2Y₁₄ mRNA is widely distributed in the human body, with moderate to high levels observed in placenta, adipose tissue, stomach, intestine, selected brain regions (e.g., corpus striatum, cerebellum, caudate nucleus, hippocampus, and hypothalamus), spleen, lung, heart, bone marrow, and thymus. RT-PCR also revealed expression in brain glial cells and prominent expression in neutrophils, lymphocytes and megakaryocytic cells (Chambers et al., 2000; Moore et al., 2003). P2Y₁₄ receptor mRNA was high in immature monocyte-derived and low in ma-

ture monocyte-derived dendritic cells, suggesting a role for the receptor and its agonists in dendritic cell activation (Skelton et al., 2003).

Antiserum to a sequence in the first extracellular domain of P2Y₁₄ was used to isolate a population of hematopoietic cells restricted to bone marrow. Conditioned media from bone marrow stroma induce receptor activation and chemotaxis, suggesting a role for P2Y₁₄ as a chemoattractant receptor (Lee et al., 2003a). Antibody against the carboxyl terminus of the P2Y₁₄ receptor was used to demonstrate broad distribution in post-mortem human brain, with glial cells as the primary site of expression, suggesting a role for this receptor in neuroimmune function (Moore et al., 2003). In the rat, P2Y₁₄, which is abundantly expressed in brain, has been reported to be regulated by immunological challenge (Charlton et al., 1997; Moore et al., 2003), suggesting that this receptor may link the humoral and nervous system responses to infection and inflammation. Consistent with this result, exposure of murine N9 cells to LPS resulted in a highly significant increase of receptor function, as suggested by the increase in the percentage of cells responding to UDP-glucose with intracellular calcium transients (Bianco et al., 2005).

For more information on the P2Y₁₄ receptor, see Table 12.

VII. Receptor Distribution and Function

A recent article reviews at depth the cellular distribution and functions of P2 receptor subtypes in different systems and cell types based on studies of subtype mRNA and proteins as well as functional data (Burnstock and Knight, 2004). Additional information can also be found above in individual P2Y receptor subsections and in individual summary tables. A general review of P2Y receptor distribution and function in mammalian organs and systems is presented below. (For references, see Burnstock and Knight, 2004.)

A. Excitable Cells, Nerves, Glial Cells, and Muscle

P2Y₁ receptors are widespread in many regions of the brain, whereas the P2Y₂ receptors have been localized on pyramidal neurons in the hippocampus and prefrontal cortex, on supraoptic magnocellular neurosecretory neurons in the hypothalamus, and on neurons in the dorsal horn of the spinal cord. In addition, mRNA but not protein has been reported for P2Y₄ and P2Y₆ receptor subtypes in the cerebellum and hippocampus, whereas P2Y₁₂ receptor mRNA has also been described in the cerebellum and P2Y₁₃ in the cortex. In the periphery, P2Y_{1,2,4,6} receptors on subpopulations of sympathetic neurons, P2Y₂ and P2Y₄ receptors in intracardiac ganglia, and P2Y₁ and P2Y₂ receptors on sensory neurons (although P2Y₄ and P2Y₆ mRNA have also been reported) have been described, whereas P2Y₁ receptors appear to be the dominant subtype on enteric neurons.

P2Y_{1,2,4,6} functional receptors have been found on astrocytes in the central nervous system (Neary et al., 1994, 1996, 1999, 2003; Bolego et al., 1997; Centemeri et al., 1997; Fumagalli et al., 2003) and also on microglia where functional P2Y₁₂ receptors have also been identified (for review, see Abbracchio and Verderio, 2006). P2Y₁ and P2Y₂ receptors have been located in Schwann cells and oligodendrocytes, in which functional P2Y₁₂ receptors also appear to be present. P2Y₂ (and/or P2Y₄) receptors are expressed on enteric glial cells. There is also emerging evidence for P2Y receptors on stem cells.

All cloned P2Y receptors have been found in both healthy and failing human hearts (Banfi et al., 2005) to support functional roles in myocardial function (Vassort, 2001). Whereas the dominant P2 receptor subtype on smooth muscle is P2X₁, P2Y receptor subtypes are also present, notably P2Y₁ and/or P2Y₂ receptors on visceral smooth muscle and P2Y_{1,2,4,6} receptors on vascular smooth muscle. P2Y₁ receptors are present in developing skeletal muscle with evidence for P2Y₂ receptors on the myotube C2C12 cell line.

B. Immune Cells

ATP stimulates production of prostaglandin and has mitogenic actions on thymocytes largely via P2Y₂ receptors, although P2Y₁ receptor mRNA is also present. P2Y_{2,6,11,13} receptor mRNAs have been identified in whole spleen.

P2Y₂ receptors are the dominant subtype in macrophages, although the presence of P2Y_{1,4,12} receptors on alveolar macrophages and P2Y₁₁ receptors on human macrophages has also been noted. ATP and UTP acting via P2Y₁ and P2Y₂ receptors promote adhesion of neutrophils to endothelial cells.

P2Y_{1,2,4,6,11} receptor mRNA have been identified with RT-PCR in both eosinophils and lymphocytes. Several P2Y subtypes are expressed in human monocyte-derived dendritic cells, in particular the P2Y₁₁ receptor that mediates the semimaturation of these cells in response to ATP (Wilkin et al., 2001; La Sala et al., 2002; Schnurr et al., 2003).

P2Y_{1/2/4/6} receptors have been identified in monocytes, and UDP leads to production of IL-8 (Warny et al., 2001). ATP action via P2Y₁ and P2Y₂ receptors leads to degranulation and release of histamine from mast cells as well as cell migration and chemoattraction.

C. Endocrine, Adipose, and Exocrine Cells

The P2Y₂ receptors are dominant in the anterior pituitary in which they modulate prolactin release, and P2Y₁ receptors have been identified in the pineal gland. P2Y₂ receptors mediate regulation of aldosterone secretion in the adrenal gland. In the thyroid gland, P2Y_{2,4,6} receptors mediate cell proliferation, whereas P2Y₂ receptors are involved in insulin release from pancreatic islet β cells.

P2Y₂ receptors are present in testicular Sertoli and in Leydig cells involved in estradiol and testosterone secretion. P2Y₂ receptors are expressed in ovary and placenta. P2Y₂ receptors mediate antagonism of estradiol and progesterone secretion from granulosa cells in the ovary.

Brown adipocytes express P2Y_{1,2,4} receptors involved in regulation of lipogenesis. P2Y₂ receptors involved in regulation of ionic balance are dominant in salivary and lachrymal glands.

In sweat glands and exocrine pancreas, P2Y_{1,2,4} receptors are present. They have been implicated in regulation of secretion.

D. Gut, Liver, and Biliary System

P2Y₁ and P2Y₁₁ receptors appear to mediate the purinergic component of nonadrenergic, noncholinergic relaxation of smooth muscle in the gastrointestinal tract, whereas P2Y_{1,2,4,6} receptors on gut epithelial cells may mediate ion secretion, although P2Y₄ seems to play a major role, as demonstrated in knockout mice. P2Y₁ receptors have also been identified in intrinsic enteric neurons in both myenteric and submucosal plexuses. P2Y_{1,2,4,6,13} receptors have been described in hepatocytes and may serve to regulate gluconeogenesis and glycolysis. P2Y_{2,4,6} receptors are present on bile duct epithelium; in addition, P2Y₁ receptor mRNA is present in gallbladder epithelium. UTP has been shown to stimulate Cl⁻ secretion and modulate bile release.

E. Kidney and Bladder

P2Y receptors are richly expressed in all regions of the kidney tubule and glomerulus. P2Y_{1,4,6} receptors in the proximal convoluted tubule and loop of Henle are involved in reabsorption of water, ions, and nutrients. The P2Y₂ receptor is dominant in the distal convoluted tubule concerned with secretion of ions, acids, and toxins whereas P2Y_{1,2,4,6} receptors appear to be involved in transport of water and ions in the collecting duct. In the glomerulus, P2Y_{1,2,4,11,12} subtypes are all expressed on mesangial cells, P2Y_{1,2,6} on podocytes, and P2Y₂ on endothelial cells. In rat bladder, ATP exerted an immediate and transient contraction, followed by a slower sustained relaxation, which was completely abolished by the G protein blocking agent, guanosine 5'-O-(2-thiodiphosphate), suggesting a functional role for P2Y receptors (Bolego et al., 1995). Evidence for purinergic functional regulation of human urinary bladder, which may have potential therapeutic implications for human incontinence, has been also provided (Palea et al., 1993).

F. Lung

P2Y₂ and P2Y₆ receptors on cultured smooth muscle cells from the lung induce an increase and a decrease in smooth muscle proliferation, respectively. P2Y₂ receptors are the predominant receptor subtype in lung epithelial cells and are involved in mucin secretion and

mucociliary clearance. In addition, P2Y₁ and P2Y₆ receptor subtypes have been identified on respiratory epithelium cell lines.

G. Bone and Cartilage

P2Y₂ receptors on osteoblasts mediate inhibition of bone formation, whereas ADP acting via P2Y₁ receptors on osteoclasts increase bone resorption. Chondrocytes express P2Y₁ and P2Y₂ receptors, which mediate cartilage resorption and prostaglandin production.

H. Skin

P2 receptors play a major role in cell turnover of keratinocytes in stratified epithelium. P2Y₁ and P2Y₂ receptors modulate cell proliferation in basal cells.

I. Endothelial Cells

P2Y₁ and/or P2Y₂ receptors are dominant on vascular endothelial cells, where they mediate release of NO and subsequent vasodilatation as well as cell proliferation and the expression of adhesion proteins for monocytes. In some blood vessels, P2Y_{4,6,11,12} receptors have also been found.

J. Special Senses

In the eye, P2Y₂ receptors mediate trophic events in the retina and cornea, whereas P2Y₁ receptors have been implicated in regulation of fluid secretion in the ciliary body. In the inner ear, P2Y receptors have been implicated in cochlear function, particularly P2Y₂ and P2Y₄ receptors in vestibular dark cells and stria vascularis marginal cells. P2Y receptors also appear to be involved in olfactory function, especially P2Y₂ and P2Y₆ receptors on nasal epithelium.

K. Platelets

Platelets express three nucleotide receptors: the P2X₁ cation channels activated by ATP, and two GPCRs, P2Y₁ and P2Y₁₂, both activated by ADP, which perhaps represent the most studied P2Y receptors in a native system (see also above). Each of these receptors has a selective role during platelet activation (Hechler et al., 2005), which has implications for their role in thrombosis (Gachet and Hechler, 2005). We will not expand here on the role of the P2X₁ receptor, which is not negligible, being involved in platelet activation by collagen and in the thrombosis of small arteries (Hechler et al., 2003; Mahaut-Smith et al., 2004). However, its deficiency does not induce a defect in normal hemostasis and does not modify the platelet responses to ADP. ADP-induced platelet aggregation is under the control of P2Y₁ and P2Y₁₂. Coactivation of both receptors is necessary for normal ADP-induced platelet aggregation since separate inhibition of each of them by selective antagonists results in dramatic inhibition of aggregation (Jin and Kunapuli, 1998; Gachet, 2001; Hechler et al., 2005).

1. The P2Y₁ Receptor Initiates Platelet Activation and Aggregation. The presence and role of the P2Y₁ receptor in platelets, initially suggested by the detection of P2Y₁ mRNA in both megakaryoblastic cells and platelets (Léon et al., 1997), was confirmed using selective P2Y₁ antagonists (Léon et al., 1997; Hechler et al., 1998b) and by studies in P2Y₁ receptor-deficient mice (Fabre et al., 1999; Léon et al., 1999a). There are ~150 P2Y₁ receptors/platelet (Baurand et al., 2001; Baurand and Gachet, 2003), a number that is very low compared, for instance, with thromboxane prostanoid receptors or with the thrombin receptor PAR-1 (1000–2000 receptors/platelet). The P2Y₁ receptor coupled to G_{α_q} triggers calcium mobilization from internal stores, which results in platelet shape change and weak and transient aggregation in response to ADP (Hechler et al., 1998a,b; Jin et al., 1998a; Savi et al., 1998). In addition, the P2Y₁ receptor participates in aggregation induced by collagen, as shown by the reduced amplitude of aggregation and the prolongation of the lag phase from the addition of collagen to the onset of aggregation in P2Y₁ knockout platelets (Léon et al., 1999a). P2Y₁ receptors also play a key role in collagen-induced shape change when thromboxane (TX) A₂ formation is prevented (Mangin et al., 2004). Examination of the morphological changes during platelet aggregation indicates that the P2Y₁ receptor is involved in the centralization of platelet granules induced by ADP and the formation of filopodia in platelets activated with low concentrations of agonists such as TXA₂ or thrombin (Eckly et al., 2001). Overall, the P2Y₁ receptor mediates weak responses to ADP and has a crucial role in the early steps of platelet activation induced by ADP or collagen.

2. The P2Y₁₂ Receptor Completes and Amplifies Platelet Activation and Aggregation. Several lines of evidence suggested the presence of a second platelet ADP receptor coupled to G₁₂ and responsible for adenylyl cyclase inhibition, well before its identification (Gachet, 2001). Indeed, molecules such as clopidogrel or the ATP analogs of the AR-C series selectively inhibit the G₁-coupled ADP response without any impact on the P2Y₁ receptor-mediated effects (Hechler et al., 1998a; Jin et al., 1998a; Savi et al., 1998), whereas P2Y₁ receptor antagonists inhibit ADP-induced platelet aggregation without inhibiting the effect of ADP on adenylyl cyclase activity (Daniel et al., 1998; Hechler et al., 1998b; Savi et al., 1998). In addition, the P2Y₁ receptor is normal, at the genetic and pharmacological levels, in a patient with congenital impairment of platelet responses to ADP (Cattaneo et al., 1992; Léon et al., 1999b). Lastly, in P2Y₁ receptor-deficient mice, ADP is still able to inhibit platelet adenylyl cyclase activity (Fabre et al., 1999; Léon et al., 1999a). The requirement of this receptor to complete aggregation to ADP was confirmed by the generation of P2Y₁₂ receptor-deficient mice, which display a defect in platelet aggregation in response to ADP, although shape change is conserved (Foster et al., 2001;

André et al., 2003). Numerous studies established the key role of this receptor not only for completion of aggregation in response to ADP but also for the ADP-dependent amplification of platelet aggregation induced by other agents acting at the G_{α_q} -coupled serotonin receptor 5HT_{2A} (Savi et al., 1998) or the G_{α_q} and $G_{12/13}$ -coupled TXA₂ and PAR-1 receptors (Dorsam et al., 2002). Similarly, the P2Y₁₂ receptor is an important cofactor of platelet aggregation and secretion induced by cross-linking of the Fc γ RIIIa receptor with specific antibodies (Gratacap et al., 2000), or when platelets are activated by collagen through the GPVI/tyrosine kinase/PLC- γ 2 pathway (Nieswandt et al., 2001). The P2Y₁₂ receptor is also involved in potentiation of platelet secretion (Cattaneo et al., 1997, 2000) and mediates the stabilization of platelet aggregates induced by thrombin (Trumel et al., 1999) or TXA₂ (Eckly et al., 2001).

