

Signaling at Purinergic P2X Receptors

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Key Words

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Abstract

P2X receptors are membrane cation channels gated by extracellular ATP. Seven P2X receptor subunits (P2X_{1–7}) are widely distributed in excitable and nonexcitable cells of vertebrates. They play key roles in inter alia afferent signaling (including pain), regulation of renal blood flow, vascular endothelium, and inflammatory responses. We summarize the evidence for these and other roles, emphasizing experimental work with selective receptor antagonists or with knockout mice. The receptors are trimeric membrane proteins: Studies of the biophysical properties of mutated subunits expressed in heterologous cells have indicated parts of the subunits involved in ATP binding, ion permeation (including calcium permeability), and membrane trafficking. We review our current understanding of the molecular properties of P2X receptors, including how this understanding is informed by the identification of distantly related P2X receptors in simple eukaryotes.

AMPA: α -amino-3-hydroxy-5-methylisoxazole-4-propionic acid

EPSP: excitatory postsynaptic potential

INTRODUCTION

P2X receptors came to our attention as the receptor involved in transmitting the actions of ATP released from sympathetic nerves onto the smooth muscles of the vas deferens and small intestinal arterioles (1). These receptors were distinguished from G protein-coupled metabotropic P2Y receptors (2). The historical development of thinking as to P2X receptors, and the initial associated skepticism, has been recently reviewed (3).

The cloning of P2X receptor subunit cDNAs in 1994 (4, 5) led directly to two major avenues of progress. The first is the demonstration that the proteins have a very widespread tissue distribution in vertebrates and indeed in eukaryotes. This finding in turn has led to the development of receptor antagonists and of mice in which receptor subunits have been genetically deleted; the use of such tools has indicated wide-ranging and hitherto unexpected functional roles for P2X receptors. Second, P2X receptors do not resemble other ion channels at the molecular level. This has spurred many studies to determine the molecular details of the operation of these receptors in the cell membrane. This review focuses on these two areas of research.

PHYSIOLOGICAL ROLES FOR P2X RECEPTORS

Central Nervous System

P2X receptors are widely distributed on central nervous system neurons and glia at the mRNA (6–8) and protein (8, 9) levels. At the protein level, the inadequacy of several of the existing antibodies has handicapped progress, particularly in the case of the P2X₇ receptor (10, 11). Two approaches dominate the literature: the use of either slices of brain tissue or cells dissociated and maintained in culture. The first includes studies of the effects of exogenous ATP and analogs to nerve cells and glia, measuring either the direct effects on membrane currents or intracellular calcium (postsynaptic effects) or the alteration of transmitter release (presyn-

aptic effects). The second complementary approach is to deduce roles for endogenous ATP in intercellular signaling by the use of pharmacological antagonists or knockout mice.

Postsynaptic effects. A careful immunohistochemical study using the postembedding technique for electron microscopy showed that P2X₄ and P2X₆ subunits are found in perisynaptic locations on the hippocampal CA1 pyramidal and cerebellar Purkinje cells (9). There are several reports of residual excitatory postsynaptic potentials (EPSPs) in the presence of high concentrations of blockers of AMPA (α -amino-3-hydroxy-5-methylisoxazole-4-propionic acid), kainate, and NMDA subtypes of glutamate receptor. These EPSPs are rather small in amplitude, and their attribution to ATP is based on blockade by antagonists of limited specificity [habenula (12), hippocampus (13, 14), somatosensory cortex (15–17)]. Pankratov et al. (17) concluded that in the mouse cortex, some glutamate-containing vesicles also contained ATP, whereas others did not. In the lateral hypothalamus of the mouse (and neonatal chick), GABA and ATP are co-released, and each evokes its own postsynaptic current that can be distinguished pharmacologically (18). There is a clear need for a systemic comparison of the properties of such EPSPs between wild-type mice and mice lacking P2X receptor subunits. In the case of the P2X₄ knockout mouse, long-term potentiation in the hippocampus is of reduced amplitude as compared with the wild-type animals, but the underlying cellular mechanism for this remains unclear (19).

Direct postsynaptic effects on neurons of exogenously applied ATP have been described in many parts of the central nervous system and are not reviewed here (see References 8, 20, and 21). More recently, there have been reports of ATP actions on glial cells that are attributable to P2X receptors. Fellin et al. (22) reported ATP-evoked currents in hippocampal astrocytes, although Jabs et al. (23) did not. Astrocytes in the mouse cortex respond to ATP with inward currents exhibiting many of the properties of

the P2X_{1/5} heteromer, and single-cell RT-PCR shows the predominant expression of the two subunits (24).

Presynaptic effects. Presynaptic effects of ATP have been reported at several sites in the central nervous system. The most prevalent effect is an increase in the spontaneous release of glutamate, as detected by increased frequency of spontaneously occurring excitatory postsynaptic currents (EPSCs) in the presence of tetrodotoxin (21). For example, Khakh & Henderson (25) found that ATP enhances glutamate release onto neurons in the motor nucleus of the fifth cranial nerve in the rat but that the receptor involved has properties different from those of the cell bodies of the sensory input neurons that give rise to these presynaptic terminals (located in the mesencephalic nucleus of the fifth nerve) (26). The presynaptic facilitatory effect is rather specific in the hippocampus: ATP facilitates glutamate release at synapses onto inhibitory interneurons, but not at synapses onto CA1 pyramidal cells. This effect was not seen in mice lacking the P2X₂ subunit (27).

In the nucleus tractus solitarius, calcium entry through presynaptic P2X receptors elicits glutamate release sufficient to drive postsynaptic cell firing (28). This effect does not involve voltage-gated calcium channels or tetrodotoxin-sensitive sodium channels. The spontaneous EPSCs elicited by ATP are larger in average amplitude than those occurring spontaneously, suggesting that P2X receptor-induced calcium entry evokes the release of a distinct subpopulation of glutamate-containing vesicles. $\alpha\beta\text{meATP}$ (α,β -methylene adenosine 5'-triphosphate) ($\geq 30 \mu\text{M}$) mimics the effect of ATP, and TNP-ATP [2',3'-O-(2,4,6-trinitrophenyl) adenosine 5'-triphosphate] ($\text{IC}_{50} \approx 1 \mu\text{M}$) blocks these effects. These results are consistent with the involvement of P2X₁- or P2X₃-containing receptors but do not completely exclude other subtypes. It will be important to determine the molecular basis through which P2X receptors appear to access a pool of releasable glutamate distinct from that

which occurs spontaneously. Release of GABA by presynaptic P2X receptors has also been described in the cerebellum (29), and spontaneous release of glycine release has been observed in the dorsal horn of the spinal cord (30).

ATP depolarizes pituitary gonadotrophs, and the properties of the corresponding inward current conform to those expected of P2X₂ and/or P2X₄ subtypes (31). This results in a release of luteinizing hormone. In the neurohypophysis, ATP elicits inward currents on vasopressin terminals, but not on oxytocin-releasing terminals (32, 33). The response has features most similar to those expected for P2X₂ receptors. In both the anterior pituitary and posterior pituitary, the overall physiological context of these ATP effects remains obscure.

Afferent Signaling

Action potentials in afferent nerve fibers are often initiated by the release of a specific chemical from specialized sensory cells that then acts on receptors near the peripheral terminals of the nerve. ATP has been implicated as one such chemical.

Taste. Recent work has indicated that homomeric and/or heteromeric P2X₂ and P2X₃ receptors are essential players in taste transduction. A single mammalian taste bud consists of approximately 100 cells of four distinct types. Type I supporting cells wrap around type II and type III cells to form the characteristic taste bud clusters. Type II cells are the taste receptors expressing distinct G protein-coupled receptors specific for the five differentiated tastes (sweet, sour, salty, bitter, and umami), but type II cells do not synapse with afferent nerves within the taste buds. Presynaptic type III cells synapse, via 5-HT₃ neurotransmission, onto afferent taste nerves. The physiology of type IV cells is not clear, although they are known to function in the regeneration of taste cells (34, 35).

Single-cell RT-PCR of isolated mouse taste bud cells found P2X₂, P2X₄, and P2X₇ subunits (36). Immunohistochemical studies using well-characterized antibodies localized P2X₂ to

EPSC: excitatory postsynaptic current

$\alpha\beta\text{meATP}$: α,β -methylene adenosine 5'-triphosphate

TNP-ATP: 2',3'-O-(2,4,6-trinitrophenyl) adenosine 5'-triphosphate

SNARE: a multiprotein scaffold between exocytotic vesicle and plasma membrane that controls vesicle/plasma membrane fusion and transmitter release

afferent nerve fibers and type III presynaptic cells, localized P2X₃ to afferent fibers only, and showed no P2X₄ protein in taste bud cells or nerve fibers (36–38). ATP release in response to tastants has been measured by luciferase assays from isolated segments of lingual epithelia (39) and also from single isolated type II taste cells via a biosensor assay (40, 41). Release of ATP from these cells is not via classic SNARE-based exocytosis, and both Huang et al. (40) and Romanov et al. (41) have suggested that release is via a hemichannel-like pore made up of proteins of the pannexin or connexin family. Finger et al. (39) recorded afferent firing from both chorda tympani and glossopharyngeal gustatory nerve trunks in mice lacking both P2X₂ and P2X₃ receptors. These researchers found an almost complete absence of nerve response to tastants

applied to the oral cavity; responses to mechanical and other chemical stimuli were not altered. Moreover, behavioral responses to sweeteners, glutamate, and bitter substances were diminished or absent in these double-knockout mice. Finger et al. (39) also found moderately diminished neural and behavioral responses to tastants in single-knockout (P2X₃ or P2X₂) mice. Taken together, these studies provide convincing evidence for the role of ATP as the primary neurotransmitter in taste perception, acting on homomeric P2X₂ receptors, homomeric P2X₃ receptors, and heteromeric P2X_{2/3} receptors.

