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Neuromodulation by extracellular ATP and P2X receptors in the CNS

Baljit S. Khakh¹ and R. Alan North²

¹Departments of Physiology and Neurobiology, David Geffen School of Medicine, University of California Los Angeles CA 90095-1751, USA

²Faculty of Medical and Human Sciences, University of Manchester, Michael Smith Building, Oxford Road, Manchester, M13 9PT, UK

Abstract

Extracellular adenosine 5' triphosphate (ATP) is a widespread cell-to-cell signaling molecule in the brain, where it activates cell surface P2X and P2Y receptors. P2X receptors define a protein family unlike other neurotransmitter-gated ion channels in terms of sequence, subunit topology, assembly and architecture. Within milliseconds of binding ATP, they catalyze the opening of a cation-selective pore. However, recent data show that P2X receptors often underlie neuromodulatory responses on slower time scales of seconds or longer. Herein, we review these findings at molecular, cellular and systems levels. We propose that, while P2X receptors are fast ligand-gated cation channels, they are most adept at mediating slow neuromodulatory functions that are more widespread and more physiologically utilized than fast ATP synaptic transmission in the CNS.

Keywords

purinergic; glia; synapse; signaling; neuron; secretion; pain; drug; structure; biophysics; astrocyte; channel

The discovery of ATP signaling

The field of ATP signaling has taken nearly a century to evolve since the discovery of ATP in 1929 (Khakh and Burnstock, 2009). The first evidence for the release of ATP from sensory nerves was provided by Pamela Holton in the 1950s (Holton and Holton, 1954; Holton, 1959). In 1972, Geoffrey Burnstock proposed the existence of “purinergic nerves” (Burnstock, 1972), laying the foundations of a new field. Forty years later, there is little doubt that extracellular ATP signaling is widely utilized in cell sensing.

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Correspondence: BSK (bkhakh@mednet.ucla.edu) and RAN (r.a.north@manchester.ac.uk).

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Much of the early data on purinergic nerves derived from physiological experiments on smooth muscle preparations. By the early 1990s, however, ATP was also shown to depolarize neurons (Jahr and Jessell, 1983; Krishtal et al., 1983), to open single ion channels in patches of membrane (Benham and Tsien, 1987; Kolb and Wakelam, 1983) and to mediate fast synaptic transmission at synapses in the peripheral and central nervous systems (Edwards et al., 1992; Evans et al., 1992; Silinsky et al., 1992). By this time, ATP receptors had also been subdivided as belonging to two major families: metabotropic P2Y receptors and ionotropic P2X receptors. With the subsequent cloning of the genes encoding P2X receptors came a new era. In this review, we focus on P2X receptor mediated ATP signaling in the brain, discussing general themes pertinent to mechanisms and neuromodulation at the molecular, cellular, systems and disease levels.

Cell surface receptors for ATP and its metabolites

ATP activates a family of metabotropic P2Y and ionotropic P2X receptors. Collectively, the actions of ATP and its breakdown products produce responses that last from milliseconds to minutes, and even longer time scales through changes in gene regulation via second messengers. Signaling diversity is further increased by the fact that ATP receptors display a very broad range of ATP sensitivities, ranging from nanomolar in the case of P2Y receptors, to hundreds of micromolar for P2X7 receptors (Surprenant et al., 1996). Thus, ATP receptors respond over remarkably broad spatiotemporal scales, and ATP signaling is very dynamic.

The first P2X receptor genes were identified in 1994 (Brake et al., 1994; Valera et al., 1994) and within two years the whole family had been identified (Buell et al., 1996; Chen et al., 1995; Collo et al., 1996; Le et al., 1997; Lewis et al., 1995; Soto et al., 1996; Surprenant et al., 1996). This was an exciting period that culminated with the realization that P2X receptors defined a unique protein family. Each of the homomeric P2X receptors also displays distinct functional properties (Fig. 1, Table 1). The available data on the subunit composition and properties of heteromeric P2X receptors (Coddou et al., 2011) are not considered in depth here.

P2X receptors are non-selective cation channels with high Ca^{2+} permeability that carry a depolarizing current under standard physiological conditions. In some cells, P2X channels are also significantly permeable to anions. For example, the full length P2X5 receptor is permeable to Cl^- (North, 2002). This remains the exception rather than the rule. Thus at its most fundamental level, P2X receptor mediated neuromodulation starts with chemistry: ATP rapidly gates P2X receptor pores, triggering transmembrane fluxes of selected ions.

The seven mammalian P2X subunits range from 379 (P2X6) to 595 (P2X7) amino acids in length. Each subunit contains two hydrophobic membrane-spanning segments (TM1 and TM2) separated by an ectodomain which contains ten conserved cysteine residues that form disulfide bonds. Representing the simplest known architecture for ligand-gated ion channels, P2X receptors adopt a relatively simple fold, with intracellular N and C termini and most of the molecule forming an extracellular loop. Multiple lines of evidence have established that the channels are trimeric (Barrera et al., 2005; Nicke et al., 1998; Stoop et al., 1999). This

contrasts with eukaryotic glutamate-gated cation channels, which form as tetramers, and with the large family of pentameric "Cys-loop" receptors which includes members gated by acetylcholine, glycine, γ -aminobutyric acid, 5-hydroxytryptamine and glutamic acid. TM1 and TM2 both contribute to the function of the P2X pore, but the major pore lining segment is TM2 (Egan et al., 1998; Haines et al., 2001; Jiang et al., 2001; Kracun et al., 2010; Li et al., 2008; Rassendren et al., 1997; Samways et al., 2008).