The P2Y₁₂ receptor is coupled to the $G_{\alpha_{i2}}$ G protein subunit (Ohlmann et al., 1995; Bodor et al., 2003). Downstream of $G_{\alpha_{i2}}$, several signaling pathways are involved in amplification mechanisms. First, inhibition of cAMP production, although not sufficient to trigger platelet aggregation, has a facilitating effect on activation (Haslam, 1973) at least by inhibition of the cAMP-dependent PKA-mediated phosphorylation of vasodilator-stimulated phosphoprotein (VASP). VASP is an actin regulatory protein and a negative modulator of α IIB β 3 integrin activation. Thus, levels of VASP phosphorylation/dephosphorylation reflect the P2Y₁₂ inhibition/activation state, which might constitute a sensitive marker to identify patients insufficiently protected by clopidogrel treatment (Schwarz et al., 1999; Aleil et al., 2004). Second, P2Y₁₂ stimulates PI3-K activity, which is important to sustain aggregation (Trumel et al., 1999; Kauffenstein et al., 2001). In addition, P2Y₁₂ is known to activate the small GTPase Rap1b through a PI3-K dependent mechanism (Lova et al., 2002, 2003; Woulfe et al., 2002; Larson et al., 2003). These multiple pathways explain why P2Y₁₂ plays such a central role in hemostasis and thrombosis (Conley and Delaney, 2003). Thus, overall, the P2Y₁₂ receptor appears to be responsible for most of the cofactor role of ADP in amplification of platelet activation induced by low concentrations of agonists such as TXA₂, thrombin, collagen, chemokines, or immune complexes.

The P2Y₁ and P2Y₁₂ receptors are differentially involved in the procoagulant activity of platelets. Both receptors are indirectly involved through platelet P-selectin exposure and formation of platelet-leukocyte conjugates, leading to leukocyte tissue factor exposure, whereas the P2Y₁₂ receptor is also directly involved in the procoagulant activity of platelets through phosphatidylserine exposure at the surface of platelets (Léon et al., 2001, 2003). An interesting feature of these receptors is their regulation after activation. Once activated by a first application of ADP, platelets become unresponsive to a second stimulation with ADP. This so-called refrac-

tory state of platelets to ADP has been recently shown to be caused by desensitization of the P2Y₁ receptor with a resultant loss of shape change and aggregation (Baurand et al., 2000), whereas the P2Y₁₂ receptor remains functional. Further studies established that the P2Y₁ and P2Y₁₂ receptors are differentially regulated and relocated upon activation, the P2Y₁ receptor after clathrin-dependent internalization, whereas the P2Y₁₂ receptor mainly stays at the plasma membrane (Baurand et al., 2005). This action may be of major consequence *in vivo*, since even in platelets refractory to stimulation by ADP, the P2Y₁₂ receptor would be able to ensure platelet reactivity at sites of injury, thus preventing loss of the hemostatic function. In summary, both P2Y₁ and P2Y₁₂ receptors are necessary for normal hemostasis and both play a key role in arterial thrombosis. P2Y₁₂ receptors targeting antithrombotic drugs already exist, and new compounds have been developed and are under clinical evaluation.

For P2Y₁, studies with knockout mice as well as pharmacological inhibition with antagonists such as MRS2179 and more recently MRS2500 clearly demonstrated its relevance as a possible target for new antiplatelet drugs (Gachet and Hechler, 2005). P2Y₁₂ receptor-deficient mice have been generated. They are viable, fertile, and display no spontaneous bleeding or other overt abnormalities (Foster et al., 2001; André et al., 2003). They display prolonged bleeding time compared with wild-type as well as P2Y₁ receptor-deficient mice. Platelet aggregation in response to ADP is inhibited, but shape change is conserved. Finally, these mice are insensitive to clopidogrel treatment (Foster et al., 2001), confirming that the P2Y₁₂ receptor is the target of this drug. These mice have been used in several models of arterial thrombosis and were shown to be resistant to thrombosis. The role of the P2Y₁₂ receptor in the stabilization of the thrombus was confirmed in these mice. Whether other phenotypic modification could be observed has not yet been reported.

VIII. Source of Naturally Occurring Ligands and Mechanisms of Transport and Breakdown

ATP is released into the extracellular space as an autocrine/paracrine molecule in response to neuronal stimulation, platelet aggregation, stress, mechanical stimulation and a number of receptor agonists. As previously indicated, known P2 receptors are stimulated by ATP, ADP, UTP, UDP, or UDP-glucose, and concentrations of these nucleotides in the extracellular space ranging from 0.1 to 10 μ M are required for activity.

The dynamic regulation of extracellular nucleotides manifested by the balance between release, catabolism, and interconversion in the extracellular space has imposed a layer of complexity to the understanding and quantification of the pharmacologically active nucleotides. Additional confounding factors such as cell lysis or

tissue damage contribute to the complexity by releasing large quantities of intracellular nucleotides. In addition, most of the measurements of extracellular levels of ATP have been performed in the bulk phase of the extracellular environment of the cells, and, therefore, these measurements largely underestimate the true concentration of extracellular nucleotides at the receptor level in the pericellular environment of the cells. Attempts to overcome some of these difficulties have been addressed by the development of sensitive assays capable of estimating subnanomolar concentrations of ATP and UTP (Levitt et al., 1984; Chen et al., 1994; Yang et al., 1994; Taylor et al., 1998; Beigi et al., 1999; Lazarowski and Harden, 1999; Huang et al., 2001; Lazarowski et al., 2003a) and by the development of specific biosensors such as the attachment of the luciferase enzyme to the surface of the cell (Beigi et al., 1999; Joseph et al., 2003), and by the use of nucleotide receptors to "sense" the pericellular nucleotide concentrations (Homolya et al., 2000; Sorensen and Novak, 2001; Coco et al., 2003; Hayashi et al., 2004). These techniques have allowed the assessment of nucleotide concentrations at the pericellular level and have demonstrated that the measurements of nucleotides in the bulk phase could underestimate the values at the receptor level by at least 20-fold (Joseph et al., 2003).

Millimolar concentrations of ATP exist in the cytoplasm of all cell types. However, because of the net negative charge of the molecule, ATP and other nucleotides do not permeate the cell membrane and therefore, require specialized transport mechanisms to reach the extracellular space and to act on the target receptors. Most of the available information on extracellular nucleotides refers to ATP and the products of its catabolism; however, the other agonist of P2Y receptors, UTP (and probably other nucleotides as well), is also released to the extracellular space by mechanisms similar to those for ATP. Several forms of nucleotide release have been proposed to date, as summarized below.

A. Basal Unstimulated Nucleotide Release

1. Constitutive Release of ATP. Extracellular nucleotides have been detected in multiple cell types under unstimulated conditions (Lazarowski et al., 2000, 2001a; Ostrom et al., 2000; Joseph et al., 2003). Stress or mechanical stimulation of many cell types results in a large elevation in the concentration of extracellular nucleotides, followed by a rapid decrease to a constant level in resting conditions that ranges between 0.5 and 10 nM in most cells. Lazarowski et al. (2000, 2001a) and Donaldson et al. (2000) demonstrated that resting levels of ATP and UTP are maintained by a steady-state equilibrium between release and metabolism. This observation suggests that a constitutive unstimulated release of ATP and UTP exists in many cells. Consistent with this observation, the resting levels of ATP increased gradually by inhibition of the metabolism of ATP with decreasing

the temperature of primary cultures of endothelial cells (Schwiebert et al., 2002) or by treatment of BAC1.2F5, 1321N1, PC12, and C6 cells with the ecto-ATPase inhibitor α,β -meATP (Beigi and Dubyak, 2000; Joseph et al., 2004).

Now that the tools have been developed to assess low levels of extracellular nucleotides including UTP, UDP, and UDP-glucose, along with sensitive "biosensors" and methodology to monitor the levels of ATP in the pericellular space, the physiological relevance of the resting levels of extracellular nucleotides can be explored. For example, Ostrom et al. (2000) demonstrated in MDCK cells that basal levels of ATP acting on P2Y receptors and thus promoting the formation of prostaglandins is the key determinant of the "set point" of the cAMP signal transducing pathway, illustrating the relevance of constitutive release of ATP on the autocrine/paracrine regulation of other signal transduction mechanisms (Ostrom et al., 2000).

Another example of the physiological relevance of constitutive release of extracellular ATP is observed in the airway epithelium. This steady state modulates some of the important functions of CFTR via activation of A_{2B} receptors, including regulation of the airway surface liquid volume to afford efficient cilia movement and optimal viscoelastic properties of the mucus layer to maintain efficient transport and clearance of the airways (Lazarowski et al., 2004).

2. Constitutive Release of UDP-Glucose. The identification of UDP-glucose as the endogenous ligand of an orphan GPCR (Chambers et al., 2000; Freeman et al., 2001) that has been incorporated into the family of P2Y receptors on the basis of the sequence homology and structural motifs (Abbracchio et al., 2003) prompted the development of a sensitive and selective assay for UDP-glucose (Lazarowski et al., 2003b). This assay was subsequently used for the assessment of extracellular levels of the receptor agonist and to explore the potential mechanisms of release of UDP-glucose into the extracellular space. Mechanical stimulation of 1321N1 astrocytoma cells resulted in an increase of extracellular levels of ATP and a modest increase of UDP-glucose. In marked contrast with ATP, the levels of UDP-glucose remained elevated for an extended period of time. Under resting conditions at steady state, the levels of UDP-glucose are severalfold higher than those of ATP or UDP. UDP-glucose is released in primary cultures of human airway epithelial cells, Calu-3, COS-7, CHO-K1, and C6 glioma cells with resting levels similar to or higher than those of ATP. These studies suggest that UDP-glucose, is released constitutively in a manner similar to ATP, and it is metabolized at a slower rate than ATP in all cells studied.

UDP-glucose participates in the quality control of protein synthesis in the lumen of the endoplasmic reticulum where it is actively transported by an UDP-glucose/UMP antiporter. The authors speculate that UDP-

glucose transported to the endoplasmic reticulum lumen could be released from cells as cargo as part of the constitutive pathway that delivers glycoproteins to the plasma membrane (Lazarowski et al., 2003b, 2004).

B. ATP Release by Excitable and Secretory Tissues

Exocytosis of ATP-containing vesicles occurs in neurons, platelets, adrenal medullary chromaffin cells, neuroendocrine cells, and mast cells. Neuronal release of nucleotides was first identified 50 years ago by Holton and Holton (1954) and is now one of the best-understood sources of extracellular nucleotides (Bodin and Burnstock, 2001a). ATP and probably other nucleotides are stored at high concentrations in dense granules in platelets, synaptic vesicles in neurons, and chromaffin granules in adrenal glands. The concentration of nucleotides in these vesicles could reach up to 150 to 1000 mM. Stimulation of these cells results in the calcium-dependent fusion of these vesicles with the plasma membrane and the release of vesicular contents into the extracellular space. The specific mechanisms of exocytotic granule release have been extensively investigated (Sperlágh and Vizi, 1996).

C. ATP Release by Nonexcitatory Cells

Extracellular nucleotides are released by multiple nonexcitatory cell types such as endothelial and epithelial cells, smooth muscle cells, fibroblasts, astrocytes, red blood cells, lymphocytes, monocytes, and a number of transformed cell lines. The release of nucleotides by nonexcitatory cells is not completely understood and involves the participation of multiple mechanisms. No evidence for the existence of specialized ATP-containing vesicles in most of these cells has been demonstrated, and, therefore, alternative mechanisms for the release of nucleotides have been postulated. These mechanisms are outlined below in the context of 1) stress/hypoxia/mechanical stimulation, 2) vesicular trafficking, and 3) agonist-promoted stimulation.

1. Stress/Hypoxia/Mechanical Stimulation. Evidence for non-neuronal, nonlytic release of ATP has been documented from a large number of tissues under a variety of physiological and experimental conditions. One of the first reports of the physiological non-neuronal release of ATP dates from 1969 by Forrester and Lind, who demonstrated the presence of ATP in the venous effluent of a forearm subjected to a sustained contraction.

Release of ATP and UTP induced by mechanical stimulation has also been documented in vitro in cell lines expressing P2Y receptors. Normal cell culture manipulation and/or medium changes resulted in marked P2Y receptor-dependent accumulation of inositol phosphates in cells expressing the recombinant P2Y₁, P2Y₂, and P2Y₄ receptors, but not in nontransfected cells (Filtz et al., 1994; Parr et al., 1994; Lazarowski et al., 1997b). The effect of medium changes and cell manipulation

were eliminated by treatment of cells with apyrase. Because no evidence of cell lysis was observed in these and other studies, these observations indicate that mechanical stimulation results in the nonlytic release of sufficient concentrations of ATP and UTP to effectively trigger the signaling of nucleotide receptors.

One of the most relevant responses to physiological mechanical stimulation occurs in epithelial cells. The bladder is subjected to marked changes in tension and pressure during normal filling and voiding. Burnstock (1999) proposed that the distension of the bladder and other visceral tissues such as ureter, vagina, and gallbladder leads to the release of ATP in the vicinity of the epithelium, where sensory nerve endings containing P2X receptors (P2X₃ or P2X_{2/3}) are stimulated to convey nociceptive stimuli to the central nervous system (Bodin and Burnstock, 2001a).

Several studies have demonstrated that perfusion of endothelial cells at high shear force leads to the release in a nonlytic fashion of significant amounts of nucleotides to the extracellular space, and hypoxia potentiates the release of nucleotides induced by shear force (Bodin and Burnstock, 1995). Hypoxia induces ATP release from pulmonary artery adventitial fibroblasts and endothelial cells (Gerasimovskaya et al., 2002), and the resulting autocrine activation of P2 receptors appears to be involved in the proliferation of fibroblasts under hypoxic conditions. Under hypoxia, erythrocytes release ATP that induces vasodilatation in skeletal muscle and pulmonary artery (Sprague et al., 2001; Gonzalez-Alonso et al., 2002).

Extracellular nucleotides play an important role in the regulation of the function of airway epithelial cells. Activation of P2Y₂ receptors located in the apical membrane results in calcium-activated chloride channel activity important in the regulation of fluid secretion, mucociliary clearance, and the regulation of the volume of the periciliary liquid, enabling efficient cilia movement and mucus layer transport (Schwiebert and Kishore, 2001; Leipziger, 2003).

The lack of nerve endings in airway epithelial cells, led to the suggestion that the epithelial cell itself is the main source of extracellular nucleotides. This topic has been extensively studied during the past few years and evidence suggests that ATP and UTP release from epithelial cells occurs preferentially at the luminal face (Lazarowski et al., 2004).