However, ATP neurotransmission in mouse taste buds (**Figure 1**) may be atypical. Rong et al. (42) used an isolated, intra-arterially perfused rat tongue with separated gustatory (chorda tympani) and general sensory (lingual) nerves. They observed P2X₂ receptor and P2X₃ receptor responses only from lingual nerves that did not respond to tastants applied to the tongue, whereas chorda tympani nerves responded robustly to salty or acidic solutions but not to ATP or ATP analogs. There may be technical difficulties in the use of an isolated tongue preparation that can account for the contrasting results, or this study may serve as a caution against the overwhelming domination of mouse tissue and mouse models in physiology research today.

Hayato et al. (36) have demonstrated the presence of functional P2X₇ receptors on subpopulations of taste bud cells, although the specific cell type has not been ascertained. Taste bud cells are among the shortest-lived of all mammalian cells, having turnover rates of only several days, whereby old taste cells are removed via apoptosis. Because prolonged P2X₇ receptor activation leads to apoptotic cell death, this demonstration of functional P2X₇ receptors in taste buds indicates a possible role for P2X₇ receptors in taste cell regeneration.

Hearing. Immunohistochemical and electrophysiological studies have been carried out in mouse, rat, and guinea pig cochlea (reviewed in References 43 and 44). These studies have provided evidence for ATP as a neurotransmitter

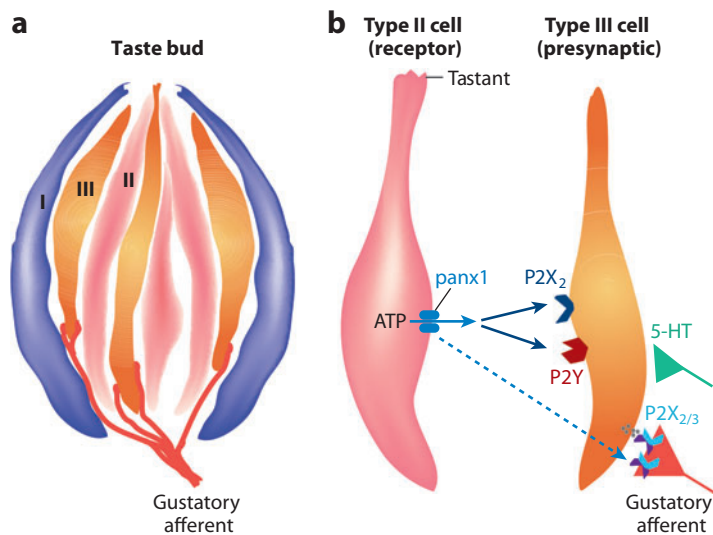


Figure 1

P2X₂ and P2X₃ receptors are involved in taste transduction. (a) Diagram of a single taste bud indicating three of the four types of taste cells: type I supporting cells, type II receptor cells, and type III presynaptic cells; type IV basal cells are not shown (35). (b) In response to tastants, specific type II cells release ATP via pannexin-1 (panx1) hemichannels (40, 41). The released ATP acts in a paracrine fashion to activate P2X₂ and/or G protein-coupled P2Y receptors in type III cells; this leads to increased intracellular calcium, which evokes classical synaptic release of ATP onto presynaptic P2X_{2/3} receptors on afferent fibers (39). ATP released from type II cells may also act in a paracrine manner to activate presynaptic afferent P2X_{2/3} receptors (dashed arrow). Depolarization of type III cells by ATP also leads to the release of 5-HT onto gustatory afferents (35).

or neuromodulator in afferent (but not efferent) auditory transmission. Patch-clamp recordings from inner and outer hair cells show ATP-gated currents that are likely mediated by P2X₂ receptors, whereas neuronal spiral ganglion cells show P2X₃ receptor-mediated inward currents (43–48). The electrophysiological data correspond to the immunolocalization of these receptor subunits. Housley and coworkers have amassed a large body of data that implicates P2X receptors, primarily P2X₂, in modulation of the endocochlear potential (43–45). However, there are currently no publications regarding the hearing phenotype of mice lacking P2X₂ and/or P2X₃ receptors or studies using selective P2X₃ antagonists to hearing behavior in animals or humans.

Chemoreceptors. The carotid body contains glomus cells that release a primary afferent transmitter onto primary afferent nerves in response to hypoxia. The identity of this transmitter (as also the primary hypoxia sensor) has been the subject of long-standing debate (49). Rong et al. (50) reported that ATP is released from type 1 glomus cells in response to hypoxia, that ATP activates primary afferent nerves (probably through a P2X_{2/3} heteromeric receptor), and that the excitation of primary afferent nerves by hypoxia is much reduced in mice lacking the P2X₂ subunit (see Reference 51).

Bladder. Accumulating evidence indicates a role for P2X receptors in the initiation of primary afferent signaling in hollow viscera and particularly the urinary bladder. Distension of the bladder results in ATP release from urothelial cells (52). ATP and $\alpha\beta$ meATP increase the excitability of bladder afferent nerves (53). Bladder afferent nerves express P2X₃ receptor subunits, and in the anesthetized P2X₃ knockout mouse the bladder must be distended to a greater volume than in wild-type mouse to initiate the voiding reflex (54, 55).

Pain. ATP acting at P2X receptors has been implicated in pain sensation at both periph-

eral and central sites within the nervous system. ATP applied within the skin evokes pain (56), and P2X₃ subunits are expressed on a subset of primary afferent neurons implicated in pain sensation (57, 58). Three lines of evidence implicate P2X₃ receptor subunits in some aspects of neuropathic and inflammatory pain perception: Some, although by no means all, pain-related behavior is blocked by antagonists selective for P2X₃ subunit-containing receptors (59), by reduction in P2X₃ subunit expression by RNA knockdown (60), and by genetic elimination of the P2X₃ subunits (reviewed in detail in Reference 61).

A role for P2X₄ receptors in neuropathic (but not inflammatory) pain has more recently been proposed. P2X₄ subunit expression is increased in microglia of the dorsal horn of the spinal cord following spinal nerve ligation, a commonly used model for neuropathic pain. The mechanical allodynia resulting from the ligation was reduced by intrathecal TNP-ATP and by intrathecal antisense oligonucleotides that somewhat reduced P2X₄ receptor expression (62). Conversely, intrathecal administration of cultured microglia that had been stimulated with ATP could induce allodynia in naive rats. It was proposed that spinal nerve ligation activates dorsal horn microglia by a mechanism involving P2X₄ receptors and that they release brain-derived nerve growth factor (BDNF). BDNF then acts on spinal cord interneurons to stimulate the expression of the potassium-chloride cotransporter KCC2, and the resultant reduction of intracellular chloride converts the usually inhibitory action of GABA into excitation (63). One difficulty with the attribution of these effects to the activation of P2X₄ receptors is that TNP-ATP has several nonspecific actions and is in any event a weak blocker of P2X₄ receptors (64). It will be important to repeat these studies in mice in which the P2X₄ and P2X₇ receptors have been knocked out, or with currently available P2X₇ receptor antagonists.

More recently, P2X₇ receptors have also been implicated in pain perception. Disruption of the P2X₇ receptor gene markedly reduces both neuropathic and chronic inflammatory

BDNF: brain-derived nerve growth factor

UVEC: umbilical vein endothelial cell

IL-1 β : interleukin-1 beta

IL-1Ra: a naturally occurring protein that binds to IL-1 receptor without activating it and prevents binding of IL-1 β

pain in mice (65), and antagonists with high selectivity for P2X₇ receptors strongly inhibit neuropathic pain in rats (66). The mechanism of these actions is as yet unclear but presumably involves the rearrangement of sensory processing pathways and decreased release of proinflammatory cytokines from microglia in the spinal cord.

Cardiovascular System

P2X receptors have widespread expression throughout tissues of the cardiovascular system (6). Our understanding of their function in two tissues—(a) the endothelium and its role in the control of blood pressure and (b) the platelet with respect to blood coagulation—has advanced rapidly in recent years.

Vascular endothelium. Vascular endothelial cells coexpress P2X₄ and P2X₇ receptors: mRNA and protein levels for the former are typically 5–20-fold higher than for the latter (67). The heterogeneity of vascular endothelia should be kept in mind when functional roles of any endothelial protein are discussed. One of the clearest examples of this heterogeneity is a microarray study performed by Chi et al. (68) on 53 different human endothelial cell lines obtained from various large and small arteries and veins; distinct patterns of gene expression were detected on the basis of not only tissue site, developmental stage, and artery versus vein type but also specific functions such as left/right asymmetry. Nevertheless, in endothelial cells increased intracellular calcium directly leads to nitric oxide (NO) production and release with consequent smooth muscle vasodilation; therefore, calcium assays (using Fura2 or Indo1) are confidently used in endothelial cells as indirect readouts of NO activity (69). Human and bovine umbilical vein endothelial cells (UVECs) have been the most commonly used preparations for *in vitro* studies of purinoceptor function in endothelia.