Molecular Mechanisms

Flukes, fins and beaks

The crystal structure of the zebrafish P2X₄ receptor has provided a detailed picture of the atomic anatomy of the receptor (Fig. 3A) (Hattori and Gouaux, 2012; Kawate et al., 2009). In overall shape, a single P2X receptor subunit resembles a dolphin rising from the ocean surface (the cell membrane). The narrow distal part of the dolphin (the fluke) is formed from the two membrane spanning domains as they run through the cell membrane. The body region, with its attached fins and flippers, rises and curves over so that the rostral region corresponding to the head and beak run almost parallel to the membrane surface. The three subunits curl around each other on an axis of symmetry projecting as a perpendicular from the cell membrane, enclosing a central space or cavity.

The tip of the P2X receptor stands some 70 Å proud of the cell membrane, its turret formed by loops from each subunit surrounding a central aperture that is too narrow for hydrated ions to pass (Fig. 3). This turret is formed by the upper hairpin of a two-stranded beta sheet, the three copies of which form the wall of a slightly widened central cavity as they pass down toward the cell membrane (the upper vestibule).

Extending laterally from the upper body is the head region, a relatively poorly conserved tangle of loops and short β -strands that is stabilized by three disulfide bonds in mammalian receptors. Three prolines cluster at the central axis, corresponding to P89 of the rat P2X₂ receptor, which is highly conserved (all numbering refers to rat P2X₂). The surrounding region appears to stabilize the upper body as a "brace" with respect to the movements of the lower body that occur during channel opening. Below P89, the central vestibule widens again, its wall formed by three-stranded β -sheets (Fig. 3B). This forms the lower body region, and amino acid side chains projecting into the central vestibule give it a strongly acidic surface. The β -sheets extend down to join directly to the outer ends of the transmembrane domains and, as they do, lateral portals open between them just above the level of the outer membrane surface (Fig. 3A).

Where ATP binds

Facing outwards from the β -sheets that form the wall of the lower body, some 45 Å from the cell surface, one finds the key residues involved in ATP binding. The triphosphate chain of the ATP bends in a U-shape around the ammonium nitrogen atom of K69, and oxygen atoms of the α , β and γ phosphates make several interactions with K69, K70, N288, R290 and K308 (Hattori and Gouaux, 2012) (Fig. 3D). At least one oxygen atom of the γ phosphate is exposed to surrounding solution. The ribose ring of ATP likely makes hydrophobic contact with L211 and I226 (Fig. 3D), leaving its 2' and 3' oxygen atoms

exposed to the extracellular solution. The adenine moiety lies deeper in the binding cavity, its side-chain nitrogen hydrogen bonding to backbone carbonyl oxygens of K70 and T184, and its main ring nitrogen atom hydrogen bonding with the side-chain oxygen of T184. These interactions account for the observed selectivity of P2X receptors for ATP, rather than CTP, UTP or GTP (Hattori and Gouaux, 2012).

How ions permeate

The lateral portals between the β -strands of the lower body provide the access route for ions into the central vestibule of the protein (Hattori and Gouaux, 2012; Kawate et al., 2011; Samways et al., 2011). These portals are situated just above the outer membrane surface, ringed by polar side-chains of Q52 and Q321 (A chain), and I328, D57, E59, and K195 (B chain) (Fig. 3E). They are likely partially occluded by lipid at their lower edge. They are sufficiently wide in the closed channel to allow passage of hydrated ions, and they enlarge markedly with channel opening (Fig. 3E). Ions passing through these portals enter the central vestibule, where the excess of negative charges on its inner wall probably serves to concentrate cations. A role in ion selectivity for the portals themselves is indicated by the observation that mutation of some of their negatively charged side-chains changes the relative permeability of calcium (Samways et al., 2012; Samways and Egan, 2007).

The permeation pathway through the cell membrane forms down the central axis of the three tilted TM2 domains. They are widely separated at the outer membrane surface (Fig. 3B), but the extracellular vestibule formed within them narrows as the TM2s approach and cross each other. In the closed structure, the first narrowing occurs at the level of T336, but the narrowest part of the transmembrane pore is formed one helical turn further along TM2 at T339 (Fig. 3A,B). At this point, side-chains occlude the channel when closed, and project into it when open. When T339 is replaced by lysine in one, two or three of the subunits, the unitary conductance is reduced stepwise, and the relative chloride permeability is similarly increased, suggesting a critical interaction with permeating ions (Browne et al., 2011). Thus, the side chains of T339 which occlude the permeation pathway in the closed channel (Fig. 3B) also form the narrowest part of the constriction of the open channel (Hattori and Gouaux, 2012). In the P2X2 receptor, the side chain of the next residue along the helix (S340) also becomes accessible to the permeating ions as the helices separate during channel opening: positively-charged substitutions here markedly increase outward currents (Cao et al., 2009), probably as a result of increasing permeability to chloride ions (Browne et al., 2010).

Passing further into the cell, the splaying TM2 domains enclose a widening intracellular vestibule. A conserved glycine residue situated one helical turn interior to T339 (G342) allows TM2 to kink during channel opening (Fujiwara et al., 2009; Hattori and Gouaux, 2012). The importance of this helix bend is unclear, but it is known that addition of a side chain at this position by mutagenesis does not prevent the response to ATP (Cao et al., 2009; North, 2002). Indeed, in the P2X4 receptor, the identity of such a side-chain profoundly alters kinetic behavior (Khakh et al., 1999a). The side-chain at this position would not protrude towards the permeation pathway of the closed channel, and these effects likely result from stabilizing one of the two open conformations.