Several mechanisms for the release of ATP in response to nonlytic stimuli have been contemplated: transport by ATP binding cassette (ABC) proteins, stretch-activated channels, and voltage-activated channels and release via vesicular trafficking (Bodin and Burnstock, 2001b; Knight et al., 2002; Lazarowski et al., 2003a).

a. ATP binding cassette proteins. The CFTR and the multidrug resistance (MDR) gene product, P-glycoprotein, have been suggested to participate in the transport

of ATP into the extracellular space (Abraham et al., 1993; Reisin et al., 1994; Schwiebert et al., 1995). However, further studies did not confirm that CFTR functions as an ATP transport protein (Li et al., 1996; Reddy et al., 1996; Grygorczyk and Hanrahan, 1997; Watt et al., 1998). An alternative hypothesis to explain the conflicting results has been proposed, suggesting that CFTR and the MDR1 gene product participate in the release of ATP by regulating the activity of alternative nucleotide transporters associated with the ABC protein (Jiang et al., 1998; Sugita et al., 1998; Hazama et al., 2000; Braunstein et al., 2001; Roman et al., 2001). Evidence against a direct role of CFTR on ATP release has been demonstrated in ocular ciliary epithelial cells (Mitchell et al., 1998).

b. Stretch and voltage-activated Cl^- channels. Cell volume regulation represents a unique opportunity to study a physiological mechanism of nucleotide release by epithelial cells. ATP release stimulated by treatment with hypo-osmotic solutions has been demonstrated in a number of cell types including epithelial and endothelial cells, suggesting that a mechanosensitive ATP channel tightly associated with CFTR is implicated in the efflux of ATP. These conclusions were based on the pharmacological effects of gadolinium (Gd^{3+}), a nonselective inhibitor of stretch-activated channels (Braunstein et al., 2001). Further studies by Boudreault and Grygorczyk (2002) caution that these observations could be produced by interference of Gd^{3+} with the luciferin-luciferase bioluminescence assay used in these studies. Consistent with this conclusion, Knight et al. (2002), failed to demonstrate an effect of Gd^{3+} (or glibenclamide) on ATP release during nonlytic distension of the ureter. In contrast, these authors found that monensin and brefeldin A inhibited the distension-evoked ATP release, suggesting that nucleotide efflux from the ureter epithelium occurs by vesicular exocytosis (Knight et al., 2002).

Another potential mechanism for the regulation of cell volume is by the electrogenic translocation of ATP via an anion channel independent of CFTR. Recent studies have provided evidence for an ATP-conducting maxi-anion channel involved in volume regulation (Sabirov et al., 2001; Dutta et al., 2002; Bell et al., 2003b). This channel has been named the volume (and voltage)-dependent ATP conductive large conductance anion channel (see Sabirov and Okada, 2004). This channel has a large unitary conductance and a permeability ratio (P_{ATP}/P_{Cl}) of 0.14. This permeability, although apparently low, is sufficient to produce physiologically significant ATP efflux in the presence of high intracellular Cl^- concentrations. Further studies by these investigators indicated that this channel fits the pharmacological profile of swelling-induced ATP release, namely, sensitivity to Gd^{3+} , 4-acetamido-4'-isothiocyanostilbene-2,2'-disulfonate, and 5-nitro-2-(3-phenylpropylamino)benzoic acid, and insensitivity to phloretin, niflumic acid, and glibenclamide (Sabirov and Okada, 2004).

Activation of this ATP-conducting maxi-anion channel has been demonstrated in macula densa cells in response to an increase in tubular NaCl concentration. Under these conditions, ATP is released at relevant physiological concentrations from the basolateral membrane of macula densa cells in close proximity to the mesangial cells that express $P2Y_2$ receptors (Bell et al., 2003a).

A cell-to-cell paracrine signal transduction mediated by ATP and this anion channel plays a central role in stress sensory transduction for cell volume regulation and tubuloglomerular feedback in the kidney. The molecular entity of this channel has not been established; however, a candidate molecule with similar electrophysiological properties is the mitochondrial voltage-dependent anion channel (VDAC-1) porin.

Recently, Okada et al. (2004) reported that a splice variant of mitochondrial VDAC-1 is expressed in airway cells and demonstrated that VDAC-1 contributes to the release of ATP in murine cells in response to hypo-osmotic challenge. However, the authors also observed that cells from VDAC-1 knockout mice exhibit significant release of ATP, suggesting that additional ATP transport molecules play a role in volume regulation in murine cells (Okada et al., 2004).

2. Nucleotide Release via Vesicular Trafficking. An alternative mechanism for the release of nucleotides into the extracellular space via vesicular trafficking has been proposed (Burnstock, 1999; Bodin and Burnstock, 2001b). Distension of the guinea pig ureter by increasing the intraluminal pressure resulted in the release of ATP in a pressure-dependent manner. Treatment with different drugs added to the perfusion medium was used to determine the possible mechanism of ATP release. Monensin, an inhibitor of vesicular formation from the Golgi apparatus, and brefeldin A, a drug that disrupts the vesicular traffic between the endoplasmic reticulum and the Golgi apparatus, inhibited the distension-evoked ATP release by the urothelium. Drugs that inhibit ABC proteins or nucleoside and nucleotide transporters had no effect on the release of ATP, suggesting that the release of ATP from the ureter epithelium is mediated by vesicular exocytosis (Knight et al., 2002).

3. Agonist-Promoted ATP Release. Activation of signaling pathways by some receptor agonists can result in the specific release of ATP at sufficient concentrations to elicit a physiological response in the same cell or in a neighboring cell. Receptor-mediated ATP release has been observed in aortic endothelial and smooth muscle cells treated with thrombin (Pearson and Gordon, 1979). Bradykinin, acetylcholine (ACh), and serotonin also promote release of ATP by endothelial cells and bradykinin and phenylephrine promote release from MDCK cells (Yang et al., 1994; Ostrom et al., 2000).

Increases of extracellular levels of ATP have been reported in response to treatment with ADP in coronary endothelial cells (Buxton et al., 2001) and with UTP in

COS-7 and HEK-293 cells (Ostrom et al., 2000); however, in both cases the appearance of ATP could result from NDPK-dependent phosphorylation of added ADP in the first case and from phosphorylation of endogenous ADP in the second case. The intrinsic mechanisms for the agonist-mediated ATP release have not been elucidated and are probably related to the activation of the specific signaling pathways promoted by these agonists. In MDCK, COS-7, and HEK293 cells, the Ca^{2+} mobilizing hormones, bradykinin and phenylephrine, but not the cAMP-elevating forskolin and prostaglandin E_1 promoted the release of ATP, suggesting that a Ca^{2+} -dependent event downstream of G_q and PLC was involved in the release of ATP (Ostrom et al., 2000). Joseph et al. (2003) recently demonstrated that the release of ATP by thrombin was only partially dependent on Ca^{2+} . This observation, together with the evidence for the participation of PI3-K (Feranchak et al., 1998, 1999) and Rho or Rho kinase in the release of ATP (Koyama et al., 2001) suggest that, in the case of thrombin, ATP release could be mediated by G_{12} -dependent activation of Rho. Because Rho and PI3-K are involved in the regulation of vesicle traffic and cytoskeleton organization, an interpretation of the effects of thrombin described above is that the mechanism of ATP release could involve vesicle transport (Lazarowski et al., 2003a).

Pathogenic bacteria (*Shigella* and *Escherichia coli*) induce the release of ATP from gut epithelial cells (Crane et al., 2002; Tran Van Nhieue et al., 2003). The released ATP correlates with bacterial invasion and spreading, but also promotes secretory diarrhea. The release of ATP by bacteria is partially due to cell lysis but also involves specific mechanisms of recognition, e.g., the ligation of host glycolipid asialo-GM1 by the flagellin of *Pseudomonas aeruginosa* flagellum (McNamara et al., 2001).

D. ATP Release by Tissue Damage

Cell lysis or tissue damage results in the release of the intracellular nucleotide contents into the extracellular space providing nucleotide concentrations sufficient to activate nucleotide receptors. This source of extracellular nucleotides accounts for only some of the known physiological actions of ATP, including transmission of nociceptive signals and onset of inflammation.

E. Extracellular Nucleotide Metabolism

The physiological effect of extracellularly released nucleotides depends mainly on the particular receptor subtype(s) expressed in the tissue. In addition, an important modulatory effect of the physiological response is imposed by the presence of soluble and membrane-bound ecto-nucleotidases that rapidly metabolize the nucleotides with the potential to terminate or initiate receptor-specific signaling cascades.

Extracellular ecto-nucleotidases are expressed in most if not in all cell types. Several families of ectoen-

zymes participate in the degradation and interconversion of extracellular nucleotides. The most prominent enzyme classes are the ecto-nucleoside triphosphate diphosphohydrolase (E-NTPDase) family, the ecto-nucleotide pyrophosphatase/phosphodiesterase (E-NPP) family, ecto-5'-nucleotidase, alkaline phosphatase, the nucleotide-converting enzyme ecto-NDPK, and adenylate kinase. Ecto-nucleotidases exist as membrane-associated forms or soluble forms. Some of the soluble ecto-nucleotidases are produced by the proteolytic cleavage of the extracellular domains of membrane-bound forms or by the release from a glycosylphosphatidylinositol-anchoring phospholipid. The characterization of the ecto-nucleotidases has uncovered significant overlapping substrate specificities and tissue expression (Zimmermann, 2000; Goding et al., 2003).

1. *Ecto-Nucleoside Triphosphate Diphosphohydrolases.* E-NTPDases previously known as ecto-apyrases, NTPases, or E-ATPases can hydrolyze both nucleoside triphosphates (NTPs) and nucleoside diphosphates (NDPs). Eight members of this family (E-NTPDases 1–8) have been described. E-NTPDases 1, 2, 3, and 8 are cell surface-associated, whereas E-NTPDases 4, 5, 6, and 7 are intracellular enzymes. E-NTPDases 1, 2, 3, 4, 7, and 8 have N- and C-terminal transmembrane domains, whereas E-NTPDases 5 and 6 have only a single N-terminal transmembrane domain that can be cleaved, releasing the extracellular domain as soluble protein.

E-NTPDase 1, previously known as apyrase or CD39, hydrolyzes both NTPs and NDPs with a 1:1 selectivity ratio. E-NTPase 2 also known as ecto-ATPase or CD39L1, selectively hydrolyzes NTPs over NDPs with a selectivity ratio of 1:0.03 and E-NTPDase 3, known as ecto-ATPDase, CD39L3, and HB6, has an intermediate selectivity ratio of 1:0.3. Recently, a new member of this family, E-NTPDase 8, has been described. This enzyme hydrolyzes adenine and uridine nucleotides with low apparent K_m values (Bigonnesse et al., 2004). All members of this family share five highly conserved sequence domains, known as apyrase conserved domains that are essential for the catalytic activity (Handa and Guidotti, 1996; Vasconcelos et al., 1996; Zimmermann, 2000).

2. *Ecto-Nucleotide Phosphophosphates/Phosphodiesterases.* The E-NPP family comprises five members. E-NPP1–3 are transmembrane metalloenzymes with a structure consisting of a short N-terminal intracellular domain, a single transmembrane domain, and a large extracellular domain containing the catalytic active site. Soluble forms of these enzymes have been found and are presumably originated by proteolytic cleavage of the extracellular domains of the membrane-bound forms. E-NPP1–3 are classified as alkaline ecto-nucleotide pyrophosphatase/phosphodiesterase I and catalyze the hydrolysis of pyrophosphate and phosphodiester bonds. Therefore, these enzymes hydrolyze ATP into AMP and PP_i ; 3',5'-cAMP into AMP; ADP into AMP and P_i and

NAD⁺ into AMP and nicotinamide mononucleotide. E-NPP isoenzymes can hydrolyze purine and pyrimidine nucleotides, as well as dinucleotides and UDP-sugars (see below). Genes encoding putative E-NPP4 and 5 have been identified in searches of databases. The gene products of these enzymes have not been characterized (Goding et al., 2003).

Different investigators have independently characterized members of the E-NPP family, and several names have been given to the same protein. E-NPP1 is also known as PC-1 (Van Driel et al., 1985; Van Driel and Goding, 1987), E-NPP2 is also known as PD-I α and autotaxin, and E-NPP3 is known as PD-I β , B10, and gp130^{RB13-6} (Van Driel et al., 1985; Van Driel and Goding, 1987; Stracke et al., 1992; Narita et al., 1994; Deissler et al., 1995; Jin-Hua et al., 1997; Scott et al., 1997). PD-I α and autotaxin are splice variants of the same gene (Murata et al., 1994; Kawagoe et al., 1995), whereas PD-I β , B10, and gp130^{RB13-6} correspond to the same protein.

In addition to their role in the metabolism of extracellular nucleotides, E-NPP isoenzymes have other, probably related, physiological roles including nucleotide recycling, regulation of pyrophosphate levels, bone mineralization, and cell motility (see Goding et al., 2003). Recently, the substrate selectivity for E-NPP2 has been expanded with the observation that this enzyme, but not E-NPP1 or E-NPP3, hydrolyzes lysophosphatidylcholine into lysophosphatidic acid, linking the activity of the enzyme with tumor cell growth and motility (Tokumura et al., 2002; Umezu-Goto et al., 2002).

3. Hydrolysis of UDP-Glucose. UDP-glucose, the natural agonist of the P2Y₁₄ receptor, and other nucleotide sugars are also substrates of E-NPP isoenzymes. Lazarowski et al. (2003b) demonstrated that cultures of 1321N1 human astrocytoma cells hydrolyzed UDP-glucose into UMP and glucose 1-phosphate. Similar findings were observed in COS-7, Calu-3, and C6 cells (E. Lazarowski, personal communication). Whether UDP-glucose is hydrolyzed by an E-NPP isoenzyme or a novel enzymatic activity requires further investigation.

4. Hydrolysis of Diadenosine Polyphosphates. Diadenosine polyphosphates via stimulation of P2Y (and P2X) receptors regulate a large variety of physiological functions, such as synaptic transmission (Miras-Portugal et al., 1999, 2003), platelet aggregation (Zamecnik et al., 1992), vascular tone (Ralevic et al., 2001), and intraocular pressure (Pintor et al., 2004). Diadenosine polyphosphates are stored and released into the extracellular space from chromaffin cells, neurons, and platelets among other cell types. Upon release, diadenosine polyphosphates are rapidly hydrolyzed, presumably by members of the E-NPP family. Studies of the hydrolysis of dinucleoside polyphosphates conducted with adrenal, endothelial, airway epithelial, blood cell, and synaptic membranes (Lüthje and Ogilvie, 1988; Ramos and Rotllan, 1995; Mateo et al., 1997a,b; Picher and Boucher,

2000), demonstrated an asymmetric hydrolysis of the dinucleoside polyphosphates into AMP and Ap_{n-1}. Identical degradation products from several dinucleoside polyphosphates were found in a recent study by Vollmayer et al. (2003) using E-NPP1, E-NPP2, and E-NPP3, suggesting that these enzymes are the major candidates for the hydrolysis of dinucleotides.

5. 5'-Nucleotidase. Ecto-5'-nucleotidase is a glycosylphosphatidylinositol-anchored enzyme also known as CD73. This enzyme converts nucleoside monophosphates into the corresponding nucleoside and inorganic phosphate. 5'-Nucleotidase is the major enzyme involved in the formation of adenosine in the extracellular space (Zimmermann, 1992, 1996b). A soluble form of 5'-nucleotidase released from the glycosylphosphatidylinositol anchor has been described.

6. Nucleoside Diphosphokinase Extracellular nucleotides are also substrates of enzymes that participate in the interconversion and preservation of high-energy nucleotides. Extracellular NDPK activity has been described in 1321N1 astrocytoma and C6 glioma cells, as well as in airway epithelial cells. This enzyme catalyzes the reversible transphosphorylation of nucleoside diphosphates into nucleoside triphosphates such as ATP + UDP \rightleftharpoons ADP + UTP (Lazarowski et al., 1997a, 2000; Grobber et al., 1999; Yegutkin et al., 2000, 2001, 2002).