Ando and colleagues (70) have presented evidence for autocrine activation of P2X₄ receptors in human and bovine UVECs via shear

stress-induced release of ATP from endothelial cells. The key evidence for P2X₄ receptor involvement is that antisense oligonucleotides that decreased protein levels abrogated shear stress-induced calcium transients and that heterologous expression of P2X₄ receptors in human embryonic kidney 293 cells resulted in a shear stress-induced calcium transient. Surprisingly, the ATP-induced responses blocked by P2X₄ receptor antisense oligonucleotide treatment were observed only with low ATP concentrations (<1 μ M), concentrations that do not activate homomeric human or rodent P2X₄ receptors or heteromeric rat P2X_{4/6} receptors (71).

In contrast, Wilson et al. (67) found no evidence for P2X₄ receptor-mediated membrane currents or calcium transients in human UVECs even under conditions (interferon- γ and TNF α stimulation) that upregulated P2X₄ receptor mRNA and protein by 10–75-fold; these investigators found that >95% of total P2X₄ protein is intracellular. The predominantly intracellular localization of the P2X₄ receptor has been documented in several other cell types and appears to be the default expression pattern of this receptor (72, 73). After activation by interferon- γ and TNF α , these human UVECs released small amounts of proinflammatory interleukin-1 beta (IL-1 β) in response to P2X₇ receptor stimulation but simultaneously released anti-inflammatory IL-1Ra; the net effect was anti-inflammatory.

Although these *in vitro* studies of P2X receptor function in UVECs may seem contradictory, Ando and colleagues have provided conclusive evidence for P2X₄ receptor involvement in endothelial function in studies, using wild-type and P2X₄ receptor knockout mice (74). First, they found that flow-induced increases in calcium and NO production in cultured endothelia obtained from lung microvessels were virtually absent in preparations from P2X₄ receptor knockout mice and that these responses were regained with the reintroduction of P2X₄ receptor by adenoviral constructs. Second, these researchers found that *in vivo* vasodilation of precontracted cremaster

arterioles induced by intra-arterial ATP (but not acetylcholine), and the associated increased blood flow, was decreased in knockout mice; the decrease (30–100%) depended on ATP concentration and flow rate. A similar inhibition of ATP-induced vasodilation occurred in isolated mesenteric arteries from knockout mice. Third, P2X₄ receptor knockout mice had significantly higher blood pressure and lower basal plasma concentrations of nitrites and nitrates (an indicator of NO levels) than did wild-type controls. Fourth, Ando and coworkers found that the resting carotid artery diameter of P2X₄ receptor knockout mice was significantly smaller and that their smooth muscle layers were thicker compared with wild type. Moreover, these investigators used the carotid artery ligation to elicit neointimal depositions and long-term decreased vessel diameter, which is commonly employed as an animal model of atherosclerotic disease; P2X₄ receptor knockout mice showed no further decrease in vessel diameter. Chamberlain et al. (75) performed similar carotid artery ligation experiments on P2X₇ receptor knockout mice and found no differences in responses, thus ruling out P2X₇ receptor involvement in this animal model of vascular disease. Taken overall, the vascular phenotype of P2X₄ receptor knockout mice resembles that seen in endothelial nitric oxide synthase (eNOS) knockout mice (74), supporting the idea that the critical functional role of P2X₄ receptors in mouse vascular endothelia resides in the modulation of NO production and release. This finding has implications for the possible use of P2X₄ receptor antagonists in atherosclerosis.

Thrombosis. The study of ADP acting on P2Y receptors on platelets has a long history that has brought considerable clinical utility. The main players are the P2Y₁ and P2Y₁₂ receptors; the latter is blocked by the antithrombotic agents such as clopidogrel (76). Platelets also express P2X₁ receptors, and their activation by $\alpha\beta$ meATP induces a rapid calcium influx associated with a transient shape change (77, 78). The bleeding time of P2X₁ receptor knockout mice is normal, suggesting lit-

tle impairment of normal hemostasis; however, thrombosis associated with injury of the walls of small arterioles is reduced in the knockout mice (79). Thus, the P2X₁ receptor may play a role in thrombus formation under conditions of high shear stress, suggesting that a P2X₁-selective antagonist may be particularly effective in inhibiting thrombosis in conditions of stenosis, in which shear stress is high, while having little effect on normal hemostasis, in which shear stress is less (76).

Respiratory System

An important recent development has been the discovery that P2X receptors expressed by the ciliated epithelial cells of the bronchi can control ciliary beat frequency and thus the clearance of mucus from the airways. The subject has obvious importance for a range of lung diseases, including cystic fibrosis. A second area of substantive progress has been the central control of respiration. Sophisticated assays of ATP release combined with electrophysiological recordings have strongly supported a role for P2X receptors in the ventral surface of the medulla.

Ciliated epithelia. Ciliated epithelia play a critical role throughout the airways in clearing mucous secretions. Agents that increase ciliary beating, such as β_2 -adrenergic and P2Y₂ receptor agonists, provide key therapies in several acute and chronic respiratory diseases (80). ATP increases ciliary beating in ciliated epithelia from nose, trachea, and other airway epithelia in several species via activation of both P2Y and P2X receptors (80, 81). Silberberg and coworkers have determined the properties of P2X responses in rabbit airway ciliary epithelia (82, 83). They have concluded that a novel P2X receptor, which they term P2X_{cilia}, underlies the ionotropic purinergic response. The functional responses reported for rabbit P2X_{cilia} are the same as those reported for rodent P2X₇ receptors in most respects. These responses include agonist dose responses, divalent cation sensitivity, facilitation of response

***N*-methyl-D-glucamine (NMDG):** the largest soluble cation (molecular weight 195.2) used in biophysical studies to estimate channel pore size

during prolonged agonist application, and inhibition by extracellular sodium ions. The key differences are that the rabbit cilia do not show uptake of dyes (such as propidium), do not show increases in *N*-methyl-D-glucamine (NMDG) permeability with prolonged agonist application, and have currents that are potentiated by zinc (1 μ M) (83). Because dye-permeable pore formation and zinc inhibition are typical properties of P2X₇ receptors, these workers ruled out (homomeric) P2X₇ receptor involvement. However, the dye/NMDG-permeable properties of P2X₇ receptor function depend on the density of P2X₇ receptor protein expression (84) and/or the presence of other associated interacting proteins (85), and zinc modulation of many ligand-gated ion channels, including P2X receptors, commonly shows biphasic potentiation and inhibition of agonist-evoked responses depending on zinc concentration (21). The complete concentration dependence of the action of zinc in airway epithelial cells remains to be determined.

The Silberberg group (82, 83) also found that ivermectin potentiates the amplitude of the ATP-induced cationic current. Because ivermectin potentiation is a selective characteristic of P2X₄ receptors (21), this research group's general conclusion was that P2X_{cilia} most likely resulted from a heteromeric P2X_{4/7} receptor. However, ivermectin potentiated only the amplitude, and not the duration, of the ATP response in rabbit ciliary epithelia, whereas studies on ectopic or endogenous P2X₄ receptors have also shown a marked prolongation of the duration of P2X₄ receptor responses. We have recently observed that ivermectin significantly potentiates the amplitude of mouse (but not rat) P2X₇ receptor-induced current (J. Sim, unpublished observations) in a manner similar to that observed by Silberberg and colleagues in rabbit cilia.

Central control of respiration. Dale and colleagues (86) have cleverly developed an enzymatic/amprometric electrode to measure local ATP concentration. In one application of this device (87), they measured ATP concentrations

in the ventral medulla during hypoxia. In vivo, they detected ATP release only in those discrete regions of the medullary surface known to correspond to classical CO₂ chemosensitivity. Slices of medulla maintained in vitro also released ATP in response to increased CO₂. Higher concentrations of CO₂ were required to stimulate respiratory activity in the presence of P2X receptor antagonists. Moreover, exogenous ATP itself mimicked the effect of increased CO₂ levels. These findings led the authors to suggest that ATP released in response to increased CO₂ acts on P2X receptors expressed on the dendrites of ventral respiratory column neurons and thus determines the adaptive responses in breathing (51, 87). This implies a key role for ATP released from peripheral chemosensors to initiate action potentials in primary afferent nerves (see above section, Afferent Signaling) and a very similar role for ATP in the ventral medulla to drive the central respiratory response. In both cases, an understanding of the mechanism of ATP release is urgently needed.

Genitourinary System

ATP acting at P2X receptors was first recognized as the efferent transmitter process from sympathetic nerves to the vas deferens (3), and more recently it has become clear that ATP released from urothelial cells can activate afferent nerves in a bladder distention reflex. More recent work implicates P2X receptors in the initial stage of urine formation, namely in the feedback mechanism for maintaining urinary filtration at the glomerulus in a wide range of physiological conditions.