The side chain of the V343 is orientated toward the center of the permeation pathway, and this is consistent with the observation that a high affinity metal binding site may be formed here by cysteine substitution (Li et al., 2010), although the cadmium block reversed with washout. Cadmium also has some blocking effect at S345 and D349, but the further splaying of the helices as one passes into the cell makes it unlikely that this results from stable coordination within the central axis of the pore (Kracun et al., 2010). At S345 C β radii are 12 Å and 11 Å for open and closed channels and at D349 the radii are 10 Å (open) and 16 Å (closed) (Fig. 3B). The electronegativity at D349 may serve a "cation concentrating" role in the inner vestibule (Cao et al. 2009).

From closed to open

ATP binding initiates several substantial rearrangements (Jiang et al., 2012; Lörinczi et al., 2012; Roberts et al., 2012). The first step may be the disruption of the strong repulsion between the quaternary nitrogen atoms of K308 (on β 14 of chain A) and K69 (on β 1 of chain B) as the highly negative phosphate tail of ATP is drawn into the binding cleft. These nitrogen atoms are 8 Å apart in the closed structure, and approach to 4 Å in the open structure when they interact with negative oxygen atoms on the phosphate chain. R290 (on β 13 of chain A) forms a salt bridge with E167 (projecting down from the head domain of the same subunit) in the closed channel: this is exchanged for a salt bridge with the γ phosphate of ATP (Hausmann et al., 2012) (Fig. 3C). The lateral movement of Q138 allows the head domain to drop down. At the same time, S284 on the left flipper retracts clear of the binding cleft. On the B chain an upward movement of the dorsal fin allows it to make hydrophobic interactions with the ribose moiety and adenine base, principally through L211 and I226 (Fig. 3E). Of interest, the dorsal fin of the P2X7 receptor lacks one helical turn, which includes the first of these two hydrophobic residues: this might contribute to the relative insensitivity to ATP shown by this subtype.

The downward movement of the head domain can be followed by the approach of H120 to H213. Nagaya et al. showed that these residues contributed to an intersubunit zinc binding site (Nagaya et al., 2005) and that, when H120 and H213 are both replaced by cysteine, a new intersubunit disulfide can form which inhibits channel opening. The intimate approximation of the head domain (chain A) with the dorsal fin (chain B) appears to be in itself sufficient to lead to channel opening in a suitably mutated "reporter" receptor (Jiang et al., 2012).

The movements of the left flipper (chain A) and dorsal fin (chain B) exert tension on the β -sheeted wall of the lower body of the same subunit, causing it to flex outwards by a slight separation of its constituent β -sheets and increasing its circumference by a progressively greater amount as it passes down toward the outer surface of the membrane (Roberts et al., 2012). Here, at the outer end of TM2 the "diameter" increases from 18 to 32 Å (Fig. 3C). This widening of the lower body causes the transmembrane helices to separate at their outer ends, and expands the pore like a three-leafed iris (Fig. 3F).

Caveats and limitations

The foregoing interpretation and description gives a compelling account of how a P2X receptor can bind ATP and transform from a closed to an open state. Yet there are caveats and limitations. The first is that the determination of a single open structure does not preclude the existence of other stable forms, and obviously it does not address the movements that occur in the most flexible parts of the protein. Studies using normal mode analysis and molecular dynamic simulations (Du et al., 2012; Jiang et al., 2012) are likely to be most informative in providing insight into conformational dynamics. Additionally, the zebrafish P2X4 receptor used for crystallography lacked intracellular N and C termini, which will likely have an effect on channel properties. Likewise, the membrane proximal regions of these N- and C-terminal domains contain conserved motifs in which minor substitutions can impair receptor function (North, 2002). There is also evidence that these membrane proximal intracellular regions are involved in desensitization (Werner et al., 1996), and the molecular rearrangements underlying desensitization remain unclear. Further, P2X receptors have three ATP binding sites (Bean et al., 1990), and the present structures provide little insight into the molecular basis of the observed cooperativity (Ding and Sachs, 1999; North, 2002). Finally, the stationary snapshots and nanosecond simulations leave us with much to learn about the kinetics of receptor activation, and how these proteins may be better suited to respond to longer-lasting diffusing signals than to the short, sharp pulses of a fast transmitter.

Drugs and allosteric modulators

There has been considerable activity in the drug development world focused on P2X receptors, and this has been well reviewed (Coddou et al., 2011; Ford, 2012). The ATP-bound structure of Hattori and Gouaux (2012) shows clearly how molecules such as TNP-ATP could be accommodated in the ATP binding pocket of the receptor. Wolf and colleagues (2011) have studied NF770, a derivative of suramin, which blocks P2X2 receptors at about 20 nM (Wolf et al., 2011). By homology modelling and docking, they demonstrated a direct hydrogen bond between R290 (critical for ATP binding, see Fig. 3D–F) and the methoxy oxygen atom of NF770, thus accounting for its high affinity as a competitive antagonist.

Ivermectin, strongly potentiates ATP-induced currents at P2X receptors, and this effect is largest for P2X4 receptors (Khakh et al., 1999b). Residues involved in ivermectin binding have been identified on both TMs as lipid-facing (Jelínková et al., 2008; Silberberg et al., 2007). The ATP-bound structure confirms the outward facing orientation of these residues around the helix crossing point, and substantiates the suggestion by Silberberg et al. (2007) that ivermectin interacts at the protein-lipid interface so as to stabilize an open state.