The presence of this transphosphorylation activity in a tissue or cell line could be higher than that of the ecto-nucleotidase activity, as in the case of 1321N1 or C6 cells, and this condition could lead to erroneous interpretation of the effects of administration of exogenous nucleotides to these cells. For example, the addition of UTP in a system with high levels of endogenous ADP and NDPK could result in the formation of sufficient ATP to elicit a pharmacological response that could be misinterpreted as being produced by UTP. The molecular nature of this enzymatic activity and its tissue distribution remain to be elucidated.

7. Alkaline Phosphatase. This enzymatic activity alone can catalyze the complete sequential hydrolysis of a nucleoside triphosphate to the corresponding nucleoside. Alkaline phosphatase can also hydrolyze PP_i (Zimmermann, 1996a, 2000). The presence of this enzyme was described recently in the mucosal surface of human airway epithelium (Picher et al., 2004). Alkaline phosphatase activity and mRNA expression were elevated 3- to 6-fold above normal in the airway epithelium of patients with chronic airway diseases such as primary ciliary dyskinesia, cystic fibrosis, and α_1 -antitrypsin deficiency (Picher et al., 2004).

8. Adenylate Kinase. Adenylate kinase catalyzes the conversion of two ADP molecules into ATP + AMP in a reversible manner. In contrast with alkaline phosphatase and NDPK activities, the substrate selectivity for adenylate kinase is restricted only to adenine nucleotides. This transphosphorylation activity plays a role in the metabolism of extracellular nucleotides by counter-

acting the stepwise nucleotide breakdown by E-NTP-Dases, E-NPPs, and 5'-nucleotide. Ecto-adenylate kinase activity has been documented in endothelial cells (Yegutkin et al., 2001, 2002), human airway epithelial cells (Donaldson et al., 2002; Picher and Boucher, 2003), and rat brain synaptosomes (Nagy et al., 1989).

The mechanisms of nucleotide release, hydrolysis, and interconversion are not fully understood. The recognition of the molecular basis of nucleotide release and the catalytic mechanism of hydrolysis is essential for the understanding of nucleotide receptor signaling and its role under normal physiological and pathological conditions.

A better understanding of the regulation of these processes will prove useful in the broadening of potential therapeutic approaches to modulate extracellular nucleotide signaling and their multiple physiological consequences.

IX. Interactions between P2Y and Other Receptors

A. Modes of Interaction between G Protein-Coupled Receptors

Individual GPCRs were long considered to preferentially activate specific intracellular signaling pathways and so act in a linear fashion to produce a change in cellular activity. However, it is now clear that they can also modulate the signals initiated by another GPCR to potentiate or inhibit the activity of these receptors. Such interactions can take place at the level of the GPCR themselves, through the formation of oligomers, or downstream of the receptor through the action of second messengers. The former process is commonly referred to as receptor dimerization and is characterized by the appearance of a physical receptor complex as demonstrated by coimmunoprecipitation or other methods, which displays novel pharmacological properties and/or interactions with second messenger systems. The latter process, known as receptor cross-talk, can have either positive or negative effects and serves to integrate coincident signals from multiple types of receptors, which are not physically associated.

B. Receptor Dimerization

Until recently, individual GPCRs were assumed to exist and function as monomers. However, in the mid-1990s it started to become clear that many GPCRs can coalesce into functional dimers or oligomers. Indeed, such aggregation may be the rule rather than the exception and may be essential for the correct trafficking and membrane expression of GPCRs (Bouvier, 2001; Milligan, 2004). At present, it is difficult to differentiate experimentally between dimers and oligomers and so the term dimer tends to be used. Within a receptor family, individual subtypes can form homodimers or different subtypes can form heterodimers. An increasing

number of combinations are also being described for receptors from totally different families. As GPCRs are already major therapeutic targets, dimerization has important implications for the development of new drugs.

There is also evidence that the human P2Y₂ receptor forms homodimers. Expression in cell lines of constructs in which the receptor was tagged at the C terminus with the cyan or yellow variants of green fluorescent protein produced functional receptors that were activated by UTP and which coupled to endogenous G proteins to induce release of Ca²⁺ stores (Kotevic et al., 2005). When both variants were coexpressed in the same cells, fluorescence resonance energy transfer (FRET) imaging indicated that they were in close proximity, i.e., they formed constitutive receptor complexes. It was notable that UTP did not change the FRET signal, indicating that binding of an agonist does not induce dissociation of the oligomeric complex. Further studies are required to determine whether other P2Y subtypes similarly form homodimers.

To date, the interaction between the rat P2Y₁ receptor and adenosine A₁ receptor is the only example of dimerization involving P2Y receptors with non-P2Y receptors that has been characterized extensively. Yoshioka et al. (2001) coexpressed the rat P2Y₁ and adenosine A₁ receptors in HEK293 cells. Initial experiments showed that these receptors coimmunoprecipitated in Western blots of whole cell membrane lysates, indicating that they formed a heteromeric complex. Coexpressing the P2Y₁ receptor did not alter surface expression of the A₁ receptor, but it did inhibit the binding of radiolabeled A₁ agonists and antagonists in membrane preparations. This change was not seen in a mixture of membranes from cells expressing each receptor individually. Additionally, the binding of an A₁ agonist was displaced by the P2Y₁ agonist ADPβS and P2Y₁ antagonist MRS2179 in cotransfected cells, but not in cells expressing the A₁ receptor only. These data again indicate formation of a heteromeric complex.

A₁ receptors couple to G_i and so mediate depression of intracellular cAMP levels, whereas P2Y₁ receptors interact with G_{q/11} and have no effect on cAMP. ADPβS inhibited cAMP production in cotransfected cells only, an effect that was antagonized by the A₁ antagonist 8-cyclopentyl-1,3-dipropylxanthine, but not by MRS2179, and abolished by pertussis toxin. Thus, ADPβS appears to have acted via the A₁ receptor ligand-binding site, i.e., the P2Y₁/A₁ dimer has novel pharmacological properties compared with the parent receptors. Interestingly, although ADPβS induced inositol phosphate synthesis, the A₁ agonist cyclopentyl adenosine (CPA) did not. Thus, dimerization did not lead to a complete change in pharmacological properties in this case.

Using confocal laser microscopy to study the subcellular distribution of the P2Y₁ and A₁ receptors, Yoshioka et al. (2001) showed that both were expressed mainly

near the plasma membrane of HEK293 cells. Furthermore, there was a strong overlap in their distribution in individual cells. This was confirmed in a subsequent study using the biophysical technique of bioluminescence resonance energy transfer (Yoshioka et al., 2002b). In the absence of agonists, the receptors showed a homogeneous colocalization across the cells. Addition of ADP β S and CPA together, but not alone, induced an increase in the bioluminescence resonance energy transfer ratio over 10 min. Thus, although the receptors have a constitutive association, their coactivation increased the association. This association was also seen with native receptors in central neurons. Using confocal laser microscopy and double immunofluorescence, Yoshioka et al. (2002a) demonstrated that the P2Y₁ and A₁ receptors colocalized in neurons of the rat cortex, hippocampus, and cerebellum. A direct association was then shown by their coimmunoprecipitation in membrane extracts from these regions.

Together, these studies clearly indicate that the rat P2Y₁ and A₁ receptors physically interact to form a functional dimer with novel pharmacological properties. The structural requirements for this interaction are not known at present. The physiological roles of the P2Y₁/A₁ dimer also remain to be determined, although Nakata et al. (2003) have pointed out that its pharmacological properties resemble those of a presynaptic receptor that mediates inhibition of neurotransmitter release in some tissues. Finally, it is still not known whether other P2Y subtypes also form functional heterodimers, but this is likely to be the case as Yoshioka et al. (2001) reported that the rat P2Y₂ receptor also coimmunoprecipitated with the A₁ receptor when they were coexpressed in HEK293 cells. Thus, the formation of oligomers by P2Y receptors is likely to be widespread and to greatly increase the diversity of purinergic signaling.

Indeed, homodimers of the human P2Y₁ receptor have recently been shown to exist [Choi et al., 2005a (<http://www.pa2online.org/abstracts/Vol3Issue2abst010P.pdf>)]. Expressed receptors were labeled in culture, and FRET confocal microscopy was applied. This method can be used to report specifically on the receptors located at the cell membrane and measure the percentage dimerized. Constitutive P2Y₁ homodimers were found; dimerization was increased by exposure to agonist to 80 to 90%. An obligate determinant of the dimer formation resides in the tail of the sequence. P2Y₁ receptor agonist-induced internalization is preceded by the dimerization [Choi et al., 2005b (<http://www.pa2online.org/abstracts/Vol3Issue2abst121P.pdf>)].

C. Receptor Cross-Talk

The ability of GPCRs to modulate the signals initiated by other GPCRs, receptor tyrosine kinases and ligand-gated ion channels is widespread and has been studied extensively (Selbie and Hill, 1998; Hur and Kim, 2002). A variety of mechanisms can underlie these interac-

tions, and here we will discuss a few examples of P2Y receptor-mediated cross-talk with these three different types of receptors.

1. G Protein-Coupled Receptors. A common form of cross-talk is for two types of receptors to produce a greater than additive, or synergistic, change in the level of second messengers. Thus, coactivation of a variety of G_i-coupled receptors and G_{q/11}-coupled P2Y receptors induces a synergistic rise in intracellular IP₃ and Ca²⁺ levels (Gerwins and Fredholm, 1992; Megson et al., 1995; Selbie et al., 1997; Werry et al., 2002) and release of AA (Felder et al., 1991; Selbie et al., 1997). For example, the DDT₁ MF-2 smooth muscle cell line expresses native A₁ and P2Y₂ or P2Y₄ receptors and at concentrations that had little or no effect on IP₃ levels, the A₁ agonist CPA potentiated the response to ATP in a concentration-dependent manner (Gerwins and Fredholm, 1992). Similar effects were seen when the downstream effect of IP₃, a rise in intracellular [Ca²⁺], was measured. Low concentrations of CPA and ATP or UTP applied individually had little effect, but when administered together the increase in [Ca²⁺] was nearly maximal. A subsequent study demonstrated that PKC translocation from the cytoplasm to the plasma membrane was also potentiated in these cells (Fredholm et al., 2003). Recombinant human CXCR2 chemokine receptors expressed in HEK293 cells and the native P2Y₁ and P2Y₂ receptors also display cross-talk (Werry et al., 2002). The CXCR2 receptor couples to G_i and so its agonist IL-8 has no effect on intracellular Ca²⁺ levels when applied alone. However, after prestimulation of cells with ATP or UTP, IL-8 induced a substantial [Ca²⁺] elevation. Prestimulation of P2Y₁ and P2Y₂ receptors also induced the appearance of a Ca²⁺ response to subsequent stimulation of the endogenous, G_s-coupled β -adrenoceptor. Activation of G_{q/11}-coupled P2Y receptors also leads to synthesis of DAG and stimulation of PKC, which can induce the release of AA. This arm of the G_{q/11}-activated signaling cascade can also cross-talk with G_i-coupled receptors. Expression and activation of recombinant m₂, α_2 , and D₂ receptors in CHO-K1 cells did not induce AA release per se, but each augmented the response mediated by ATP acting at an endogenous P2Y receptor (Felder et al., 1991).

The cross-talk between the P2Y₂ and CXCR2 receptors involved an enhancement of PLC activity (Werry et al., 2003) and that between neuropeptide Y1 and P2Y receptors in CHO-K1 cells was inhibited by $\beta\gamma$ subunit scavengers (Selbie et al., 1997). Thus, the most likely possibility is that G protein $\beta\gamma$ subunits generated by G_i-coupled receptors interact with G protein α subunits generated by the P2Y agonists at the level of PLC β , thus increasing the production of IP₃ and DAG, with consequent increases in Ca²⁺ release and PKC activation.

P2Y receptors have also been reported to inhibit effects evoked by non-P2Y GPCR. Sphingosine-1-phosphate acts via sphingosine-1-phosphate receptors in re-

nal mesangial cells to activate the MAP kinase pathway and induce mitogenesis. Activation of endogenous P2Y receptors by ATP or UTP inhibits these actions (Xin et al., 2004). The nucleotide-mediated inhibition was reversed by inhibitors of protein kinase C. In rat C6 glioma, β -adrenoceptors mediate a decrease in the activity of protein kinase B and induce cellular differentiation (Van Kolen and Slegers, 2004). Costimulation of the native P2Y₁₂ receptors reversed the inhibition of protein kinase B to activation, abolished the differentiation, and induced cell proliferation. These effects were G_i- and phosphatidylinositol 3-kinase-dependent. Thus, P2Y receptors may be important modulators of cell growth and differentiation induced by other GPCR agonists.

Recently, two types of cross talk between P2Y receptors and receptors for cysteinyl-leukotrienes (CysLT) have been described. The first is a positive action, as found in human mast cells, where UDP at nanomolar levels activated the CysLT₁ and CysLT₂ receptors to stimulate Ca²⁺ mobilization and cytokine production (Mellor et al., 2001, 2002, 2003). ATP and UTP were ineffective, and the UDP preference suggests an interaction with the P2Y₆ receptor. The nature and mechanism of this interaction are at present not determined (Kanaoka and Boyce, 2004). The second type is a negative cross-talk between P2Y and CysLT₁ receptors both in human monocyte/macrophage-like U937 cells (which constitutively express P2Y₂ and P2Y₆ receptors), and in COS-7 cells transiently expressing the CysLT₁ receptor (Capra et al., 2005). In U937 cells, activation of P2Y receptors with ATP or UDP induced CysLT₁ receptor heterologous desensitization. Conversely, activation of the CysLT₁ receptor by cysteinyl-leukotrienes had no effect on P2Y receptor responses, suggesting that the latter have a hierarchy in producing desensitizing signals. Although cysteinyl-leukotriene-induced homologous desensitization of the CysLT₁ receptor was followed by receptor internalization, ATP/UDP-induced CysLT₁ receptor desensitization was unable to cause receptor internalization. Moreover, at variance with homologous desensitization, P2Y receptor-induced regulation of the CysLT₁ receptor was dependent upon PKC, suggesting that P2Y receptors may mediate PKC-dependent phosphorylation of the CysLT₁ receptor. This study demonstrates, for the first time, that CysLT₁ receptor desensitization and trafficking are differentially regulated by its cognate ligand or by extracellular nucleotides. Moreover, because both cysteinyl-leukotrienes and nucleotides accumulate at sites of inflammation, this receptor cross-talk may represent a mechanism to fine tune the inflammatory response, i.e., a feedback mechanism by which extracellular nucleotides protect cells from the vast increase of inflammatory mediators characteristic of several pathological processes. The high degree of functional interaction between P2Y and CysLT receptors is also confirmed by the demonstration that montelukast and pranlukast, two well-known selective CysLT₁

receptor antagonists (Brink et al., 2003), functionally interact with P2Y receptor signaling pathways by inhibiting nucleotide-induced calcium mobilization in a relatively nonsubtype-specific manner (Mamedova et al., 2005).