Tubuloglomerular feedback. Strong evidence now exists for a critical role of P2X₁ receptors in renal autoregulation. Autoregulation is the process whereby the kidney is able to maintain constant renal blood flow and glomerular filtration rate over changes in arterial pressure from approximately 80 to 180 mm Hg in the absence of changes in sympathetic input and/or vasoactive substances (88).

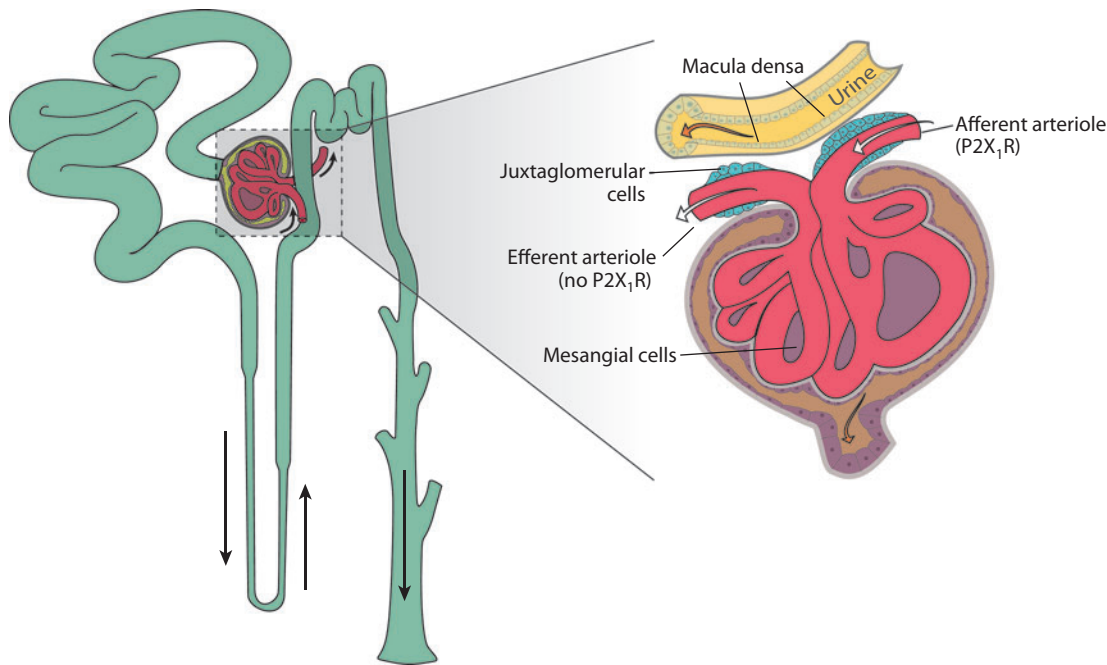


Figure 2

P2X₁ receptors are involved in renal autoregulation via tubuloglomerular feedback. To the left is a single nephron, with the glomerular apparatus enlarged on the right. Changes in tubular fluid flow rate occur in response to changes in glomerular filtration rate or the rate of reabsorption in the proximal tubule. Changes in glomerular filtration rate are sensed by macula densa cells that release ATP onto P2X₁ receptors on afferent arteriolar smooth muscle, triggering an increase in afferent arteriolar resistance. This reduces pressure in the glomerular capillaries and decreases glomerular filtration (i.e., tubuloglomerular feedback).

This autoregulation is due to both intrinsic myogenic response of smooth muscle and tubuloglomerular feedback, and P2X₁ receptors appear to be involved in the latter process. Tubuloglomerular feedback involves the juxtaglomerular apparatus, consisting of the macula densa, the glomerulus, and both efferent and afferent arterioles of the same nephron (**Figure 2**) (89). Apical macula densa cells are exquisitely sensitive to increases in NaCl concentration in the distal tubular fluid; as little as 5–10 mEq l⁻¹ leads to afferent arteriolar vasoconstriction via tubuloglomerular feedback (90). How does this occur? Bell and colleagues (91, 92) used P2X₂ receptor-expressing PC12 cells as ATP biosensors. When they placed whole-cell patch-clamped (or Fura2-loaded) PC12 cells into contact with rabbit macula densa cells, they recorded (P2X₂-activated) currents (or calcium transients) when

luminal NaCl was increased by 5- to 10-fold; the responses corresponded to the release of ~10 μM ATP. Smaller changes in NaCl concentration were not examined in this assay. However, the Navar group made *in vivo* microdialysis measurements of canine renal interstitial ATP (and adenosine) in response to changes in renal arterial pressure within the autoregulatory range (90, 93). They found a tight correlation between interstitial concentrations of ATP (but not adenosine) and tubuloglomerular feedback in response to small stepwise changes in renal arterial pressure. Taken together, these results show conclusively that ATP is released from the macula densa in response to luminal changes in NaCl. Although the molecular mechanism of the ATP release remains unknown, it is currently hypothesized that this ATP release triggers tubuloglomerular feedback.

However, there is considerable debate as to how such tubuloglomerular feedback is achieved. On one side of the debate is the adenosine A₁ receptor hypothesis: Specific ectonucleotidases hydrolyze released ATP to adenosine with subsequent activation of vasoconstrictor A₁ receptors on afferent arterioles (94). On the other side of the debate is the P2X₁ receptor hypothesis: Released ATP directly activates vasoconstrictor P2X₁ receptors on afferent arteriolar smooth muscle (89). The main support for the former hypothesis consists of the following. (a) The specific ectonucleotidases required to break down ATP to adenosine are highly expressed in the juxtaglomerular apparatus, (b) specific A₁ receptor antagonists inhibit or abolish tubuloglomerular feedback, and (c) A₁ receptor knockout mice do not exhibit tubuloglomerular feedback (94–97). Equally strong support for the latter model comes from findings that (a) P2X₁ receptors are expressed only on afferent arterioles but not on efferent arterioles or other kidney cells, (b) a selective P2X₁ receptor antagonist (NF279) abolishes tubuloglomerular feedback, and (c) P2X₁ receptor knockout mice do not exhibit tubuloglomerular feedback (89, 98, 99). Interpretations of results from blockade of A₁ receptor with antagonists or knockout mice may be more complicated than for results from similar blockade of P2X₁ receptors because of the widespread expression of A₁ receptors in the kidney and renal vasculature and because of the concomitant vasodilatory actions of the similarly widespread expression of A₂ receptor throughout the renal vasculature (100). It is also likely that either or both ATP and adenosine underlie tubuloglomerular feedback, depending on subpopulations of nephrons and their individual metabolic state; there are more than 300,000 individual nephrons in a rodent kidney. Moreover, subtle differences in *in vivo* microdialysis and other *in vitro* techniques and protocols among laboratories may also contribute to experimental differences. This A₁ receptor/P2X₁ receptor puzzle may be resolved by exchange of the knockout mice among the relevant groups and parallel experiments in the

same laboratory using both sets of mice, along with a more quantitative comparison of the kinetics and pharmacology of inhibition of tubuloglomerular feedback by A₁ and P2X₁ receptor antagonists.

Renal fibrosis. The abnormal accumulation of extracellular matrix (i.e., fibrosis) is associated with all chronic renal diseases. Renal mesangial cells are primarily responsible for extracellular matrix deposition via the secretion of transforming growth factor β (TGF β), which leads to increased accumulation of fibronectin and other extracellular matrix proteins (101). Thus, mesangial cell dysfunction is a key contributor to interstitial fibrosis and, as such, has attracted recent interest in regard to the potential roles purinoceptors may play in normal and diseased mesangial cells.

Schulze-Lohoff et al. (102) first showed that P2X₇ receptor activation induces mesangial cell apoptosis. Harada et al. (103) showed that ATP exerts dual actions on isolated rat mesangial cells; P2Y₂ receptor and/or P2Y₄ receptor activation stimulated mesangial cell proliferation, and P2X₇ receptor stimulation with BzATP [2',3'-O-(4-benzoyl-benzoyl) adenosine 5'-triphosphate] led to increased apoptosis. Soloni et al. (104) confirmed these findings and found that P2X₇ receptor stimulation increased secretion of TGF β and subsequent fibronectin levels, effects that were significantly enhanced when cells were grown in a high-glucose medium (mimicking hyperglycemic conditions). Goncalves et al. (105) used an animal model of interstitial fibrosis, unilateral ureteral obstruction, in both wild-type and P2X₇ receptor knockout mice. This group found that the classical features of unilateral ureteral obstruction (high macrophage invasion into the interstitium, high TGF β secretion, increased mesangial cell apoptosis, and interstitial fibrosis) were all significantly abrogated in the P2X₇ receptor knockout mice. Taken together, these results suggest a physiologically significant role for P2X₇ receptors in renal fibrosis and may point to the therapeutic

potential of P2X₇ receptor antagonists in the treatment of chronic renal diseases.