Divalent and trivalent cations have been used extensively to probe the ectodomain function of P2X receptors (Coddou et al., 2011). There are two salient areas. The first is that all P2X receptors are exquisitely sensitive to the concentrations of normal extracellular ions: all the receptors respond to lower ATP concentrations when the concentrations of extracellular calcium and magnesium are reduced, effects generally more marked for P2X7 receptors (Surprenant et al., 1996). Reduction of extracellular magnesium will increase the fraction of

ATP that is in the uncoordinated (ATP^{4-}) state, which is now known to be the active binding form, but this effect should be equal for all receptors. The importance of the divalent ions is particularly seen with P2X7 receptors, where the reduction in calcium and magnesium concentrations strongly promotes "dilatation" and divalent ions appear to serve as "co-agonists" (Jiang et al., 2005; Shinozaki et al., 2009; Surprenant et al., 1996). A similar behavior underlies the astonishing property of P2X3 receptors to "remember" for many minutes a brief exposure to a higher than normal concentration of calcium ions (Cook et al., 1998). This action was ascribed to an acceleration of recovery from a very long-lived desensitized state.

Gadolinium was present in the solution used to grow the first crystals of the zebrafish P2X4 receptor, and gadolinium (100 μM) completely inhibits ATP currents at those receptors (Kawate et al., 2009). X-rays showed it unequivocally to be present in the upper part of the central vestibule, coordinated by E98 from each of the three subunits, but also on the external surface of the body domain, one ion per subunit. In this latter position, the gadolinium is coordinated by carboxylates from D184 and N187. Any general significance of this finding seems unlikely: zfp2X4 D184 is not well conserved among P2X receptors, whereas the Asp at position 187 is conserved but glycosylated (Lenertz et al., 2010; Newbolt et al., 1998; Roberts and Evans, 2006).

Pore dilation

Although all functional P2X receptors undergo conformational changes that result in the opening of a cationic pore within milliseconds of ATP binding, some P2X receptors (notably P2X2, P2X4 and P2X7) also undergo additional slower conformational changes (Fig 4). Pore dilation follows several seconds of ATP activation and is characterized by increases in permeability to organic cations and several dyes (Khakh et al., 1999a; Khakh and Lester, 1999; North, 2002; Virginio et al., 1999). In other P2X receptors, notably P2X1 and P2X3, extended activation by ATP results in channel closure through desensitization. Pore dilation is of interest because it occurs over seconds, endowing P2X receptors with slow signaling capabilities and potentially providing the ability to release intracellular constituents such as ATP itself.

Two mechanisms have been proposed for pore dilation (Fig 4). For P2X2, P2X4 and P2X7 receptors, pore dilation appears to involve an intrinsic conformational change in the protein itself (Chaumont and Khakh, 2008; Khadra et al., 2012; Yan et al., 2010; Yan et al., 2011; Yan et al., 2008). However, for natively expressed P2X7 channels, an accessory protein may also be required, and pannexin-1 channels may be involved in receptor pore dilation (Jiang et al., 2005; Pelegriin and Surprenant, 2006; Pelegriin and Surprenant, 2007; Surprenant et al., 1996) in a manner that varies with the particular splice variant being studied (Xu et al., 2012). In all cases the dilated pore state is regulated by cellular processes and mechanisms that involve the C terminal tail. In the case of P2X4 receptors, fast scanning atomic force microscopy has been used to image a slow conformational change that may underlie the phenomenon within single protein molecules (Shinozaki et al., 2009).

Pore dilation may allow P2X receptors to function as intrinsic frequency detectors, by switching to the larger pore state with altered signaling upon repeated ATP activation

(Khakh et al., 1999a). Recent data suggest that this particular state of P2X7 receptors may be involved in susceptibility to chronic pain in rodents and humans (Sorge et al., 2012), raising the possibility that pore dilation of other P2X receptors in the brain may also mediate important slow responses. Further structural as well as physiological studies are needed to evaluate precisely how pore dilation and dynamic selectivity filters occur and what their functions are *in vivo*.

Functional interactions of P2X receptors with other ion channels

P2X and nicotinic receptors undergo functional interactions (Barajas-Lopez et al., 1998; Nakazawa, 1994; Nakazawa et al., 1991; Searl et al., 1998; Searl and Silinsky, 1998; Zhou and Galligan, 1998). Co-expression of P2X2 and $\alpha 3\beta 4$ nicotinic receptors in *Xenopus* oocytes provided evidence for functional interactions resulting in cross inhibition: the activation of one channel type affected distinct kinetic and conductance states of the other, and co-activation resulted in non-additive responses owing to inhibition of both channel types (Khakh et al., 2000). This study also showed that the functional interactions occurred in synaptically coupled myenteric neurons where nicotinic fast excitatory postsynaptic currents were occluded during activation of endogenously co-expressed P2X channels. Similar experiments have now been repeated with several ion channel combinations showing that cross inhibition between P2X receptors and members of the nicotinic receptor-like family are common (Barajas-Lopez et al., 1998; Barajas-Lopez et al., 2002; Boue-Grabot et al., 2003; Boue-Grabot et al., 2004a; Boue-Grabot et al., 2004b). Most recently, functional interactions have been reported for P2X receptors and acid sensing ion channels (ASIC) (Birdsong et al., 2010) as well as between P2X3 receptors and TRPV1 channels (Stanchev et al., 2009). We comment here on general themes that emerge.