2. Receptor Tyrosine Kinases. P2Y receptors can modulate the activity of receptor tyrosine kinases. For example, LPS acts through a member of this family to activate I κ B kinase, which controls the activity of the transcription factor NF- κ B. In murine J774 macrophages, UTP, acting via P2Y₆ receptors, slightly induced I κ B kinase activation and greatly potentiated the effect of LPS (Chen and Lin, 2001). The release of intracellular Ca²⁺ stores and activation of calmodulin-dependent kinase by UTP played a major role in this synergistic interaction. P2Y receptors can also act to inhibit receptor tyrosine kinase signaling. The growth hormone PDGF acts via a receptor tyrosine kinase to increase human vascular smooth muscle cell proliferation, as measured by [³H]thymidine incorporation into DNA and an increase in cell number (White et al., 2000). UTP and UDP had no effect on [³H]thymidine incorporation on their own, but significantly reduced the response to PDGF. Similarly, the nucleotides had no effect on cell number per se, but the increase induced by PDGF over 7 days was abolished. Interestingly, ATP potentiated the [³H]thymidine incorporation elicited by PDGF. The mechanism underlying these excitatory and inhibitory effects of P2Y receptors is not known, but the authors demonstrated clearly that changes in Ca²⁺ levels or p42/p44 MAPK activity were not involved. The fact that different nucleotide agonists had opposing effects also indicates that the roles of P2Y receptors in cell proliferation can be complex. P2Y₆ receptors have also been reported to protect against apoptosis induced by the receptor tyrosine kinase agonist tumor necrosis factor α in 1321N1 astrocytoma cells (Kim et al., 2003b). Interestingly, activation of the P2Y₄ receptor was ineffective, again indicating that the interactions of P2Y receptors with receptor tyrosine kinases can be complex.

3. Ligand-Gated Cation Channels. P2X receptors are ligand-gated cation channels and are often expressed in the same cells as P2Y receptors. Thus, there is a great scope for bidirectional cross talk between these two families of nucleotide-sensitive receptors. When ATP is applied to *Xenopus* oocytes expressing the recombinant human P2X₁ receptor, it induces a transient inward current (Vial et al., 2004). Repeated application at 5-min intervals caused rundown of this response, which was reversed by coexpression and coadministration of the P2Y₁ receptor, plus ADP, or the P2Y₂ receptor, plus UTP. This effect was mediated by activation of a protein kinase but did not involve direct phosphorylation of the P2X₁ receptor. Rather, an accessory protein appeared to be the target.

Cross-talk can also take place in the opposite direction, e.g., P2X₁ receptors have been shown to modulate

P2Y receptor activity. P2X₁ receptors are Ca²⁺ permeable and in megakaryocytes, the P2X₁ agonist α,β -meATP induced a rapid and transient Ca²⁺ influx (Vial et al., 2002). In contrast, ADP, acting via P2Y₁ receptors, evoked a slower, larger, and more maintained Ca²⁺ influx. The response seen when the two agonists were coapplied was markedly accelerated and the peak amplitude was potentiated, suggesting that the P2X₁ receptor may have a priming role in activation of P2Y₁ receptors during platelet stimulation.

P2Y receptors also interact with non-nucleotide ligand-gated cation channels. The TRPV1 receptor is the target for capsaicin and is coexpressed with P2Y₁ receptors in a population of nociceptive sensory nerves (Kennedy et al., 2003). ATP, 2-MeSATP, and ADP shifted the capsaicin concentration-response curve 2-fold to the left, with no change in maximum response, and decreased the threshold for activation by heat from 42 to 35°C (Tominaga et al., 2001). The potentiation was inhibited by the PKC inhibitor calphostin and mimicked and then occluded by the PKC activator PMA. Thus, phosphorylation of TRPV1 receptors by P2Y₁ receptor-activated PKC can modulate the sensitivity of sensory neurons to noxious stimuli.

X. Gene Activation Regulated by P2Y Receptors

A. Scope of the Gene Activations

GPCRs possess the important capability of maintaining signal transduction processes over extended periods where needed, allowing some transductions to proceed to the level of nuclear gene regulation and protein synthesis. Nearly all the cases for which this is known involve G_s-linked receptors, through their activation of the cAMP response element-binding transcription factor (CREB). Only one G_s-linked P2Y receptor is known: P2Y₁₁ (see section VI.E.); its responses in dendritic cells of the immune system (Wilkin et al., 2001; Marteau et al., 2004, 2005) suggest that it is likely there to promote transcription through the CREB pathway, since the activation of P2Y₁₁ led to production of specific cytokines via a cAMP increase. Furthermore, another study (van der Weyden et al., 2000a) made in renin-expressing cells has shown that activation of the P2Y₁₁ receptor therein increased renin mRNA and protein 3-fold over 36 h and gave strong luciferase-reporter activity when linked to a renin gene construct. This response was established to be through the CREB pathway, since the P2Y₁₁ activation stimulated CREB phosphorylation and its effects were blocked by mutation at the cAMP response element within the renin gene promoter.

There is a small amount of definitive information on gene transcription control by identified G_q- or G_{i/o}-linked P2Y receptors. One such case is represented by the P2Y₂ receptor. When stably expressed in 1321N1 cells, this receptor was found to signal through the p38 MAPK cascade to phosphorylate CREB, which then mediated

cis-activation of target genes, including the antiapoptotic *bcl-2* and *bcl-xl* genes (Chorna et al., 2004). UTP incubation also up-regulated expression of a range of genes for neurotrophins and neuropeptides and induced proliferation of the astrocytoma cells. It is interesting that P2Y₁ behaves differently from P2Y₂ in the same cells. In a study performed with the P2Y₁ receptor therein (Sellers et al., 2001), agonist treatment led, via a different intracellular cascade, to the activation instead of the transcription factor Elk-1. The outcome of that transduction route was apoptosis and inhibition of cell proliferation. In another study performed on rat astrocytes, activation of a native PTX-sensitive P2Y-like receptor increased the binding to DNA of the activator protein-1 and NF- κ B transcription factors (Brambilla et al., 2003). The outcome in that case was the de novo synthesis of cyclooxygenase-2 (confirmed at the protein level), whose gene transcription has been shown to occur through these factors (Brambilla et al., 1999, 2000, 2002, 2003), followed by elongation of astrocytic processes, an index of reactive astrogliosis.

B. Synaptically Released ATP Can Act in the Control of Gene Transcription

At many synapses, ATP is stored and released with another transmitter throughout life; hence, it has the potential to serve as an additional nerve-released trophic factor, which would imply its regulation of gene transcription. This possibility has recently been explored using initially the neuromuscular junction (NMJ) of skeletal muscles; this type of synapse has the greatest accessibility and uniformity in its population and is currently the best understood of all types in structure, operation, and synaptogenesis. ATP is costored in vesicles of the nerve terminals with ACh and is coreleased quantally with it (at a ratio of approximately 1 ATP to 6 ACh) (Silinsky and Redman, 1996). Also, UTP is present in those vesicles at ~10% of the ATP content (Zimmermann, 1994).

Functional postsynaptic P2Y₁ and P2Y₂ receptors colocalized at the NMJs with the nicotinic ACh receptors (AChRs) have been demonstrated in mammalian, chicken, and amphibian muscles (Choi et al., 2001; Tsim et al., 2003; Tung et al., 2004). There is earlier phenotypic evidence that muscle AChR and acetylcholinesterase (AChE) have some controls of their expression and clustering at the NMJ in common (Sanes and Lichtman, 1999); hence, the genes of both those effectors have been studied (Choi et al., 2001, 2003), using differentiated myotubes when the P2Y receptors are present throughout the cell surface before being clustered at nerve contact when in vivo. Exposure to 2-MeSADP or to UTP each produces an activation of the genes of the multiple subunits of the AChR and also of the AChE catalytic subunit gene. Properties of the gene activations mediated by the native P2Y₁ and P2Y₂ receptors in the muscle cells (Choi et al., 2001; Tung et al., 2004) are 1) a

strong increase (up to 550%) in each mRNA (confirmed at the protein level where tested), related to the concentration of the P2Y agonists; 2) a dose-dependent agonist-induced increase in the activity of a promoter-luciferase reporter construct for each of the subunit genes; 3) approximate equivalence for the P2Y₁ and P2Y₂ receptor activities in stimulating the promoter, for each gene (one example is shown in Fig. 8C); 4) a total block of the P2Y₁-coupled action at the gene promoters by the specific P2Y₁ antagonist MRS2179 (Fig. 8C) or by suramin for both receptors; and 5) a great increase in the 2-MeSADP-activated promoter activities of the AChR genes when the P2Y₁ receptor protein content was boosted by a transfection of its cDNA.

Examination of several potential transcription factors for this action has identified Elk-1 as a candidate, which becomes maximally phosphorylated and nuclear-localized when myotubes are exposed for 30 min to 2-MeSADP or UTP. Four potential DNA binding sequences for Elk-1 were identified in the first intron of the AChE gene (Fig. 8A). Two were proven to be active by gel mobility shift assays, and each of these sites was used in the reporter assay, replacing the entire 2.2-kb DNA of the promoter-containing region by a 20-nucleotide Elk-1 binding site. Each again showed AChE gene promoter activity dependent on P2Y₁ or P2Y₂ receptor activation on the myotube (Fig. 8B). This was enhanced when the two sites were combined. In confirmation of their activity in the native gene, when the entire 2.2-kb promoter-containing DNA was mutated at six bases in those two sites, its P2Y₁/P2Y₂-mediated promoter activity was abolished.

The significance of the location at the synapse of two P2Y receptors that signal to the transcription of genes encoding effectors of the postsynaptic machinery should be considered in the context that a few subsynaptic nuclei at each junction become transcriptionally specialized during skeletal muscle development to sustain the local synthesis of the operational proteins of the junction, including the postsynaptic AChR and AChE (Duclet and Changeux, 1995). The P2Y₁ and P2Y₂ receptors at the synapse are thus part of a local circuit allowing efficient genetic control. ATP would not be the sole regulator of the expression of the AChR and AChE genes at the NMJ, since the neurally released trophic protein neuregulin is another major one, which also has its receptor, the ErbB protein, at the postsynaptic NMJ membrane (Fischbach and Rosen, 1997) and signals through a transduction cascade to activate a specific transcription factor in the nearby subsynaptic nuclei (Schaeffer et al., 2001). There is some evidence for an equivalent cascade route for the P2Y₁ receptor there (Choi et al., 2003), but this is not known fully yet, nor for the P2Y₂ receptor at this location. The primary action of neuregulin is in the development of the NMJ, although it later contributes to the maintenance of the NMJ (Buonanno and Fischbach, 2001). It is a protein secreted in

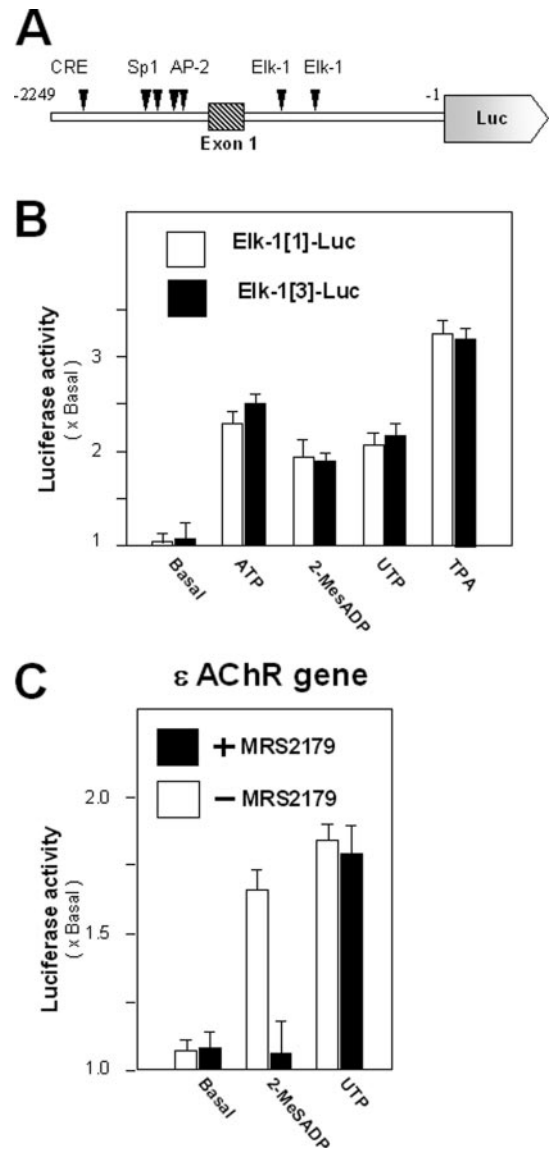


FIG. 8. Examples of analysis of gene activations controlled through P2Y receptor action. A, promoter region of the gene for the catalytic subunit of AChE. The first exon is noncoding, within the 5'-untranslated region. Promoter sites lie on either side of that, illustrated for four of the transcription factors active in muscle cells. Shown are two of the four predicted sites here for the transcription factor Elk-1 (Elk-1 [1] and Elk-1 [3]), which were identified as specific Elk-1 binding sites by gel mobility shift analyses on extracts of myotube nuclei. DNA sequences at those sites were inserted upstream in a luciferase reporter vector for assays as in B. B, two reporter constructs controlled by the specific binding sites for Elk-1 from the AChE gene promoter are used to show that Elk-1 is activated when the P2Y₁ or P2Y₂ receptors in a muscle cell are stimulated; hence, Elk-1 binds to activate transcription of the AChE gene there. Mouse myotubes transfected with either construct were incubated at 37°C with the agonists shown, or with none (Basal), or with the phorbol ester 12-O-tetradecanoylphorbol-13-acetate (TPA), which produces a nonspecific activation of AChE synthesis in muscle cells. Both Elk-1 sites are seen to have receptor-stimulated promoter activity (and when combined in tandem they are synergistic; not shown). Elk-1 is activated by specific phosphorylation through P2Y₁ (2-MeSADP) or P2Y₂ (UTP) action. The effect of ATP is significantly higher than effects for the other two (with all three at a saturating concentration), since it acts at both subtypes. C, similar analysis performed using the promoter region of the ε subunit of the nicotinic AChR gene. Both the P2Y₁ and P2Y₂ receptors act through Elk-1 in activating the multiple genes for the AChR in muscle cells (α, δ, and ε subunit genes have been tested). Note that in confirmation, the P2Y₁-specific antagonist MRS2179 blocks this action by 2-MeSADP but not that by UTP (data from Choi et al., 2003; Tung et al., 2004).

low amount, and it may be that the constant large quantity of ATP released is also needed for an additional role in the maintenance, specifically, of synaptic AChRs and AChE.

In conclusion, for some P2Y receptors a contribution to gene transcription control over a longer time frame may commonly be a role. Synapses elsewhere should be examined for a situation similar to that at the NMJ. In short-term phenotypic analyses, it may be difficult to recognize all of the functions of those receptors. Cases in which a P2Y gene knockout seems to have no obvious effect, other than where that is shown to be by compensation by another subtype, may be in this category.

XI. Potential Therapeutic Applications

There is increasing interest in the expression of P2Y receptors in pathological conditions and in the therapeutic potential of P2Y receptor-related compounds (see also reviews by Burnstock and Williams, 2000; Boeynaems et al., 2001, 2005a,b; Yerxa, 2001; Burnstock, 2002, 2006; Jacobson et al., 2002; Ralevic and Burnstock, 2003; Gachet, 2005).