However, Soloni et al. (106) used human mesangial cells from normal segments of kidneys obtained at nephrectomy and repeated experiments they had carried out on rat mesangial cells, with surprising results. They found that ATP clearly induced significant apoptosis in human mesangial cells, but could find no evidence for the presence or involvement of P2X₇ receptor; moreover, their pharmacological data were not readily compatible with the involvement of known human P2Y or P2X receptors. Only high concentrations of ATP (>1 mM), but not of BzATP, induced the apoptotic response, suggesting that ATP breakdown to adenosine may have been responsible; it has been shown that apoptosis of a hepatic epithelial cell line by ATP and ADP is due to the activation of adenosine A₃ receptors in response to hydrolysis of these nucleotides to adenosine (107). In any event, results from this study serve as caution against overreliance on, and overinterpretation of, results obtained from rodent tissues and cells.

Gastrointestinal System

The nonadrenergic, noncholinergic inhibitory transmission to the smooth muscle of the intestine was one of the very first demonstrations of a functional role for extracellular ATP (reviewed in References 1 and 3). Further, quite distinct roles in the gastrointestinal tract are now emerging. These include accessory structures such as salivary glands, liver, and exocrine pancreas. They also include signaling between nerve cells of the enteric nervous system within the gut wall.

Exocrine glands. Luminal ATP has long been known to activate purine receptors on several secretory epithelia with the consequent secretion of chloride, potassium, bicarbonate, or hormonal peptides (80, 81). The majority of studies on nucleotide-induced epithelial secretion show clearly that the P2Y family of purinocep-

tors, particularly the P2Y₂ receptor, is primarily responsible (80, 81). Additional involvement of P2X₇ receptors and/or P2X₄ receptors has also been suggested, but the interpretation is confounded by marked species differences in the physiological and pharmacological properties of these two receptors. Submandibular gland acinar and ductal cells from rat and mouse have been the most studied. Patch-clamp recordings show both P2X₄ receptor- and P2X₇ receptor-mediated currents, whereas Fura2 intracellular calcium measurements show predominantly P2X₇ receptor-mediated responses, with a very small contribution via P2X₄ receptors (<5% of total calcium response). Stimulation of P2X₇ receptors, but not of P2X₄ receptors, induces kallikrein secretion from ductal cells (108). Dehaye and colleagues (108–110) have built up a strong case for P2X₇ receptor signaling to salivary secretion through the modulation of phospholipid signaling processes, particularly phospholipase A2 and phospholipase D. The strongest evidence comes from a study comparing responses from P2X₇ receptor wild-type and knockout mice, in which (a) ATP no longer induced any phospholipid signaling, (b) the ATP-induced calcium signal was decreased by ~90–95%, and (c) the saliva from the knockout mice showed a significantly decreased potassium content (110). This study thus directly demonstrated the involvement of P2X₇ receptor in salivary secretions, but the evidence for phospholipid signaling as the underlying mechanism remains correlative.

The difficulty of matching P2X phenotype in epithelia with P2X protein expression is highlighted in a recent study on human gallbladder and rat liver bile duct cells, in which ATP and BzATP potently stimulated chloride secretion when applied to the apical surfaces of these epithelia (111). In these bile ducts, P2X₄ but not P2X₇ receptors were expressed by RT-PCR, immunoblot, and immunohistochemistry. However, the properties of the secretory response and whole-cell patch clamp recordings were more typical of activating P2X₇ than P2X₄ receptors. The recent availability of

BzATP: 2',3'-O-(4-benzoyl-benzoyl)adenosine 5'-triphosphate

highly selective and potent rodent and human P2X₇ receptor antagonists may resolve some of these discrepancies.

Intestinal motility. Afferents traveling by the vagus nerve or the dorsal root ganglia signal intestinal activity to the central nervous system. The role of P2X receptors on such nerves is described above (see Afferent Signaling section). In contrast, considerable evidence exists for a key role for P2X receptors in signaling within the enteric nervous system that underlies coordinated movements of the intestine. The principal intrinsic afferent neuron of the enteric nervous system is the AH cell (112). AH cells (which can be identified by immunoreactivity for calbindin D28K) express P2X₂ and P2X₃ subunits; there is evidence that the afferent terminals of these neurons, in the mucosal layer and in the muscle layers, are activated by ATP through P2X receptors, but the subtype has not been conclusively identified (113). Another main class of neuron, the descending interneurons on the pathway to the circular muscle motoneurons, also expresses P2X₂ subunits; in these cells ATP substantially contributes to the fast EPSP when presynaptic fibers are stimulated electrically, as shown by comparing wild-type and knockout mice (114). Peristalsis of the small intestine recorded *in vitro* is clearly impaired in the ileum taken from P2X₃ receptor knockout mice, suggesting most likely that the afferent neurons are less sensitive to the initiating distension (115).

Immune System

P2X₁, P2X₄, and P2X₇ receptor proteins coexist in most immune cells, including mast cells, B and T lymphocytes, monocytes, macrophages, microglia, and osteoclasts (8). Cation currents and calcium influx with the features expected for each of these P2X receptors have been demonstrated in all these immune cells. However, only P2X₇ receptors are thus far established as having physiological roles in the immune system. There is a long history of P2X₇ receptor involvement in inflammation via the

unique ability of the P2X₇ receptor to induce the rapid activation of caspase-1 with subsequent release of the proinflammatory cytokine IL-1 β from activated macrophage and microglia (116, 117). The underlying mechanism(s) by which P2X₇ receptor activation leads to caspase-1 activation and IL-1 β release remains a matter of considerable debate and study, but abrogation of inflammatory responses by inhibition of P2X₇ receptors is directly linked to concomitant decreased levels of IL-1 β in plasma or in the area of injury (116, 117).

Two lines of mice have been bred with disrupted P2X₇ receptors. In one [Glaxo mice (65)] 5' exons were disrupted by a neomycin-resistant vector. In the other [Pfizer mice (118)] the gene was disrupted in the portion encoding the cytoplasmic tail, leaving open the possibility of expression of a truncated receptor. In both lines, ATP did not induce the release of IL-1 β or other proinflammatory members of the IL-1 family (IL-18 and IL-1 α) from endotoxin-activated macrophages. Moreover, mechanical hypersensitivity (allodynia) in response to adjuvant injection into the paw, as well as in response to partial nerve ligation, was absent, and levels of IL-1 β from the area of paw inflammation systemically were reduced. Recently, similar results have been obtained from wild-type mice treated with the selective P2X₇ antagonist A740003 (119). Researchers have described other potent and selective P2X₇ receptor antagonists that show either human or rodent specificity (119, 120). One such molecule (AZD9056) produces clinically relevant improvements in patient assessments of symptoms, and reduced swollen and tender joints, in a Phase II clinical trial of 75 patients with active rheumatoid arthritis treated for one month (121). These early results hold promise that P2X₇ receptor antagonists in inflammation may be the first therapeutic benefits to emerge from P2X receptor research.

Killing intracellular mycobacteria. Stimulation of macrophage P2X₇ receptors can lead to the death of several types of intracellular bacteria, and the killing of mycobacterium

tuberculosis is of much clinical relevance (122). The killing of intracellular bacilli directly parallels P2X₇ receptor-mediated macrophage apoptosis (122). Because IL-1 β leads to apoptosis of surrounding cells, the direct cause of macrophage apoptosis and death of intracellular pathogens may be the P2X₇ receptor-induced release of IL-1 β . Studies with RAW 264.7 murine macrophages have ruled out this possibility. These macrophages do not show caspase-1 activation/IL-1 β release be-

cause they lack a key adaptor protein (ASC) required to form the multiprotein complex (the inflammasome; see **Figure 3**) (123). Yet these macrophages express high levels of P2X₇ receptor, and prolonged activation leads to macrophage apoptosis and concomitant killing of intracellular pathogens (123, 124). It is not known whether macrophage apoptosis is necessary for killing of intracellular mycobacteria. However, the P2X₇ receptor-mediated apoptotic process is associated with the formation of