Overall, the data suggest P2X receptors form molecular scale partnerships with distinct ion channels. Fluorescence resonance energy transfer (FRET) experiments show close interactions between P2X2 and $\alpha 4\beta 2$ nicotinic, P2X5 and ASIC, as well as P2X2 and GABA_A receptors, which provides a basis for functional interactions within the plasma membrane (Birdsong et al., 2010; Khakh et al., 2005; Shrivastava et al., 2011). Cross-inhibition between P2X receptors and nicotinic channels can occur in the absence of ion flow through P2X2 during a closed-desensitized state, and is likely due to conformational coupling (Khakh et al., 2000). Similarly, the interaction between P2X5 and ASIC channels is independent of ion flow through P2X5 receptors (Birdsong et al., 2010). In the case of spinal neurons, activation of P2X2 receptors increases the lateral mobility of GABA_A receptors, adding a previously unknown facet to the interactions between these receptor types (Shrivastava et al., 2011). An important question for future exploration is to determine if cross inhibition has behavioral consequences, either physiologically or during disease states, and it will also be important to nail down the molecular basis for the interactions.

Activation-dependent regulation of P2X receptors

P2X receptors are regulated in a use-dependent manner and it is likely these mechanisms contribute in important ways to their neuromodulatory responses in the brain. To date, two mechanisms have emerged: regulation of trafficking and regulation by Ca²⁺ sensors.

P2X4 receptors display several types of dynamic trafficking including endocytosis, lysosomal secretion and lateral mobility. A robust observation has been the role of trafficking of P2X4 receptors through dynamin-dependent endocytosis, and P2X4 receptors undergo constitutive and regulated endocytosis mediated by a novel non-canonical endocytic motif (YXXGL) (Bobanovic et al., 2002; Royle et al., 2002; Royle et al., 2005). Additionally, owing to their extensive glycosylation, P2X4 receptors are resistant to lysosomal degradation and can be secreted from lysosomes to deliver a pool of functional ion channels to the cell surface (Qureshi et al., 2007). Most recently, these insights have been extended to studies of P2X4 receptors within microglia, and it has been shown that both lysosomal secretion and plasma membrane lateral mobility of P2X4 receptors are increased by activation of the microglia (Toulme et al., 2010; Toulme and Khakh, 2012).

A second form of activity-dependent regulation has been demonstrated for P2X2 and P2X7 receptors and is mediated by the Ca^{2+} sensor proteins VILIP1 and calmodulin, respectively (Chaumont et al., 2008; Richler et al., 2011; Roger et al., 2008). In both cases, Ca^{2+} fluxes mediated by these P2X receptors result in the recruitment of the cognate Ca^{2+} sensor to the C-terminal domain of the channel to regulate functional responses. The consequences are subtle for P2X2 receptors, but result in profound facilitation of P2X7 receptor responses (Roger et al., 2008). The interaction with VILIP1, which occurs during endogenous ATP release, requires slow conformational changes resulting in exposure of a VILIP1 binding site in the cytosolic C terminal tail of P2X2 receptors (Chaumont et al., 2008; Chaumont and Khakh, 2008). Single-molecule experiments reveal that the interaction between P2X2 receptors and VILIP1 regulates plasma membrane lateral mobility of P2X receptors in neuronal dendrites (Richler et al., 2011), perhaps serving to affect recovery from desensitization by controlling the supply of receptors. Determining the full repertoire of proteins that interact with P2X2, P2X4 and P2X7 will help illuminate how these receptors are tuned to perform their tasks *in vivo*.

Single-molecule imaging experiments now provide accurate and consistent values for P2X2, P2X4 and P2X7 receptor diffusion coefficients in the plasma membrane (0.027, 0.023 and 0.021 $\mu\text{m}^2/\text{s}$, respectively) (Arizono et al., 2012; Richler et al., 2011; Toulme and Khakh, 2012). In the case of P2X2 and P2X4 receptors, activation by ATP causes the receptors to diffuse twice as fast in a cell- and subunit specific manner (Richler et al., 2011; Toulme and Khakh, 2012). Accurate P2X receptor diffusion coefficients will be invaluable in modeling receptor movement and plausible roles for lateral mobility in recovery from desensitization during physiological activation such as would occur during point source-like ATP release *in vivo*.

Cells and Mechanisms

Some general points

P2X receptors are often expressed at low levels, generally in specific compartments such as the edges of spines and within nerve terminals (Le et al., 1998; Rubio and Soto, 2001; Vulchanova et al., 1996), and are activated by quite high amounts of ATP. It seems that sufficient ATP to activate extrasynaptic P2X2 receptors is only released during bursts of action potentials (Richler et al., 2008), suggesting that P2X receptors underlie

neuromodulatory responses. Also, we are aware of no example in the brain or spinal cord where endogenous ATP release stimulates postsynaptic P2X receptors to trigger action potential firing (i.e. is a primary fast synaptic transmitter). In this respect, ATP shares features in common with other neuromodulators, such as ACh and serotonin, which also appear to mediate fast synaptic transmission only rarely in the central nervous system. In these settings, P2X receptor responses may require downstream signaling or protein interactions and not necessarily depolarization of the membrane. Indeed, P2X receptors carry significant Ca^{2+} fluxes at resting membrane potentials (Egan and Khakh, 2004).

Multiple P2X receptor knockout mice have been generated, all of which survive to adulthood (Chessell et al., 2005; Cockayne et al., 2005; Cockayne et al., 2000; Mulryan et al., 2000; Souslova et al., 2000; Ulmann et al., 2008). Few immediately obvious CNS phenotypes have been reported, yet these same mouse models show that P2X channels are strongly involved in a whole host of pathologies. Thus, it appears that endogenously released ATP does not generally affect the immediate integrative properties of neuronal circuits, but pathological alterations in signaling can have profound effects.