Based on the firmly established role of P2Y₁ and P2Y₁₂ receptors in platelet aggregation (see above), one of the most interesting therapeutic applications is currently represented by the use of antagonists of these receptors as antithrombotic agents in the prevention of recurrent stroke and heart attacks. Thus, both P2Y₁ and P2Y₁₂ receptors constitute targets for antithrombotic therapy, and compounds with a dual action might also be of interest. However, the agents currently on the market (ticlopidine and clopidogrel) or known to be in development (prasugrel, cangrelor, and AZD6140) all target the P2Y₁₂ receptor. The thienopyridines (ticlopidine, clopidogrel, and prasugrel) inactivate the P2Y₁₂ receptor irreversibly via covalent binding of an active metabolite generated in the liver. The other compounds are competitive antagonists. Cangrelor (AR-C69931MX), an ATP derivative, is suitable for intravenous perfusion, whereas AZD6140 is in clinical development as an orally active agent (Boeynaems et al., 2005b). Human myocardium expresses a full panel of both P2Y and P2X receptors (Banfi et al., 2005), and it has also been suggested recently that antagonists to P2Y₂, P2Y₆, and P2Y₁₁ receptors could be beneficial in the treatment of hypertension and congestive heart failure (Balogh et al., 2005). P2Y₂ receptor up-regulation has been reported to contribute to neointimal hyperplasia and inflammation of rabbit carotid arteries through the increased proliferation of smooth muscle cells and the recruitment of monocytes via the endothelium-dependent up-regulation of vascular cell adhesion molecule-1 (Seye et al., 2002, 2003, 2004). These findings may open up further therapeutic possibilities in the cardiovascular field, in particular, in preventing and treating atherosclerosis.

P2Y receptors appear to be involved in insulin release from pancreatic β cells. Thus, they are a candidate target for the design of innovative antidiabetic drugs (Solini et al., 2003).

Purinergic signaling plays major roles in different activities in the gut. For example, T cells are thought to play a primary role in the induction of epithelial cell damage in IBD, and the P2Y₆ receptor was found to be highly expressed in the T cells infiltrating bowel involved in IBD but absent in T cells in the unaffected bowel. P2Y₆ receptors appear to be involved in monocytic release of inflammatory cytokines. P2Y receptors on smooth muscle and ATP production in myenteric plexus increase in postoperative ileus, probably contributing to delayed colonic transport (Wang et al., 2004b).

P2Y₂ receptors are present in both quiescent and activated hepatic stellate cells. Quiescent cells express P2Y₂ and P2Y₄ receptors whereas activated cells express P2Y₆ receptors (Dranoff et al., 2004).

P2Y₁, P2Y₂, and P2Y₁₁ receptors located on immune and inflammatory cells play a pivotal role in inflammation and immunomodulation (Schnurr et al., 2003; Luttikhuisen et al., 2004). Multiple P2Y receptors on osteoblasts, osteoclasts, and chondrocytes are being considered as potential targets for the development of therapeutics to inhibit bone resorption in diseases such as rheumatoid arthritis, osteoporosis, tumor-induced osteolysis, and periodontitis (Dixon and Sims, 2000).

There is a substantial presence of P2Y receptors in the kidney, in different regions of the nephron, in the glomerulus, and in the renal vascular system. P2Y receptors mediate renin secretion, and P2Y receptor-related compounds are being explored for the treatment of chronic renal failure and transplantation-induced erythrocytosis (Jackson, 2001). In polycystic kidney disease, there is an increase in expression of P2Y₂ and P2Y₆ receptors and P2Y antagonists and inhibitors of ATP release are being explored as therapeutic agents (Turner et al., 2004).

P2Y receptor antagonists have been proposed as potential neuroprotective agents in the brain, by modulation of α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid-induced currents and by excessive activation of glutamate receptor systems being implicated in cell death associated with stroke, epileptic seizures, and neurodegenerative diseases such as Alzheimer's disease, Parkinson's disease, and amyotrophic lateral sclerosis (Zona et al., 2000). Endogenous ATP has been claimed to be involved in the regulation of anxiety via stimulation of P2Y₁ receptors in the dorsomedial hypothalamus in rats (Kittner et al., 2003).

Multiple P2X and P2Y receptor subtypes are expressed by astrocytes, oligodendrocytes, and microglia (see Abbracchio and Verderio, 2006). It has also been claimed that P2Y₂ receptors activate neuroprotective mechanisms in astrocytic cells (Chorna et al., 2004). P2Y receptors mediate reactive astrogliosis via induction of

cyclooxygenase-2, and P2Y receptor antagonists might counteract excessive cyclooxygenase-2 activation in both acute and chronic neurological disease (Brambilla et al., 1999, 2002; Brambilla and Abbracchio, 2003). It has been reported that ATP continually modulates the cerebellar circuit by increasing the inhibitory input to Purkinje neurons probably via P2Y₂ and/or P2Y₄ (and possibly P2X₅) receptors, thus decreasing the main cerebellar output activity that contributes to locomotor coordination (Brockhaus et al., 2004).

The anticancer activity of adenine nucleotides has been known since 1983. Intraperitoneal injection of ATP into tumor-bearing mice results in significant anticancer activity against several fast-growing, aggressive carcinomas (Agteresch et al., 2003). The subtypes of P2 receptors involved are largely unresolved, but P2Y₁ and P2Y₂ receptors are considered as targets for tumor cell antiproliferation.

ATP is clearly involved in the initiation of pain in nociceptive sensory nerves, and although the main focus of attention has been P2X₃ receptors, there is increasing evidence for a role of P2Y receptors in the pain pathways. P2Y₁ and P2X₃ receptors are colocalized in a subpopulation of dorsal root sensory neurons (Ruan and Burnstock, 2003) and up-regulation of P2Y₁ receptor expression occurs after transection of sciatic nerves (Xiao et al., 2002). Analgesic effects of intrathecal administration of P2Y receptor agonists UTP and UDP in normal rat and neuropathic rat pain models have been reported, suggesting that P2Y₂ (and/or P2Y₄) and P2Y₆ receptors produce inhibitory effects on spinal pain transmission (Okada et al., 2002).

In airway epithelial cells, activation of P2Y₂ receptors stimulates the secretion of chloride via outwardly rectifying chloride channels and independently from the CFTR. This constitutes the rationale for the development of aerosolized P2Y₂ receptor agonists for improving clearance of secretions from the bronchi in the treatment of cystic fibrosis (Bennett et al., 1996), chronic bronchitis (Olivier et al., 1996), chronic obstructive pulmonary disease, and sputum expectoration in smokers (Yerxa, 2001). ATP and UTP stimulate P2Y₂ receptor-mediated surfactant secretion and transepithelial chloride secretion in type II alveolar cells, but there are abnormalities in this mechanism in cystic fibrosis. In view of the bronchoconstrictor action of adenosine, uracil nucleotides were preferred to adenine nucleotides, and dinucleotides were selected because of their longer half-life: first Up₄U (INS365) and later Up₄dC (INS37217) (Yerxa et al., 2002). Various clinical trials have been performed or planned in cystic fibrosis, chronic bronchitis, the common cold, and allergic rhinitis. However, in ophthalmology the first convincing results have been obtained. P2Y₂ receptors stimulate chloride and water secretion from conjunctival epithelial cells, leading to a better lubrication of the ocular surface. In a phase III clinical trial, Up₄U drops improved corneal lesions over

a 6- to 24-week period in patients suffering from dry eye syndrome (Nichols et al., 2004). A new drug application for Up₄U (diquafosol) in that indication has been submitted to the U.S. Food and Drug Administration.

Purinergic signaling is widespread in the eye, and novel therapeutic strategies are being developed for glaucoma, dry eye, and retinal detachment (Pintor et al., 2003). P2Y receptors on human corneal epithelial cells appear to play a critical role in the injury repair process (Klepeis et al., 2004). In the auditory system, ATP, acting via P2Y receptors, depresses sound-evoked compound action potentials in the auditory nerve (Sueta et al., 2003).

ATP potentially relaxes cavernosal smooth muscle via P2Y receptors that are present both on smooth muscle and endothelial cells, suggesting that P2Y receptor-mediated purinergic signaling may be involved in the pathophysiology of erectile dysfunction (Gür and Öztürk, 2000). ATP stimulates a biphasic change in trans-epithelial conductance in the human uterine cervix, phase I mediated by the P2Y₂ receptor and phase II by the P2X₄ receptor (Gorodeski, 2002), which has potential for therapeutic manipulation of fertility and contraception.

Tables 5 through 12 summarize the P2Y receptor subtypes discussed in this article.

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TABLE 5
P2Y₁ receptor (previously known as P2Y purinoceptor 1)

2.1:NUCT:1:P2Y₁.

	TM	aa	AC	Chr. Map	References
Human	7	373	NM_002563	3q25.2	Ayyanathan et al. (1996); Léon et al. (1996)
Rat	7	373	NM_012800	2q31	Tokuyama et al. (1995)
Mouse	7	373	NM_008772	3	Tokuyama et al. (1995)
Functional assays	Measure of calcium fluxes in Jurkat T lymphocytes expressed the cloned P2Y ₁ receptor (Léon et al., 1997); human blood platelets; rat and rabbit mesenteric artery; turkey platelets				
Ligand	Action	Selectivity	Endogenous	References	
2-MeSADP	Agonist	No	No	Ayyanathan et al. (1996); Palmer et al. (1998)	
ADP	Agonist	No	Yes	Ayyanathan et al. (1996); Palmer et al. (1998)	
ATP-γS	Agonist	No	No	Ayyanathan et al. (1996); Palmer et al. (1998)	
2-MeSATP	Agonist	No	No	Ayyanathan et al. (1996); Palmer et al. (1998)	
ADPβS	Agonist	No	No	Ayyanathan et al. (1996); Palmer et al. (1998)	
ATP*	Partial agonist /antagonist	No	Yes	Ayyanathan et al. (1996); Léon et al. (1997); Palmer et al. (1998)	
ATP-α-B	Agonist	No	No	Nahum et al. (2002)	
2-MeSATP-α-B	Agonist	No	No	Nahum et al. (2002)	
2ClATP-α-B	Agonist	No	No	Nahum et al. (2002)	
A3P5PS	Antagonist	No	No	Boyer et al. (1996a)	
A3P5P	Antagonist	No	No	Hechler et al. (1998b)	
A2P5P	Antagonist	No	No	Hechler et al. (1998b)	
MRS2179	Antagonist	Yes	No	Boyer et al. (1998); Camaioni et al. (1998); Baurand et al. (2001)	
MRS2279	Antagonist	Yes	No	Boyer et al. (1996a)	
PPADS	Antagonist	No	No	Lambrecht et al. (2002)	
Reactive blue 2	Antagonist	No	No	Lambrecht et al. (2002)	
Suramin	Antagonist	No	No	Lambrecht et al. (2002)	
Agonist potencies	<i>K_i</i> values for P2Y ₁ receptor agonists for inhibition of [³ H]MRS2279 binding to purified P2Y ₁ receptor: 2-MeSADP (<i>K_i</i> = 0.0099) > ADP (<i>K_i</i> = 0.92) > ATP-γS (<i>K_i</i> = 1.33) > 2-MeSATP (<i>K_i</i> = 1.87) > ADPβS (<i>K_i</i> = 2.42) > ATP (<i>K_i</i> = 17.7) (Waldo and Harden, 2004)				
Antagonist potencies	MRS2179 (p <i>K_B</i> = 6.75 (Moro et al., 1998) (selective); MRS2279 (p <i>K_B</i> = 8.10) (Boyer et al., 2002) (selective); reactive blue 2 (Lambrecht et al., 2002); PPADS (p <i>K_B</i> = 5) (Guo et al., 2002); suramin (p <i>K_B</i> = 5.5) (Ralevic and Burnstock, 1998); A3P5PS (p <i>A₂</i> = 6.5) (Boyer et al., 1996)				
Radioligand assays	Equilibrium binding of [³ H]MRS2279 to P2Y ₁ receptor expressed in membranes (Waldo et al., 2002; Waldo and Harden, 2004); binding studies with [³³ P]MRS2179 (Baurand et al., 2001)				
Radioligands	[³ H]MRS2279; [³³ P]MRS2179				
Transduction mechanism	G _q /G ₁₁ ; PI hydrolysis (PLC activation) and increasing of calcium in expression systems				
Distribution	Brain, placenta, prostate, heart, skeletal muscle, platelets, neuronal tissue; P2Y ₁ mRNA could not be detected in cartilage and bone; it was expressed at barely detectable levels in liver, kidney, stomach, lymphocytes, and bone marrow (Moore et al., 2001)				
Tissue function	Endothelium-dependent relaxation; smooth muscle relaxation (Ralevic and Burnstock, 1998); role in ADP-induced intracellular calcium mobilization, platelet shape change, aggregation, and TXA ₂ generation (Hechler et al., 1998a,b); mitogenic action in rat aorta smooth muscle (Ralevic and Burnstock, 1998); inhibition of N-type Ca ²⁺ channels (Filippov et al., 2000); role of the receptor in nucleotide-mediated calcium signaling in astrocytes (Fumagalli et al., 2003); possible role in the modulation of neuro-neural signaling transmission (von Kügelgen and Wetter, 2000); role in osteoclastic bone resorption (Hoebertz et al., 2002)				
Phenotypes	P2Y ₁ -null mice*				
Comments	In all species, the receptor is selective for adenine nucleotide;* these mice were viable with no apparent abnormalities that affected their development and survival; platelet count in these animals was identical to that of wild-type mice, but they were unable to aggregate in response to usual concentrations of ADP, whereas high concentrations of ADP induced platelet aggregation without shape change; P2Y ₁ -null mice had no spontaneous bleeding tendency but were resistant to thromboembolism induced by intravenous injection of ADP (Fabre et al., 1999; Léon et al., 1999a); molecular modeling study of the human P2Y ₁ receptor supports the idea that ATP binding to at least two distinct domains of the P2Y ₁ receptor, both outside and within the TM core; the two disulfide bridges present in the human P2Y ₁ receptor play a major role in the structure and stability of the receptor, to constrain the loops within the receptor, specifically stretching the extracellular loop 2 over the opening of the TM cleft and thus defining the path of access to the binding site (Moro et al., 2002); MRS2179 is the first P2Y ₁ receptor antagonist with antithrombotic action (Baurand and Gachet, 2003)				

aa, amino acids; AC, accession; chr., chromosome; MRS2179, 2'-deoxy-N⁶-methyladenosine-3',5'-bisphosphate; MRS2279, N⁶-methyl-(N)-methanocarba-2'-deoxyadenosine-3',5'-bisphosphate; A3P5PS, adenosine-3'-phosphate-5'-phosphosulfate; A3P5P, adenosine 3', 5'-diphosphate; A2P5P, adenosine 2', 5'-diphosphate; ATP-α-B, 5'-O-(1-boranotriphosphate).

TABLE 6
P2Y₂ receptor (previously known as P2U, P2U purinoceptor 1, and P2Y purinoceptor 2)

2.1:NUCT:2:P2Y₂.