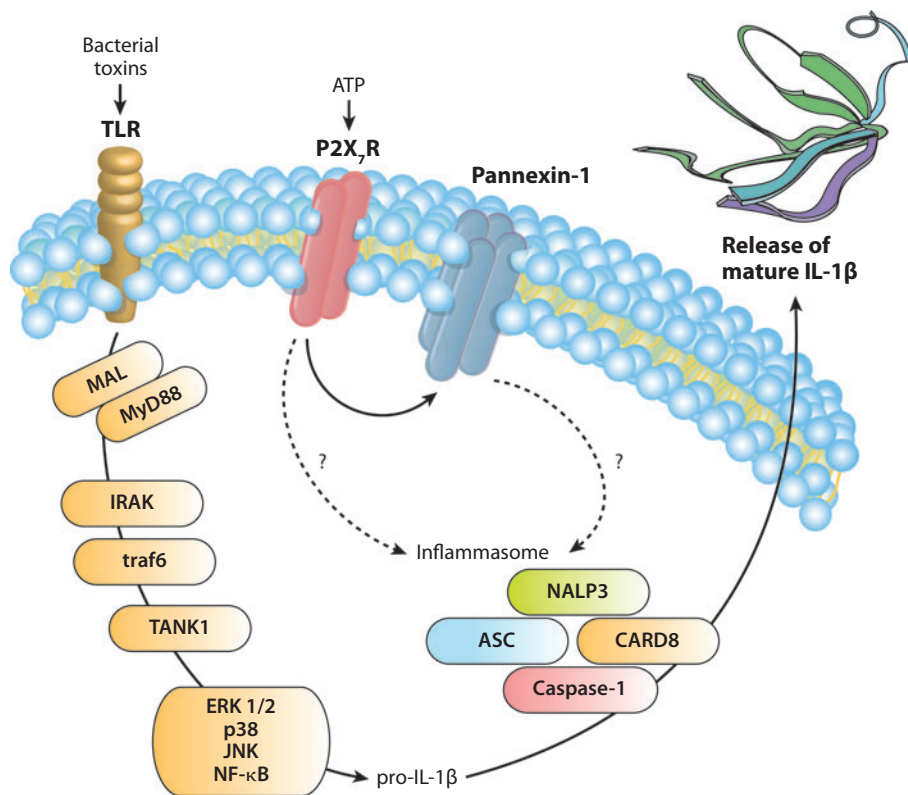


Figure 3

P2X₇ receptor (P2X₇R) and the NALP3 inflammasome. An initial inflammatory insult (e.g., bacterial endotoxins) activates Toll-like receptors (TLRs) on monocytes, macrophages, and microglia to initiate the classical nuclear factor- κ B (NF- κ B) cascade and the synthesis of pro-interleukin-1 β (pro-IL-1 β). However, in the absence of a secondary stimulus, little or no processing or release of IL-1 β occur (116, 117). High concentrations of extracellular ATP, present at sites of inflammation, activate a P2X₇R/pannexin-1 protein complex (172). This leads to the activation of caspase-1, with the subsequent formation of a multiprotein intracellular complex, the NALP3-type inflammasome (173). Activated caspase-1 within the inflammasome scaffold now cleaves pro-IL-1 β , and the release of mature, bioactive IL-1 β follows. Blockade of either P2X₇R or pannexin-1 independent of P2X₇R prevents rapid IL-1 β processing and release (117, 172). Mechanisms by which P2X₇R/pannexin-1 activation leads to caspase-1 activation and the release of mature IL-1 β remain uncertain and are likely to be tissue/cell and stimulus specific.

large intracellular vacuoles and with the fusion of lysosomes with the bacilli-containing phagosomes, thus providing a means for killing the mycobacteria via lysosomal enzymes (122, 124, 125).

Osteoclasts. Several studies have implicated P2X₇ receptors in the formation of multinucleated giant cells in granulomatous diseases and in normal and pathological formation of multinucleated osteoclasts in bone (126). In one study, the Pfizer P2X₇ receptor knockout mouse showed abnormal bone formation and resorption in the form of a reduced periosteal circumference and decreased cortical bone content (127). In another study using Glaxo P2X₇R knockout mice, there was no difference between wild-type and knockout mice in the ability of mononuclear phagocytes to generate multinucleated osteoclasts (128). The P2X₇ receptor antagonist AZ11657312 significantly reduced bone resorption, synovial inflammation, and the number of osteoclastic giant cells (129). Thus, although it has not yet been determined whether P2X₇ receptor alters bone metabolism via alterations in the formation of multinucleated giant cells, recent studies provide strong evidence for its role in modulating the function of osteoclasts in bone.

P2X₇ polymorphisms and immune function. The human P2X₇ receptor is associated with more than 500 single-nucleotide polymorphisms, some of which render the P2X₇ receptor nonfunctional. Wiley and colleagues (124, 130) have examined five specific P2X₇ receptor polymorphisms known to result in nonfunctional P2X₇ receptors when examined in heterologous expression systems. In a cohort of more than 700 mainly Caucasian adults, these investigators found that 51% showed the wild-type gene at the five alleles they examined, 40% were heterozygous at one of these alleles, and 25% of the heterozygous adults showed reduced functional readouts compared with wild type. Approximately 3% of the population was homozygous for T357S/E496A without any functional P2X₇ receptor response

in their blood-derived macrophages (130). Macrophages obtained from individuals consistently show “high, inducible, or no” response to P2X₇ receptor stimulation (125, 130), and specific polymorphisms in the P2X₇ receptor gene may explain these observations. The molecular identity of individual P2X₇ receptors may of course alter their susceptibility to newly developed therapeutic antagonists.

MOLECULAR PROPERTIES OF P2X RECEPTORS

Direct structural approaches with single-particle electron microscopy and atomic force microscopy strongly support the view that the P2X receptor channel forms as a trimer (131, 132). This stoichiometry was originally based on polyacrylamide gel electrophoresis under nondenaturing conditions (133) and is supported by measurements of single-channel kinetics (134), or currents evoked by ATP through channels formed from concatenated subunits with reporter mutations (21, 135).

In the past few years, deduced P2X receptor sequences have become available from the genomes of several simple eukaryotes. Where these have been shown to function as ATP-gated ion channels, the limited sequence relatedness helps to identify those residues or parts of the molecule that are most critical for function. **Figure 4** illustrates this for five P2X receptors. These are green algae (*Ostreococcus tauri*), choanoflagellate (*Monosiga brevicollis*), slime mold (*Dictyostelium discoideum*), trematode worm (*Schistosoma mansoni*), and the vertebrate *Rattus norvegicus* (21, 136–138). The first two are single-celled organisms that are phylogenetically close to the origins of the divergence of plant and animal cells, the third has a life cycle with both single-celled and multicellular components, the fourth is a segmented worm, and the fifth is man.

Several points can be taken from this comparison of sequences of functional receptors. First, only relatively few residues, notably tyrosines, lysines, and glycines, are completely conserved. In almost all cases, these conserved

N terminus, TM1, post-TM1

```
rP2X2 ----MVRRLARGCWSAFWDYETPKVIVVRNRRLGFVHRMVQLLILLYFVWYVFIVQKSYQDSETGPESSIITKVKG 072
schis -----MVKGIAVLFEYETPKLVQISNLIKIGVTRQLLQLIVLIVVVCWVMIYEKGYQENDI-AKSAVTTKVKG 066
choan MAASGFWGSIQGGIYSMLEYDTLTKVHIRSKVGLIFRLQLITILAYVVGYGIIYQKGYQEVDS-AVSTVTLTKVKG 075
algae -----MGLSYTKQKSVVIRSRWRLGALYYGLVAVVLAYVGFFLVYVERGYQRTSR-AVGNIGLKVKG 059
dicty -----MGFSFDWDDIFQYSTVKIVIRDRRLGILHLSFLVGVIVAYIVVYSAIIKKGYLFTEV-PIGSVRTSLKG 078
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Ectodomain

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rP2X2 -----ITMSEDKWVEEYVKPPEGGSSVSIITRIEVTPSQTLG-----TCPESMRVHSS 123
schis -----VGFTNFSHIPGIGRSWDVADYIVPPLGNALFVITNLVKTERQSLS-----KCQESSWPEAA 125
choan IAITCDNTDISSMNDCVPGDLRWDPDYIKPAQENDAFFVSNSIQTSKQTQRAEGWDEDPAA--PVTGSSAFN 149
algae -----QATLRDATTGATLVDANDLVMYEPSG---FFITALATTLQARG-----RCPGMDEDET 112
dicty -----PNTFASNLTYCSNQQHNGSTYPPTPLECNY-WDEQLALFPVGQDSTF-----TCTTRVRLSKQEAN 128

rP2X2 CHSDD-DCIAGQLDMQNGIRTGHCVPYYHG----DSKTCEVSAWCPVEDG---TSDNH-----FLGKMAPNFT 184
schis CYKDS-DCKPYFISHLGNAHTGKCIIKPGN----DIGSCEIYSWCPLENDTLPLGRKS-----FLFPMVYNT 189
choan CTSDA-DCPR--FATSRNGALTGECNT-----TTERCRIYGWGPVESKDEDRATDGLFYARHMPAVKNFT 213
algae CTDAS-ACVVGTFSPS--GRMTGQCVATALKDEDGKVKRCEVEGWCPGEPEKDEVTLEN-----VGNFT 175
dicty CNFTDPTCKFVDEPGSAKN-----IYIADIESFT 132

rP2X2 LILIKNSIHYPKFKFSKGNIASQKSDY-LKHCTFDQDS--DPYCPIFRLGFIVEKAGEN-----FTEL 243
schis LLLIKNDINFEKFGIHRRNIQNWASKKFLRTCLYNKTDPENRFCPIFQFGTIFEANVD-----QSIF 251
choan VYIKNTVFFQRFGAKFG-STDESDKVDVYTCTWSPTG-LERHCPIFKIDTILNEAGIT-----DFENQA 275
algae VFTRISVEFPGIPDEDEGGNMLWTNLNGTKP-TLGWNLWTINDLESGGMSVKEVARK-VTPYTLKEVARKGWDG 236
dicty LILIDHTMYASSSSGSQFNAVDLHGYILNQDGDEVQIDANGTSIGVSGKPDIMTIGQLLSFGGVSLDQASPV--LLDQA 224

rP2X2 AHKGGVIGIINWNCDLDLSESECNPKYSFRRLDPKYVPAS-SGYNFRFAKYYKINGTTT----- 303
schis IS-GGVIGIDIDWKCDLDWDVQYCNPTYSFRRLDDAHAKIA-SGFNFRYAHFYSENGTNY----- 309
choan MRNGALITIQVNYDCNLDSSAHTCSPTYKFTRLD-TKSDLS-AGYNFRFANYQIDPPA----- 331
algae RVDVEF-----DCNLDRGIDACAPKTPYTLKQVMHPNTLSEGFNIRWISGQNVGEPSAQAGVVYSNETANGPGKDV 307
dicty SPVDSNVSIR--YDGVLFVFITYSNTYTYSTSDFKYVYSVQQIANTIYDVPETIIESIHS----- 284
```