Fast ATP P2X receptor-mediated synaptic transmission in the CNS: small and underwhelming

The first evidence for fast ATP synaptic responses in the brain was provided in the medial habenula (Edwards et al., 1992; Edwards et al., 1997). Since then evidence for ATP as a synaptic transmitter has been provided in the locus coeruleus (Nieber et al., 1997; Silinsky et al., 1992), the hippocampus (Mori et al., 2001; Pankratov et al., 1998; Pankratov et al., 2002), in spinal neurons (Bardoni et al., 1997; Jo and Schlichter, 1999), hypothalamic neurons (Jo et al., 2011; Jo and Role, 2002) and cortex (Lalo et al., 2007; Pankratov et al., 2003; Pankratov et al., 2007). While these studies found evidence for ATP synaptic transmission, in all cases the interpretation that P2X receptors are involved is based on the use of P2X antagonists and agonists that are known to be imperfect in their selectivity (Khakh et al., 2001; North, 2002). Also, none of these studies employed P2X receptor subunit knock-out mice or provided a detailed pharmacological/biophysical characterization of the underlying P2X receptors (Table 1 shows the emerging useful pharmacopeia of P2X receptors). Additionally, ATP-mediated EPSCs detected in this manner tend to be small (about 10% of the size of EPSCs mediated by glutamate), infrequent, only observed in subpopulations of neurons within a given brain nucleus, and they generally require strong electrical stimulation to evoke. There is little evidence that the small EPSCs are physiologically effective in the neurons from which they were detected. Thus, the evidence in favor of ATP as an important synaptic neurotransmitter mediating fast synaptic potentials in the brain remains weak.

The evidence for ATP as a fast synaptic neurotransmitter with important functional roles is much stronger in the periphery, for example at neuro-effector junctions (Mulryan et al., 2000; Sneddon and Burnstock, 1984; Sneddon et al., 1982), neuro-neuronal synapses (Evans et al., 1992) and in the gastrointestinal system (Bian et al., 2003; Galligan and Bertrand, 1994). An important feature of these papers is that key experiments have been repeated in

P2X1 and P2X2 subunit knock-out mice, and the underlying receptors have been characterized in detail.

Neuromodulatory influences of postsynaptic P2X receptors: a diversity of mechanisms

Three noteworthy studies suggest that ATP signaling via postsynaptic P2X receptors plays a neuromodulatory role at brain synapses. The first concerns the role of P2X receptors in long-term synaptic potentiation (LTP) of glutamate synapses onto CA1 pyramidal neurons (Pankratov et al., 2002). In this case, Ca^{2+} flux through P2X receptors dampens NMDA receptor-dependent LTP at low frequencies of action potential firing in Schaffer collateral axons (Pankratov et al., 2002). Consequently, when P2X receptors are blocked, NMDA receptor dependent LTP occurs at lower action potential frequencies. Second, recent studies have utilized P2X4 receptor knock-out mice to analyze synaptic transmission and plasticity in CA1 pyramidal neurons, and no evidence for a role of P2X4 receptors in excitatory synaptic transmission was found (Baxter et al., 2011). Moreover, although the P2X4 deletion mice display subtly reduced LTP (Sim et al., 2006), ATP fast synaptic transmission does not seem to be the cause, and the data suggest that Ca^{2+} entry through P2X4 receptors may regulate NMDA receptor incorporation into fast synapses (Baxter et al., 2011). A third set of experiments suggest roles for P2X4 receptors in inhibitory synaptic transmission onto a specific population of steroidogenic factor 1 (SF-1) positive neurons in the ventromedial nucleus of the hypothalamus (Jo et al., 2011). Blocking P2X4 receptor endocytosis increases responses evoked by exogenous ATP in SF-1 neurons, but no evidence was found for fast ATP synaptic transmission. However, blocking P2X4 endocytosis reduced inhibitory IPSCs, which appears to be due to increased cross inhibition between P2X4 receptors and synaptic GABA_A receptors.

Demonstrating synaptic consequences for the interaction between P2X and other receptor classes is relevant to future efforts to explore the physiological roles of P2X receptors in the brain. From this perspective, the interactions between P2X5 and ASIC channels are particularly noteworthy (Birdsong et al., 2010), as they demonstrate strong functional interplay between the two ion channels in a manner that utilizes a P2X receptor subunit that is only weakly functional as a homomer (Collo et al., 1996). In this case, the interaction is independent of ion flow through the P2X receptor and dependent on a molecular interaction between the cognate subunits reminiscent of interactions between P2X and nicotinic receptors (Khakh et al., 2005). One should consider, therefore, the realistic possibility that seemingly silent P2X receptors in brain neurons may nonetheless be exerting important modulatory influences on other ion channels, particularly in cases such as ischemia when ATP is known to be released in high amounts (Birdsong et al., 2010).

Overall, recent studies that employ knock-out mice or other more specific molecular interventions suggest that P2X receptors modulate fast synaptic responses mediated by other more established fast neurotransmitters. Consistent with this suggestion, electron microscopy shows that P2X2 and P2X4 receptors are generally not located in synapses at sites directly opposite neurotransmitter release from the presynaptic terminal (Fig 5), residing instead at the edges of the postsynaptic density (Rubio and Soto, 2001). Recent single molecule imaging and tracking experiments in cell culture support these findings and

show that P2X2 receptors do not enter fast glutamatergic synapses even when heterologously over expressed and strongly activated by ATP (Richler et al., 2011). These data offer an explanation for the rarity of fast ATP neurotransmission in the brain and also raise important cell biological questions about why P2X receptors are excluded from synapses.