	TM	aa	AC	Chr. Map	References
Human*	7	377	NM_176072; NM_002564; NM_176071	11q13.5–q14.1	Parr et al. (1994)
Rat	7	374	NM_017255	1q32	Chen et al. (1996)
Mouse	7	373	NM_008773	7	Lustig et al. (1993)
Functional assays	Rat coronary artery (Godecke et al., 1996); rat salivary glands (Turner et al 1997); human airway epithelium (Clarke et al., 1992); murine gallbladder epithelium (Clarke et al., 1999); current measurement in microinjected rat superior cervical sympathetic neurons (Brown, 2000); calcium imaging in HT29 and Colo320 DM colorectal carcinoma cells (Burnstock and Knight, 2004); HeLa cells (Muscella, 2003); A6 cells (Burnstock and Knight, 2004); HEK 293 cells (Schacter et al., 1996); studies of IP ₃ accumulation and intracellular calcium mobilization in 1321N1 astrocytoma cells (Parr et al., 1994; Lazarowski et al., 1995a); [³ H]inositol polyphosphate metabolism in rat aortic smooth muscle cells (Kumari, 2003); human coronary artery endothelial cells (Seye, 2003); intracellular calcium signaling, electrophysiology, and fluid transport in human fetal, rat, and rabbit retina (Meyer, 2002); Ca ²⁺ mobilization in HL-60 human neutrophil cell line (Meshki, 2004)				
Ligand	Action	Selectivity	Endogenous	References	
UTP	Agonist	No	Yes	Lazarowski et al. (1995b)	
UTP-γS	Agonist	No	No	Lazarowski et al. (1995b)	
ATP	Agonist	No	Yes	Lazarowski et al. (1995b)	
5BrUTP	Agonist	No	No	Lazarowski et al. (1995b)	
2ClATP	Agonist	No	No	Lazarowski et al. (1995b)	
dCp4U (INS37217)	Agonist	No	No	Kellerman et al. (2002)	
Up4U (INS365)	Agonist	No	No	Pendergast, et al. (2001)	
Agonist potencies	IP ₃ : UTP (EC ₅₀ 0.14 ± 0.02 μM) > ATP (EC ₅₀ 0.23 ± 0.01 μM) > Ap ₄ A (EC ₅₀ 0.72 ± 0.02 μM) > ATP-γS (EC ₅₀ 1.72 ± 0.15 μM) > 5BrUTP (EC ₅₀ 2.06 ± 0.04 μM); Ca ²⁺ : dCp4U (INS37217) (EC ₅₀ 0.22 μM); Up4U (INS365) (EC ₅₀ 0.1 μM) (Lazarowski et al., 1995; Pendergast et al., 2001; Yerxa et al., 2002)				
Antagonist potencies	Suramin; PPADS insensitive (Charlton et al., 1996)				
Radioligand assays	None				
Radioligands	None				
Transduction mechanism	G _i /G ₁₁ , PLCβ, and possibly G _i /G _o ; PI hydrolysis (PLCβ activation) and elevated Ca ²⁺ in expression systems; activation of ICl/Ca in <i>Xenopus</i> oocytes; transactivation of integrin and growth factor receptors (Soltoff, 1998; Erb et al., 2001; Seye et al., 2003; Liu et al., 2004)				
Distribution	Quantitative RT-PCR: skeletal muscle, heart, and some brain regions; at more moderate levels spleen, lymphocytes, macrophages, bone marrow, and lung; lowest levels of mRNA in liver, stomach, and pancreas (Moore et al., 2001); vascular smooth muscle and endothelial cells (Ralevic and Burnstock, 1998)				
Tissue function	Epithelial cell Cl ⁻ secretion (Clarke, 1992; Parr, 1994); pulmonary surfactant secretion (Rice, 1995); receptors on endothelial cells mediate NO release and subsequent vasodilatation (Ralevic and Burnstock, 1998); receptors on smooth muscle in some blood vessels mediate vasoconstriction (Ralevic and Burnstock, 1998); growth inhibition and apoptosis in human colorectal carcinoma cells (Burnstock and Knight, 2004); role in bone remodeling (Hoebertz et al., 2002); monocyte recruitment by vascular endothelium (Seye et al., 2003); proliferation of HeLa cells induced by c-Fos (Muscella, 2003); proliferation of lung cancer cells (Schafer, 2003); role in dry eye disease (Yerxa et al., 2002); regulation of proliferation in human keratinocyte (Burrel, 2003; Greig et al., 2003b); role in neutrophil degranulation (Meshki, 2004); growth of the arterial cell wall (Seye et al., 2002)				
Phenotypes	Reduction of chloride secretion in airway epithelia P2Y ₂ knockout mice (Cressman et al., 1999)				
Comments	The human P2Y ₂ receptor is activated by nucleotide triphosphates but not diphosphates (Nicholas et al., 1996) and is PPADS-insensitive (Charlton et al., 1996); P2Y ₂ receptor signaling is partially inhibited by PTX, indicating some involvement of G _i in receptor coupling (Parr et al., 1994)—in addition, native and recombinant forms of rat P2Y ₂ couple to PTX-sensitive G _γ o (Filippov et al., 1997; Mosbacher et al., 1998); G _o activation by P2Y ₂ receptors requires integrin receptor transactivation (Erb et al., 2001); rabbit P2Y ₂ couples to G _i /G _o isoforms (Murthy et al., 1998); *three transcript variants that encode the same protein have been identified for this gene (Parr et al., 1994); results of preclinical research suggest that P2Y ₂ receptor agonists inhibit Na absorption, restore Cl conductance, and rehydrate the cystic fibrosis airway surface (Kellermann, 2002); dCp4U (INS37217), a metabolically stable and potent P2Y ₂ agonist given for inhalation, has been evaluated in patients with cystic fibrosis lung disease (Kellerman et al., 2002); an RGD sequence in the P2Y ₂ receptor interacts with αVβ3 integrins (Erb et al., 2001); Src homology 3 binding domains in the P2Y ₂ receptor activate Src and transactivate growth factor receptors (Erb et al., 2001; Seye et al., 2004); the C-terminal domain of the P2Y ₂ receptor mediates receptor desensitization and internalization (Garrad et al., 1998)				

aa, amino acids; AC, accession; chr., chromosome.

TABLE 7
P2Y₄ receptor (previously known as P2Y purinoceptor 4 and uridine nucleotide receptor)

2.1:NUCT:3:P2Y₄

	TM	aa	AC	Chr. Map	References
Human	7	365	NM_002565	Xq13	Nguyen et al. (1995)
Rat	7	361	NM_031680	Xq31	Bogdanov et al. (1998b)
Mouse	7	361	NM_020621	X	Lazarowski et al. (2001)
Functional assays	Calcium mobilization in 1321N1 astrocytoma cells that express recombinant receptor (Communi et al., 1995; Nguyen et al., 1995)				
Ligand	Action	Selectivity	Endogenous	References	
UTP	Agonist	No	Yes	Communi et al. (1995)	
UTP γ S	Agonist	No	No	Jacobson et al. (2002)	
5Br-UTP	Agonist	No	No	Nguyen et al. (1995)	
UDP	Partial agonist	No	Yes	Nguyen et al. (1995)	
ITP	Partial agonist	No	No	Communi et al. (1995)	
ATP*	Agonist/antagonist	No	Yes	Herold et al. (2004)	
PPADS	Antagonist	No	No	Charlton et al. (1996)	
Reactive blue 2	Antagonist	No	No	Brown et al. (2002)	
Agonist potencies	Human: UTP (EC ₅₀ = 2.5 μ M) > ITP (33 μ M) (Communi et al., 1996a); rat: ITP (EC ₅₀ = 1.4 μ M) = ATP (1.8 μ M) = UTP (2.6 μ M) (Bogdanov et al., 1998b)				
Antagonist potencies	Human: PPADS (25% reduction at 100 μ M) (Charlton et al., 1996); ATP is a competitive antagonist (Kennedy et al., 2000); rat: reactive blue 2 (IC ₅₀ = 21 μ M) (Bogdanov et al., 1998b)				
Radioligand assays	None				
Radioligands	None				
Transduction mechanism	G _q /G ₁₁ and possibly G _i ; PI hydrolysis (PLC β activation) and elevated [Ca ²⁺] _i in expression systems; activation of ICl/Ca in <i>Xenopus</i> oocytes (by increased [Ca ²⁺] _i); inhibition of N-type Ca ²⁺ and M-type K ⁺ channels (Filippov et al., 2003)				
Distribution	Quantitative RT-PCR: high levels of mRNA in intestine, pituitary, and brain; low levels in liver and bone marrow (Moore et al., 2001); monocyte and lymphocytes (Jin et al., 1998b)				
Tissue Function	Endothelial cell receptors mediate NO release and subsequent vasodilatation (Burnstock et al., 2002); mitogenic actions on vascular smooth muscle cells (Burnstock et al., 2002); regulation of epithelial chloride transport in jejunum (Robaye, 2003)				
Phenotypes	Loss of nucleotide regulation of epithelial Cl ⁻ transport in the jejunum of null mice (Robaye et al., 2003)				
Comments	When expressed in a mammalian cell line, this receptor was activated specifically by UTP > UDP but not by ATP and ADP (Communi et al., 1995; Nguyen et al., 1995); competitive; a recent study shows that ATP is a potent antagonist at the hP2Y ₄ , whereas it is a full agonist at the rat P2Y ₄ (Herold et al 2004)*; rat P2Y ₄ is not selective for uridine nucleotides and instead shows an agonist potency order of ATP = UTP; ADP, ATP γ S, 2-MeSATP, and UDP are partial agonists (Bogdanov et al., 1998b; Webb et al., 1998); both human and rat P2Y ₄ are suramin-insensitive (Charlton et al., 1996); UTP regulates ion transport in jejunum (Cressman et al., 1999)—the disappearance of the jejunal Cl ⁻ secretory response to UTP and ATP in P2Y ₄ -null mice demonstrates the involvement of the P2Y ₄ receptor; P2Y ₄ receptors might be considered a potential pharmacotherapeutic target for cystic fibrosis (Robaye et al., 2003)				

aa, amino acids; AC, accession; chr., chromosome.

TABLE 8

*P2Y₆ receptor (previously known as P2Y purinoceptor 6, pyrimidinoceptor, and uridine nucleotide receptor)*2.1:NUCT:4:P2Y₆

	TM	aa	AC	Chr. Map	References
Human*	7	328	NM_176797; NM_176798; NM_176796; NM_004154	11q13.5 (Somers et al., 1997)	Communi et al. (1996b)
Rat	7	328	NM_057124	1Q32	Chang et al. (1995)
Mouse	7	328	AF298899	N.A.	Lazarowski et al. (2001b)
Functional assays	Formation of IP ₃ in 1321N1 cells that stably express recombinant receptor (Communi et al., 1996); rat C6–2B glioma cells that natively express the receptor (Nicholas et al., 1996)				
Ligand	Action	Selectivity	Endogenous	References	
UDP	Agonist	No	Yes	Communi et al. (1996b)	
UDPβS	Agonist	Yes	No	Jacobson et al. (2002)	
5BrUTP	Weak agonist	No	No	Communi et al. (1996b)	
UTP	Weak agonist	No	Yes	Communi et al. (1996b)	
ADP	Weak agonist	No	Yes	Communi et al. (1996b)	
2-MeSATP	Weak agonist	No	No	Communi et al. (1996b)	
PPADS	Antagonist	No	No	Robaye et al. (1997)	
Reactive blue 2	Antagonist	No	No	Robaye et al. (1997)	
MRS2567	Antagonist	Yes	No	Mamedova et al. (2004)	
MRS2578	Antagonist	Yes	No	Mamedova et al. (2004)	
MRS2575	Antagonist	Yes	No	Mamedova et al. (2004)	
Agonist potencies	IP ₃ formation: UDP (EC ₅₀ = 300 nM) > 5BrUTP (EC ₅₀ = 800 nM) > UTP (EC ₅₀ = 6 μM) > ADP (EC ₅₀ = 30 μM) ≫ 2-MeSATP (EC ₅₀ = 100 μM) (Communi et al., 1996b)				
Antagonist potencies	Reactive blue 2 > PPADS > suramin (Robaye et al., 1997)				
Radioligand assays	None				
Radioligands	None				
Transduction mechanism	G _q /G ₁₁ ; PI hydrolysis (PLCβ activation) and elevated [Ca ²⁺] _i in expression systems				
Distribution	Northern blot: placenta, spleen, thymus, intestine (Communi et al., 1996b); vascular smooth muscle, lung (Ralevic and Burnstock, 1998); RT-PCR: human bone, two osteoblastic cell lines, brain-derived cell lines (Burnstock and Knight, 2004); quantitative RT-PCR: spleen, placenta, kidney (high levels), lung, intestine, adipose, bone, heart (moderate expression), and some brain regions (Moore et al., 2001)				
Tissue function	Regulation of chemokine production and release in monocytes (Warny et al., 2001); NaCl secretion in colonic epithelial cells (Burnstock and Knight, 2004); role in proliferation of lung epithelial tumor cells (Schafer et al., 2003); role in mediating contractions of human cerebral arteries (Malmsjö et al., 2003b); interaction with the TNF-α-related signals to prevent apoptotic cell death (Kim et al., 2003b)				
Phenotypes	None				
Comments	Four transcript variants that encode the same isoform have been identified for this gene*; a pseudogene was also described (Burnstock and Knight, 2004); rat and mouse homologs have also been cloned (Chang et al., 1995; Lazarowski et al., 2001b); rat, mouse, and human P2Y ₆ receptors are UDP-selective, and UTP is less potent, whereas ATP is a very weak partial agonist; the P2Y ₆ receptor has a pharmacological profile very similar to the UDP receptor on rat C6–2B glioma cells (Lazarowski et al., 1994b); rat P2Y ₆ potently inhibited N-type Ca channels when expressed in sympathetic neurons, and receptor transcripts were reported to be present in brain tissues (Filippov et al., 1999)				

aa, amino acids; AC, accession; chr., chromosome.