Pre-TM2, TM2, post-TM2/C terminus

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rP2X2 RTLIKAYGIRIDVIVHGQAGKFSLIPTIINLALTSIGVGSFLCDWILLTFMN-KNKLYSHKFDKVRTPKHS 377
schis RDLIKAYGIRFVIHVSGEAGKFHLLPLTMNIGSGLALGLAPTVCDIIALNLLR-SRDIYQRAKFETIAEEQAHL 383
choan RDLYKVYGLRFVVVSGTAGRFSMVPLLVALGSGLGLGLATVIADLLVTKCIR-NANVYYGLKYQVVDEIDR 405
algae RLLVKGYGPRIRFEVTGVGRKFDWLTLSTTVGAGVAFLGIASLVVNAVMYCSGPKSKQYESWLFAEFHDTPYGS 382
dicty RLLYKRHGIRVIFIQTGTIGSFHFQTLLLTLVSGLLAVATTVVDQLAIRLLP-QRKSYSSLKFQVTESMSNPM 358
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Figure 4

Conservation of P2X receptors among disparate organisms. Sequence alignment of five P2X receptors that form functional ATP-gated channels when expressed in HEK cells. Overlines denote proposed transmembrane domains. Red denotes residues conserved among five sequences; blue denotes residues similar among five sequences (similar are IVLM, FYWH, KRH, EQDN). Sequences (with Genbank accessions) are rP2X2, *Rattus norvegicus* P2X₂ (P49653); schis, *Schistosoma mansoni* (CAH04147); choan, *Monosiga brevicollis* (EDQ92249); algae, *Ostreococcus tauri* (CAL54489); dicty, *Dictyostelium discoideum* (XP.645378). Further C termini (5–95 residues) are divergent and not shown.

residues have previously been shown to be essential for function in mammalian receptors (21, 135). Second, the highly conserved residues are in or close to the two transmembrane domains. This applies to the highly conserved intracellular Y-X-X-X-K motif that is found both ~10 residues before the start of transmembrane domain 1 (TM1) and ~10 residues after the end of transmembrane domain 2 (TM2). Such a positioning pattern also applies to the highly con-

served extracellular positively charged amino acids that are located ~15 residues after the end of TM1 (Lys⁶⁹ and Lys⁷¹ in P2X₂) as well as ~20 residues before the start of TM2 (Arg³⁰⁴ and Lys³⁰⁸). Third, the great bulk of the ectodomain is not conserved. This includes the ten cysteine residues. The lack of conservation extends also to several residues that, on the basis of mutagenesis in vertebrate receptors, had been considered to contribute to an ATP-binding site

(see 135). The key residues that had been proposed to represent homology with the ATP-binding site of class II tRNA synthases (139) are also mostly missing in the simpler P2X receptors.

N Terminus

The most striking feature of the N terminus of the P2X receptor family is the Y-X-T-X[KR] motif that ends some 10 residues prior to the start of TM1. The T-X-K conforms to the protein kinase C consensus, and phosphorylation of Thr¹⁸ in the P2X₂ receptor has been demonstrated for P2X₂ (140) but not for P2X₁ receptors (141); mutations at this site alter peak currents and desensitization at the P2X₁ receptor (141). In the P2X₂ receptor, replacement of Tyr¹⁶ by cysteine prevents channel function, although such a substitution is tolerated at Thr¹⁸ or Lys²⁰ (142).

Transmembrane Domain 1

Each residue of TM1 has been systematically substituted by cysteine [P2X₂ (142–144)], alanine [P2X₂ (145, 146)], or tryptophan [P2X₄ (147)]. The results indicate in general that (a) TM1 is very likely α -helical, (b) residues Gly³⁰ and Tyr⁴³ (P2X₂ numbering) are completely conserved and critical for function, (c) TM1 likely moves outward with respect to TM2 during channel gating (148), and (d) residues in TM1 seem not to contribute directly to the permeation pathway (146).

Post-Transmembrane Domain 1

The 20 amino acids immediately following TM1 do not show strong conservation (Figure 4) except for Tyr⁵⁵ and Gln⁵⁶, which are essential for receptor function (142, 148), and the K Φ KG motif Lys⁶⁹ and Lys⁷¹ in the P2X₂ receptor, where Φ is any hydrophobic amino acid (142). Abundant evidence now supports the view that one or both of these Lys residues play a critical role in ATP binding. The three main lines of evidence are as follows.

(a) Changing the charge on the nearby Ile⁶⁷ can affect ATP potency in a manner that is consistent with a direct electrostatic effect on the binding of a negatively charged agonist (142), (b) comparing the effects of ATP with those of the partial agonist BzATP suggest a direct binding rather than a gating role (149), and (c) Lys⁶⁹ is also required for channel activation by TNP-ATP at the constitutively active P2X₂ receptor carrying a T339S mutation (150). Experiments with coexpressed P2X₂ subunits, in which one contains Lys⁶⁹ and the other does not (i.e., P2X₂[K69A]), strongly suggest that two rather than three such lysines are required for agonist activation of the channel and, moreover, that the ATP-binding site likely forms between lysines provided by two different subunits (151).

Ectodomain

Figure 4 illustrates that most parts of the ectodomain that are conserved among vertebrates are not found in the *Dictyostelium* sequence; this includes the ten cysteine residues. This indicates that much of the ectodomain is not critical for function as an ATP-gated channel and presumably subserves other roles such as interactions with proteins provided by the matrix or by neighboring cells. The pattern of disulfide bond formation has been deduced. However, although the cysteines play a role in transport to the cell membrane, none of the cysteines are critical for channel function (152, 153). Zinc potentiates currents evoked by ATP at P2X₂ receptors. By using concatenated cDNAs to express subunits joined in tandem, Hume and colleagues (154) showed that His¹²⁰ and His²¹³ form the zinc-binding site and that different subunits contribute to the compositions of these two histidines. When these histidines are replaced by cysteine, an intersubunit disulfide bond can form. Similar intersubunit ectopic disulfide bonds between pairs of cysteines introduced into the P2X₁ receptor (Lys⁶⁸ and Phe²⁹¹) have been demonstrated (155), as was first shown for cysteines at the outer ends of the transmembrane domains of the P2X₂ receptor [Val⁴⁸ to Ile³²⁸ (148)]. One conserved motif

is present in all known sequences, including the *Dictyostelium* sequence: [EQP]-[NDS]-F-T-Φ-Φ-Φ-[KD]-NH]-[STN] (where Φ = any hydrophobic residue). The FT (Phe¹⁸³-Thr¹⁸⁴ in P2X₂) residues may play a role in stacking against the adenine moiety of ATP (149, 156). A similar role has also been ascribed to the NFR motif (Asn²⁸⁸-Phe²⁸⁹-Arg²⁹⁰ in P2X₂) (156), but this seems less likely given that these residues are missing from the *Dictyostelium* sequence (137).

Pre-Transmembrane Domain 2

This region extends from the highly conserved Arg³⁰⁴ (P2X₂) to the beginning of TM2 (around Ile³²⁸). The first half of this region shows particularly high sequence conservation (**Figure 4**). The lysine residue at position 308 (in P2X₂; position 309 in P2X₁ and position 313 in P2X₄) is required for channel function and has been implicated in ATP binding (21, 135, 142, 149, 157). For the P2X_{2/3} heteromer, Wilkinson et al. (158) suggested that the ATP-binding site may have contributions from Lys⁶⁹ of one subunit and Lys³⁰⁸ of another. Other work suggests a more fundamental role for Lys³⁰⁸ in channel gating. Yan et al. (157) favor a role for this residue (Lys³¹³ in P2X₄ receptor) in contributing to intersubunit interactions that accompany gating, rather than contributing directly to binding. Cao et al. (150) have reported that this residue is critical for channel activation in a slightly modified channel (P2X₂[T339S]) that opens spontaneously in the complete absence of any agonist. P2X₂[T339S] is also activated by suramin, but P2X₂[T339S,K308A] is not; this finding is also best interpreted to indicate a key role for Lys³⁰⁸ in channel gating. With the exception of Arg³⁰⁴ and Lys³⁰⁸, substitution of any individual residue in pre-TM2 by cysteine (in P2X₁) does not markedly affect the ability of ATP to evoke inward currents (159). In most cases, these cysteines were accessible to labeling with MTSEA-biotin, suggesting their exposure on the protein surface. Yan et al. (157) studied alanine substitutions (in P2X₄) and have speculated that the pre-

TM2 segment functions as a “signal transduction module” from ATP-binding site to channel gate. Taken together, it appears that this highly conserved region may subserve roles other than ATP binding and channel gating.