Presynaptic P2X receptors and the control of release probability

In general, activation of presynaptic P2X receptors increases neurotransmitter release probability due to influx of calcium (Fig 5), although depression of action potential evoked neurotransmitter release can also occur as a result of action potential failure and/or shunting in axons (Engelman and MacDermott, 2004; Khakh and Henderson, 2000). Presynaptic P2X receptors may be activated by endogenous ATP release in some synapses. Although, presynaptic P2X responses have now been described in many parts of the brain, we lack a satisfactory understanding of the precise physiological function of this form of presynaptic facilitation.

A presynaptic action of ATP forms a key aspect of its overall effect in the hippocampus, within the feed-forward circuit formed between CA3 pyramidal neurons, GABAergic interneurons and output CA1 pyramidal neurons. Progress from several groups is beginning to converge and suggest ways in which ATP signaling may be involved in the integrative actions of this circuit. Presynaptic P2X2 receptors increase glutamate release onto interneurons but not pyramidal neurons (Khakh et al., 2003), whereas postsynaptic P2Y1 receptors depolarize interneurons (Bowser and Khakh, 2004; Kawamura et al., 2004). However, most likely because there are few pre- or post-synaptic ATP receptors that depolarize CA1 pyramidal neurons directly (Baxter et al., 2011; Khakh et al., 2003), the net effect on output neurons is dominated by heightened GABAergic synaptic inhibition. Thus, in the stratum radiatum region of the hippocampus, the cellular effects of ATP are excitatory, but the overall result on the network is dominated by increased inhibition, implying that ATP acts as a 'physiological brake' to excitation within this feed-forward circuit (Bowser and Khakh, 2004). Recent studies show that this type of inhibition by ATP is indeed engaged via ATP release from astrocytes through connexin hemichannels, triggered by activity-dependent decreases in extracellular Ca^{2+} levels within the hippocampal circuit and propagated by slow astrocyte Ca^{2+} signaling mediated by ATP (Torres et al., 2012).

Multiplicative scaling of synaptic efficacy by P2X receptors in magnocellular neurons of the hypothalamus

One of the best examples of neuromodulation by ATP acting at P2X receptors is in magnocellular neurons (MCNs) of the hypothalamus (Fig 6). Endogenous ATP acting at extrasynaptic P2X receptors plays a crucial role in modulating excitatory synaptic transmission in MCNs (Gordon et al., 2005; Gordon et al., 2009). Early work suggested that ATP mediates part of the excitatory drive to MCNs following activation of catecholamine containing cells (Day et al., 1993), and electrophysiological studies showed that MCNs expressed functional P2X receptors that mediated membrane potential depolarization and increases in membrane conductance (Hiruma and Bourque, 1995). Later studies found ample

evidence for the presence of P2X receptor mRNAs and proteins that mediated ATP-evoked inward currents and strong elevations in intracellular Ca^{2+} levels in MCNs. It seems MCNs express a mixture of P2X receptors containing P2X2, P2X4 and possibly P2X7 subunits (Vavra et al., 2011).

It has been shown that P2X receptor signaling mediates the effect of noradrenaline (NA) on AMPA receptor-mediated mEPSCs arriving onto MCNs of the rat paraventricular nucleus (Gordon et al., 2005). Brief applications of NA increased mEPSC amplitudes for long periods of time, perhaps permanently, by increasing AMPA receptor insertion into synapses (Fig. 6). NA did not act postsynaptically via adrenoceptors in MCNs; instead, NA acted on astrocytes to release ATP. This paper provided strong evidence that ATP acting via P2X receptors resulted in modulation of glutamatergic synaptic efficacy, and astrocytes were directly implicated as the source of endogenous ATP (Gordon et al., 2005). Bains and colleagues have established that the ATP signaling mechanism is engaged during afferent activity, thus demonstrating that the effects of endogenous astrocyte derived ATP are likely to be utilized *in vivo* during physiological action potential firing impinging on MCNs (Gordon et al., 2009).

The work on MCNs also provides several interesting points that may be relevant more broadly to the study of brain P2X receptor-mediated signaling. First, the authors failed to find fast ATP synaptic transmission. Second, they found that the sources of endogenous ATP were astrocytes rather than neurons, and that bursts of action potentials were needed to engage astrocyte signaling. Third, activation of P2X receptors resulted in multiplicative scaling of all synapses in MCNs. Multiplicative scaling is a form of homeostatic plasticity, whose underpinnings are only beginning to be explored. Perhaps P2X receptors are also involved in other areas of the brain in this form of plasticity? Fourth, the effect of P2X receptor activation on synaptic efficacy in MCNs took many minutes to develop, even though P2X receptor activation occurs in seconds once ATP is bound (Vavra et al., 2011). Fifth, recent data suggest that presynaptic P2X receptors also exist in the supraoptic nucleus, where their activation facilitates glutamate and GABA release (Vavra et al., 2011). The neurosecretory cells of the hypothalamus thus emerge as the best characterized model system to explore the dynamic neuromodulatory influences of pre- and postsynaptic P2X receptors (Fig 6).