TABLE 9
P2Y₁₁ receptor (previously known as P2Y)

2.1:NUCT:5:P2Y₁₁

	TM	aa	AC	Chr. Map	References
Human	7	374	NM_002566*	19p13.2	Communi et al. (1997)
Functional assays	cAMP measurement in CHO-K1 cells stably expressing the P2Y ₁₁ receptor; IP ₃ measurement in 1321N1 astrocytoma cells stably expressing the P2Y ₁₁ receptor (Communi et al., 1997)				
Ligand	Action	Selectivity	Endogenous	References	
ATP	Agonist	No	Yes	Communi et al. (1999b)	
ADP	Agonist	No	Yes	Communi et al. (1999b)	
ATP γ S	Agonist	No	No	Communi et al. (1999b)	
BzATP	Agonist	No	No	Communi et al. (1999b)	
dATP	Agonist	No	No	Communi et al. (1999b)	
ADP β S	Agonist	No	No	Communi et al. (1999b)	
2-MeSATP	Agonist	No	No	Communi et al. (1999b)	
AR-C67085	Agonist	No	No	Communi et al. (1999b)	
ADP γ S	Partial agonist	No	No	Communi et al. (1999b)	
AMP α S	Partial agonist	No	No	Communi et al. (1999b)	
A3P5PS	Partial agonist	No	No	Communi et al. (1999b)	
Suramin	Antagonist	No	No	Communi et al. (1999b)	
Reactive blue 2	Antagonist	No	No	Communi et al. (1999b)	
UTP	Agonist	No	Yes	White et al. (2003)	
Agonist potencies**	cAMP: ATP γ S (EC ₅₀ 3.4 \pm 0.3 μ M) > BzATP (EC ₅₀ 7.2 \pm 0.5 μ M) > dATP (EC ₅₀ 8.9 \pm 0.6 μ M) > ATP (EC ₅₀ 17.4 \pm 6.1 μ M) > ADP β S (EC ₅₀ 29.7 \pm 2.7 μ M) > 2-MeSATP (EC ₅₀ 50 \pm 4 μ M) (Communi et al., 1999b); IP ₃ : BzATP (EC ₅₀ 10.5 \pm 0.3 μ M) > ATP γ S (EC ₅₀ 13.5 \pm 2.7 μ M) > dATP (EC ₅₀ 16.3 \pm 0.7 μ M) > ATP (EC ₅₀ 65 \pm 12 μ M) > ADP β S (EC ₅₀ 174 \pm 28 μ M) > 2-MeSATP (EC ₅₀ 210 \pm 6 μ M) (Communi et al., 1999b)				
Antagonist potencies	PI hydrolysis: suramin (IC ₅₀ 1 μ M), reactive blue 2 (IC ₅₀ 9 μ M); cAMP: suramin (IC ₅₀ 16 μ M) (Communi et al., 1999b)				
Radioligand assays	None				
Radioligands	None				
Transduction mechanism	G _q /G ₁₁ and G _s (see "Comments"); PI hydrolysis (PLC β activation) and elevated [Ca ²⁺] _i in expression systems; adenylate cyclase stimulation and elevated cAMP levels (Qi et al., 2001a)				
Distribution	Brain, spleen, lymphocytes, intestine; all other tissues expressed more moderate levels of P2Y ₁₁ mRNA, with lowest levels detected in liver, cartilage, and bone (Moore et al., 2001)				
Tissue function	Role in maturation, migration of dendritic cells (Wilkin et al., 2001; Schnurr et al., 2003), and in granulocytic differentiation (Communi et al., 2000); secretory role in pancreatic duct epithelial cells (Nguyen et al., 2001)				
Phenotypes	None				
Comments	In P2Y ₁₁ gene, an intron was found (Communi et al., 1997)*; P2Y ₁₁ couples to G _q and G _s , and the pharmacological profile of the GPCR is slightly different for each transduction pathway (PLC β /IP ₃ and adenylate cyclase/cAMP) (Communi et al., 1999b; Qi et al., 2001a); the receptor does not activate PLA ₂ or PLD (Communi et al., 1999b); the P2Y ₁₂ antagonist ARC67085MX is an agonist here; a chimeric mRNA due to intergenic splicing between the P2Y ₁₁ and Ssf1 genes has been found in many mammalian cells—the function of this fusion protein is unknown (Communi et al., 2001); a canine (Qi et al., 2001a; Zambon et al., 2001) but no mouse and rat orthologs have been cloned; agonist potencies are species-dependent (Qi et al., 2001a; Zambon et al., 2001)**				

aa, amino acids; AC, accession; chr., chromosome.

TABLE 10
P2Y₁₂ receptor [previously known as *SP1999*, *P2T(AC)*, and *P2Y(AC)*]

2.1:NUCT:6:P2Y₁₂

	TM	aa	AC	Chr. Map	References
Human	7	342	NM_022788*; NM_176876	3q24–25	Hollopeter et al. (2001)
Rat	7	347	NM_022800	2q31	Hollopeter et al. (2001)
Mouse	7	343	NM_027571	2	
Functional assays	cAMP measurement in CHO cells and CHO cells stably transfected with hP2Y ₁₂ (Hollopeter et al., 2001; Zhang et al., 2001); FLIPR assay in CHO-dihydrofolate reductase-deficient transiently or stably transfected with hP2Y ₁₂ (Zhang et al., 2001); cAMP measurement in platelets (Cattaneo et al., 2003)				
Ligand	Action	Selectivity	Endogenous	References	
ATP**	Partial agonist	No	Yes	Waldo and Harden (2004)	
2-MeSADP	Agonist	No	No	Zhang et al. (2001)	
2-MeSATP	Agonist	No	No	Zhang et al. (2001)	
ADP	Agonist	No	Yes	Zhang et al. (2001)	
ADPβS	Agonist	No	No	Zhang et al. (2001)	
ATPγS	Agonist	No	No	Zhang et al. (2001)	
2ClATP	Agonist	No	No	Zhang et al. (2001)	
Reactive blue 2	Antagonist	No	No	Zhang et al. (2001)	
Suramin	Antagonist	No	No	Zhang et al. (2001)	
Ticlopidine	Antagonist	No	No	Savi and Herbert (2005)	
Clopidogrel	Antagonist	No	No	Savi and Herbert (2005)	
CS-747	Antagonist	No	No	Savi and Herbert (2005)	
AR-C66096MX	Antagonist	No	No	Gachet (2005)	
AR-M69931MX	Antagonist	No	No	Gachet (2005)	
AR-C67085MX	Antagonist	No	No	Gachet (2005)	
Agonist potencies	cAMP assays: 2-MeSATP (EC ₅₀ 3.4 ± 0.5 nM) = 2-MeSADP (EC ₅₀ 14.1 ± 2 nM) > ADP (EC ₅₀ 60.7 ± 10 nM) > ATPγS (EC ₅₀ 110 ± 100 nM) > ADPβS (EC ₅₀ 191 ± 20 nM) > 2ClATP (EC ₅₀ 636 ± 100 nM) (Zhang et al., 2001); FLIPR assay: 2-MeSADP (EC ₅₀ 2.3 ± 0.3 nM) = 2-MeSATP (EC ₅₀ 3.4 ± 0.3 nM) > ADP (EC ₅₀ 74 ± 10 nM) = ADPβS (EC ₅₀ 89 ± 15 nM) > 2ClATP (EC ₅₀ 470 ± 100 nM) > ATPγS (EC ₅₀ 1200 ± 200 nM) (Zhang et al., 2001)				
Antagonist potencies	Reactive blue 2 (K _i = 1.3 μM), suramin (K _i = 3.6 μM); AR-C66096, AR-C69931MX, ticlopidine, clopidogrel [prodrug that produces a metabolite that irreversibly binds to P2Y ₁₂ (Savi et al., 2000; Gachet et al., 2001, 2005)]; no effect of PPADS				
Radioligand assays	Binding of [³³ P]2-MeSADP (Cattaneo et al., 1997)				
Radioligands	[³³ P]2-MeSADP (Cattaneo et al., 1997)				
Transduction mechanism	Gα _i , inhibition of adenylate cyclase				
Distribution	Brain (in particular glial cells), spinal cord, platelets (Zhang et al., 2001; Sasaki et al., 2003)				
Tissue function	Role in dense granule secretion (Daniel et al., 1998), fibrinogen-receptor activation, and thrombus formation (Gachet, 2005); role in the recruitment of the platelets to the site of injury and in the enhancement of the efficiency of platelet activation by other agonists, such as thrombin and thromboxane A ₂ (Gachet, 2005); this receptor is defective in several patients with blood coagulation disorders (Cattaneo et al., 2000, 2003; Hollopeter et al., 2001)				
Phenotypes	P2Y ₁₂ -null mice***				
Comments	Two transcript variants that encode the same isoform have been identified for this gene*; ATP is a partial agonist**; when P2Y ₁₂ expression is low, as is seen in platelets, it acts as an antagonist, whereas at high expression levels, as seen with recombinant receptors, it is an agonist (Waldo and Harden, 2004); radiation hybrid analysis mapped the P2Y ₁₂ gene on 3q24–3q25 (Hollopeter et al., 2001), a region that also includes the P2Y ₁ , P2Y ₁₃ , and P2Y ₁₄ receptors; in a patient with congenital bleeding, Cattaneo et al. (2003) found compound heterozygosity for two mutations in the P2Y ₁₂ gene: neither mutation interfered with receptor surface expression but both altered function—in particular, these patients generally have normal platelet shape change responses to ADP but have impaired abilities to inhibit adenylate cyclase activity (Cattaneo et al., 1992); P2Y ₁₂ knockout mice seem normal, but they exhibit highly prolonged bleeding times, and their platelets aggregate poorly in responses to ADP and display a reduced sensitivity to thrombin and collagen (André et al., 2003)***; clinical studies using clopidogrel demonstrate a significantly reduced risk of peripheral artery disease, myocardial infarction, ischemic stroke, or vascular death compared with aspirin therapy; phase II studies of intravenous AR-C69931MX in patients with acute coronary syndromes show that this agent has a rapid onset of action, rapidly achieving steady-state inhibition of platelet aggregation with a half-life of only a few minutes (Gachet, 2005)				

aa, amino acids; AC, accession; chr., chromosome; FLIPR, fluorometric imaging plate reader.

TABLE 11
P2Y₁₃ receptor (previously known as GPR86, GPR94, and SP174)

2.1:NUCT:7:P2Y₁₃

	TM	aa	AC	Chr. Map	References
Human	7	333	NM_176894	3q24	Communi et al. (2001a)
Rat	7	336	AY639875	2	Fumagalli et al. (2004)
Mouse	7	337	NM_028808	2	Zhang et al. (2002a)
Functional Assays	Inositol phosphate measurement in 1321N1 astrocytoma cells that express recombinant receptor (Communi et al., 2001a); cAMP assays on stably transfected CHO-K1 cells (Communi et al., 2001a); binding of [³³ P]2-MeSADP in 1321N1 astrocytoma cells that express recombinant receptor (Marteau et al., 2003); assays of GTPγ[³⁵ S] binding to membranes of 1321N1 that express recombinant receptor (Marteau et al., 2003; Fumagalli et al., 2004)				
Ligand	Action	Selectivity	Endogenous	References	
2-MeSADP	Agonist	No	No	Zhang et al. (2002a); Marteau et al. (2003)	
ADPβS	Agonist	No	No	Zhang et al. (2002a); Marteau et al. (2003)	
2-MeSATP	Agonist	No	No	Zhang et al. (2002a); Marteau et al. (2003)	
ADP	Agonist	No	Yes	Zhang et al. (2002a); Marteau et al. (2003)	
Ap ₃ A	Agonist	No	No	Zhang et al. (2002a); Marteau et al. (2003)	
ATP	Agonist	No	Yes	Zhang et al. (2002a); Marteau et al. (2003)	
IDP	Agonist	No	No	Zhang et al. (2002a); Marteau et al. (2003)	
PPADS	Antagonist	No	No	Marteau et al. (2003)	
Suramin	Antagonist	No	No	Marteau et al. (2003)	
AR-C67085MX	Antagonist	No	No	Marteau et al. (2003)	
AR-C69931MX	Antagonist	No	No	Marteau et al. (2003)	
Agonist potencies	Fluorimetric imaging assay: 2-MeSADP (EC ₅₀ 19 ± 5 nM) = ADPβS (EC ₅₀ 31 ± 15 nM) = 2-MeSATP (EC ₅₀ 32 ± 8 nM) > ADP (EC ₅₀ 60 ± 26 nM) > Ap ₃ A (EC ₅₀ 72 ± 9 nM) > ATP (EC ₅₀ 261 ± 48 nM) > IDP (EC ₅₀ 552 ± 64 nM) (Zhang et al., 2002a)				
Antagonist potencies	AR-C69931MX > PPADS > suramin; MRS2179 inactive (Marteau et al., 2003)				
Radioligand assays	Binding of [³³ P]2-MeSADP (Marteau et al., 2003)				
Radioligands	[³³ P]2-MeSADP (Marteau et al., 2003)				
Transduction mechanism	G _i and G ₁₆ (Communi et al., 2001)				
Distribution	Spleen, brain regions, liver, pancreas, bone marrow, heart, peripheral leukocyte (Communi et al., 2001a; Zhang et al., 2002a)				
Tissue function	No precise evidence but presence in both immune system and brain implies roles				
Phenotypes	None				
Comments	This receptor clusters in the same region of the human chromosome 3q24–3q25, where some of the cloned P2Y receptors (P2Y ₁ , P2Y ₁₂ , and P2Y ₁₄) and several orphan GPCRs are also present (GPR87, GPR91, and H963) (Abbracchio et al., 2003); activated by diphosphate adenine nucleotides and not by uridine nucleotides; cloning and pharmacological characterization of the P2Y ₁₃ receptor from rat tissues has been recently described: by using the [³⁵ S]GTPγS binding assay in 1321N1 cells that express the rat receptor, ADP was found to be more potent than 2-MeSADP (Fumagalli et al., 2004)				

aa, amino acids; AC, accession; chr., chromosome.

TABLE 12
P2Y₁₄ receptor (previously known as GPR105, KIAA0001, and UDP-glucose receptor)

2.1:NUCT:8:P2Y₁₄

	TM	aa	AC	Chr. Map	References
Human	7	338	NM_014879	3q24–25	Chambers et al. (2000); Abbraccio et al. (2003)
Rat	7	338	NM_133577	2q31	Charlton et al. (1997)
Mouse	7	305	NM_133200	2	
Functional assays Intracellular calcium assays in HEK 293 cells that express recombinant receptor (Chambers et al., 2000)					
	Ligand	Action	Selectivity	Endogenous	References
	UDP-glucose	Agonist	Yes	Yes	Chambers et al. (2000)
	UDP-galactose	Agonist	Yes	Yes	Chambers et al. (2000)
	UDP-glucuronic acid	Agonist	Yes	No	Chambers et al. (2000)
	UDP- <i>N</i> -acetylglucosamine	Agonist	Yes	Yes	Chambers et al. (2000)
Agonist potencies	UDP-glucose (EC ₅₀ 80 ± 31 nM) > UDP-galactose (EC ₅₀ 124 ± 17 nM) > UDP-glucuronic acid (EC ₅₀ 370 ± 33 nM) > UDP- <i>N</i> -acetylglucosamine (EC ₅₀ 710 ± 27 nM) [stable HEK 293 cell line expressing both recombinant Gα ₁₆ and the human receptor (Chambers et al., 2000)]				
Antagonist potencies	No antagonists tested				
Radioligand assays	None				
Radioligands	None				
Transduction mechanism	G _i /G _o (Chambers et al., 2000)—in particular, data show a coupling to Gα subunits (Gα ₁₆ , Gα _{q05} , and Gα _{q15}) of the G _{i/o} family but not members of the G _s family or to endogenous G _{q/11} proteins in HEK 293 (Moore et al., 2003)				
Distribution	Placenta, adipose tissue, stomach, intestine, some brain regions, spleen, lung, heart, bone marrow, brain glia, and peripheral immune cells (Chambers et al., 2000; Moore et al., 2003)				
Tissue function	Role as chemoattractant receptor in bone marrow hematopoietic stem cells (Lee et al., 2003a); role in neuroimmune function (Moore et al., 2003); role in dendritic cells activation (Skelton et al., 2003)				
Phenotypes	None				
Comments	The P2Y ₁₄ receptor, from a phylogenetic and structural point of view, lies with the P2Y ₁₂ and P2Y ₁₃ receptors in the second main branch of the P2Y family—the gene for this receptor has been found in human chromosome 3q24–3q25, where a cluster of other related GPCRs, consisting of P2Y ₁ , P2Y ₁₂ , and P2Y ₁₃ receptors and of orphan GPR87, GPR91, and H963 have been found (Abbraccio et al., 2003); the complete sequences of the rat (VTR 15-20; Charlton et al., 1997; Freeman et al., 2001) and mouse orthologs (Freeman et al., 2001) of the human UDP-glucose receptor have also been described; rat P2Y ₁₄ , which is abundantly expressed in brain, has been reported to be regulated by immunologic challenge (Charlton et al., 1997; Moore et al., 2003), suggesting that this receptor may link the humoral and nervous system responses to infection and inflammation; rat primary astrocytes and murine microglia have been recently reported to respond to UDP-glucose with increases of intracellular calcium concentrations, highlighting a role of this receptors in glial cells (Fumagalli et al., 2003; Bianco et al., 2005)				

aa, amino acids; AC, accession; chr., chromosome.