Transmembrane Domain 2

The consensus view from cysteine scanning (21, 135), alanine scanning (145), and tryptophan scanning (147) is that TM2 does not cross the membrane as an uninterrupted α-helix. Cysteine is tolerated at almost all positions through TM2 when substituted individually (160, 161). Tryptophan is tolerated in the outer half of TM2, and the positions causing the greatest perturbation of function (actually an increase in sensitivity to ATP) align along one surface of a helix (P2X₄) (147). Near the center of TM2, tryptophan is not tolerated at Ala³⁴⁹, Leu³⁵², Cys³⁵³, Asp³⁵⁴, Val³⁵⁷, and Leu³⁵⁸ (P2X₄; positions correspond to P2X₂ Gly³⁴⁴, Leu³⁴⁷, Cys³⁴⁸, Asp³⁴⁹, Leu³⁵², and Leu³⁵³, respectively), suggesting to these authors (147) “a tightly packed and relatively delicate structure that poorly tolerates substitution” at this position in the protein.

Early studies with cysteine scanning and the effects of methanethiosulfonates had indicated that the gate might form in the middle of TM2 [between Thr³³⁹ and Asp³⁴⁹ on the P2X₂ receptor (160)], and studies on the permeability of the P2X receptor now support this interpretation. Migita et al. (162) identified Thr³³⁹ and Ser³⁴⁰ of the P2X₂ receptor as having the greatest effect on cation permeability. More recently, the fraction of the inward current carried by calcium ions (*Pf*^{Ca}) has been determined by combination of measurement of charge with measurement of change in calcium concentration (163). Measurements of *Pf*^{Ca} also implicate residues Thr³³⁹ and Ser³⁴⁰ as the main deep determinants of calcium flux. Calcium *Pf*^{Ca} varies among homomeric P2X receptors; P2X₁ and P2X₄ receptors show the highest values [11–15%, similar to calcium *Pf*^{Ca} for NMDA receptors; see Egan & Khakh (163)]. The difference between these two receptors and

others in the family may result in part from the calcium-concentrating effect of fixed negative charges provided by glutamate residues near the extracellular ends of both TM1 and TM2 (164).

The Asp residue at position 349 (P2X₂) is completely conserved among P2X receptors other than in the algae *O. tauri* (138). Replacement of Asn by Asp in the algae receptor causes a 50% increase in relative calcium permeability. However, the reverse mutation in P2X₂ receptors does not change relative calcium permeability (138, 162). Thus, this aspartate residue does not appear to play a major role in the determination of calcium permeability. In contrast, it does have a key role in the intramembrane interhelix interactions required for channel assembly (P2X₅) (165).

C Terminus

The cytoplasmic tails of six of the P2X receptors retain considerable sequence relatedness for the first 25 amino acids (**Figure 4**), but relatedness thereafter disappears completely. A YXXXX motif, beginning ~10 residues from the cytoplasmic end on TM2 (or 28 in P2X₇; see below), is very highly conserved among all P2X receptors, including those from single-celled organisms that have poorly conserved ectodomains. This lends a symmetry to P2X receptor sequences, the conserved region of which both begins and ends with YXXXX motifs in the cytoplasmic regions (**Figure 4**). Rassendren and colleagues (166) have shown that the Tyr and the Lys of this motif are required to stabilize the P2X receptor subunit in the plasma membrane; in heteromeric P2X_{2/3} receptors the motif needs to be present on only one of the partners. The P2X₇ receptor differs from the other six in this region by the presence of an 18-amino-acid cysteine-rich insertion. Removal of this insertion prevents the P2X₇ receptor from undergoing the increase in NMDG permeability that normally occurs when the ATP application is continued for several seconds. Deletion of this cysteine-rich segment has no effect on the properties of the channel in normal extra-

cellular sodium concentrations, and it does not affect the ability of the P2X₇ receptor to drive uptake of fluorescent dyes (167).

To the C-terminal side of the YXXXX motif, the sequences of P2X receptors are subunit specific. For the P2X₂ subunit, the C terminus binds to Fe65 (168), a multidomain protein first identified by its ability to bind to β -amyloid precursor protein. Both proteins colocalize to the postsynaptic membranes of excitatory synapses in the hippocampus. Coexpression in oocytes showed that the presence of Fe65 slows the rate of pore dilation (increase in NMDG permeability) that occurs when the ATP application is applied for several minutes (see Reference 21). P2X₄ subunits have a motif (YXXG Φ) that directly binds the μ 2 subunit of the AP2 adaptor protein (73). This interaction is responsible for P2X₄ receptor endocytosis, despite the presence of a nearby YXX Φ motif that is commonly involved in the binding of AP2 to other proteins.

P2X₇ receptors interact with several partners, including cytoskeletal proteins (85). One of the interactors is receptor protein tyrosine phosphatase β , which has a clear effect on the currents through the P2X₇ receptor when it is activated repeatedly (85). Other partners include epithelial membrane proteins (169). Motifs have been identified in conserved regions of the P2X₇ receptor C terminus that resemble sequences in the SH3 domain-binding proteins *Mycoplasma genitalium* cytoadherence accessory protein HMW3 and *Caenorhabditis elegans* protein C18H2.1 (170). However, the most intriguing binding partner may be the bacterial endotoxin lipopolysaccharide (LPS) (170). A motif very close to the tail of the P2X₇ sequence (residues 573 to 590) is well conserved among all known P2X₇ subunits and closely resembles a sequence in soluble LPS-binding protein. Bertics and his colleagues showed that a peptide corresponding to this motif (rat P2X₇ 573 to 590: CRWRIRKEFPKTQGQYSG) directly binds FITC-labeled LPS, as does the LPS-binding protein itself. The same two basic amino acids in the P2X₇

peptide and in LPS-binding protein are essential for LPS binding (170), and mutation of these two residues impairs both receptor trafficking and channel function in HEK293 cells (171). It will be important to test the possibility that this region of the P2X₇ receptor is directly responsible for those actions of bacterial LPS that are independent of Toll-like receptor 4 (TLR4).

CONCLUDING THOUGHTS AND FUTURE ISSUES

The field continues to be limited by the availability of agonists, antagonists, and modulators with high selectivity for one or another P2X receptor, but there have been notable advances. Particularly in the cases of P2X₁, P2X₃, and P2X₇ receptors, there are now antagonists available with nanomolar affinity. There remain two important caveats to their application. The first is the issue of species specificity, which seems particularly important to take into account for P2X₇ antagonists. The second is the problem associated with actions at receptors other than P2X receptors. For example, suramin analogs and ivermectin have several such effects. Physiologists studying the functional roles of P2X receptors would welcome

further progress in the development of such compounds.

The availability of mice lacking P2X₁, P2X₂, P2X₃, P2X₄, and P2X₇ receptors has impacted the discovery of new functional roles. The difficulties here are those familiar to all such approaches—the species specificity and the possibilities of compensatory changes. Tissue-specific and inducible knock-in and knockout constructs would be most valuable.

At the molecular level, progress continues to be made with respect to our understanding of the permeation pathway and the ATP-binding site. The former will benefit from single-channel recordings combined with mutagenesis, and the latter from the sequencing of receptor fragments cross-linked to appropriate agonists or antagonists. Similar mass spectroscopic approaches need to be further extended to identify the range of proteins that interact directly with P2X receptors. These will lead us to an improved understanding of the subcellular trafficking and localization of P2X receptors, in which context it will be important to seek further evidence for intracellular functional roles such as those reported for *Dictyostelium*. And finally, high-resolution structure of a crystallized receptor would of course provide the field with its greatest impetus.

LPS: bacterial lipopolysaccharide

SUMMARY POINTS

1. P2X₁ receptors are neurotransmitter receptors at sympathetically innervated smooth muscle and play functional roles on platelets and in the juxtaglomerular apparatus of the kidney.
2. P2X₂ and P2X₃ subunits (and P2X_{2/3} heteromeric receptors) are necessary for the initiation of sensory signaling in pathways subserving taste, chemoreception, visceral distension, and neuropathic pain.
3. P2X₄ receptors are involved in the function of vascular endothelium. Mice lacking the P2X₄ receptor gene are hypertensive and have smaller-diameter arteries. Microglial P2X₄ receptors in the spinal cord have also been implicated in neuropathic pain.
4. Activation of P2X₇ receptors drives the release of proinflammatory cytokines from macrophages primed with lipopolysaccharides. P2X₇ antagonists appear beneficial in the inflammation of rheumatoid arthritis.
5. In the central nervous system there are widespread presynaptic and postsynaptic actions of ATP, but the importance of P2X signaling remains unclear at the cellular level.

6. P2X receptors in a simple eukaryote are intracellular and involved in osmoregulation. Thus, functions on intracellular membranes in mammalian cells may be worth seeking. The distant sequence relatedness of these primitive P2X receptors also indicates that substantial parts of the receptor ectodomain are not needed for receptor function.
7. Single-channel recordings and mutagenesis are identifying the regions of the P2X receptor involved in ion permeation, ATP binding, and trafficking to the cell surface, but higher-resolution structural information would greatly accelerate this work.

DISCLOSURE STATEMENT

The authors are not aware of any biases that might be perceived as affecting the objectivity of this review.

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