Astrocyte P2X receptors and the regulation of cortical UP states by astrocytic ATP

Astrocytes are increasingly recognized as important cellular elements within neuronal circuits not only for providing metabolic and structural support to neurons, but also for their ability to regulate neuronal function through a variety of mechanisms (Attwell et al., 2010; Halassa and Haydon, 2010). Cortical astrocytes express functional P2X7 and P2X1/5 receptors (Lalo et al., 2008; Oliveira et al., 2011) in distinct populations of astrocytes in the somatosensory and prefrontal cortices, respectively, although genome-wide analysis of astrocyte mRNA expression did not reveal any P2X receptor as being particularly enriched within astrocytes (Cahoy et al., 2008). P2X1/5 receptors on cortical astrocytes may be activated by endogenous ATP release from neurons and mediate Ca^{2+} fluxes (Palygin et al., 2010).

A recent study demonstrated that astrocytes utilize ATP signaling to regulate cortical UP states, which are network driven membrane depolarizations recorded from cortical neurons (Poskanzer and Yuste, 2011). During an UP state, the membrane potential is depolarized for hundreds of milliseconds and individual neurons fire bursts of action potentials. The available data do not allow one to conclude whether the key signals/events are mediated by astrocytic or neuronal P2X receptors; however, given that cortical astrocytes and neurons both express P2X receptors, this study provides strong evidence for how astrocytes function as the source of ATP to regulate network phenomena that occur on a time scale of hundreds of milliseconds. In future studies, it will be interesting to explore the contributions of specific P2X receptors to cortical UP states using knock-out mice and the emerging pharmacology of P2X receptors and thus attempt to correlate altered UP state dynamics with possible behavioral deficits. Finally, one wonders if neuronal P2X receptor signaling scales synaptic efficacy within principal neurons or interneurons of the cortex and regulates the output of the cortical neurons, as seen in MCNs in the hypothalamus (Gordon et al., 2005; Gordon et al., 2009).

Central chemoreception and the control of respiratory drive

An important step in peripheral sensation is activation of P2X and P2Y receptors on primary afferent terminals. Such responses are fundamental to nociception (North, 2004) and in ventilatory responses to hypoxia mediated by the carotid body (Rong et al., 2003). Recent data suggest that ATP serves similar roles in the CNS and contributes to the regulation of respiratory drive. Hypercapnia (an increase in blood CO₂; pCO₂) increases breathing, and specific areas of the medulla function as central chemoreceptors (Feldman et al., 2003). Gourine and colleagues demonstrated ATP release in micromolar amounts from the ventral surface of the medulla during hypercapnia (Gourine et al., 2005). Using miniaturized ATP biosensors, these authors mapped the sites of ATP release to the retrotrapezoid nucleus (RTN), an area known to contain pH sensitive neurons (Mulkey et al., 2004). To explore roles for endogenous ATP in contributing to respiratory drive during hypercapnia, the authors blocked P2X and P2Y receptors and observed reduced sensitivity and gain of the respiratory response to increasing CO₂ levels (Gourine et al., 2005). Exogenous applications of ATP to chemosensitive areas of the medulla mimicked the effects of CO₂ on breathing, and a P2Y-preferring agonist produced qualitatively different effects, implying important roles for P2X receptors. Furthermore, studies in brain slices show that ATP-mediated signaling can affect the firing properties of RTN neurons, but that chemosensitivity of these neurons does not derive from ATP (the neurons responded to changes in pH even when P2X and P2Y receptors were blocked (Mulkey et al., 2006)). Taken together, these data suggest that RTN neurons respond directly to pH changes (Mulkey et al., 2006; Mulkey et al., 2004) and that another process releases ATP in response to pH changes to influence the firing properties of RTN neurons (Gourine et al., 2005).

Recent data suggest that the cellular sources of ATP mediating the purinergic component of the central chemosensory response to hypercapnia are astrocytes located within the ventral surface of the medulla (i.e. near the RTN) and that the astrocytes within this area are particularly pH sensitive (Gourine et al., 2010). Hence, “excited” astrocytes propagate a Ca²⁺ signal among them due to intercellular ATP release acting on P2X and P2Y receptors.

Additionally, the authors found that the acid pH-evoked depolarization of RTN neurons was abolished when ATP signaling was blocked, implying that the neuronal response was secondary to ATP release rather than due to intrinsic chemosensitivity of the RTN neurons themselves. Moreover, the authors found that expression and illumination of channelrhodopsin within astrocytes led to light-evoked ATP release and depolarization of RTN neurons via ATP. The use of channelrhodopsin within an *in vivo* preparation showed that light-evoked astrocyte Ca^{2+} elevations lead to respiratory activity that was blocked by a mixed P2X1 and P2Y1 receptor antagonist. Taken together, this study suggests that a key step in central chemoreception involves ATP release from astrocytes located on the ventral surface of the medulla, that this signal is further propagated by ATP release acting on P2X and P2Y receptors, ultimately arriving at RTN neurons to depolarize them via ATP receptors that are likely of the P2X1 or P2Y1 class (Gourine et al., 2010).

Subsequent studies have confirmed that astrocytes release ATP in response to elevations in pCO_2 , but in a manner that is independent of pH changes and by a mechanism involving connexin 26 hemichannels (Huckstepp et al., 2010). Astrocytes in other parts of the brain may also release ATP in response to hypercapnia and so the mechanism may not be specific for the respiratory centers (Dulla et al., 2005). Moreover, recent work has shown that the purinergic component of the respiratory drive is at most 30%, a fraction that relies on hemichannel mediated ATP release, and that the actions of endogenous ATP on RTN neurons were not mediated by P2Y1 receptors, leaving open the intriguing possibility P2X receptors play a critical role (Wenker et al., 2012). Furthermore, in newborn rats, RTN chemoreception depends only on the intrinsic chemosensitivity of neurons (Onimaru et al., 2012). In summary, the available data indicate that a component of central chemoreception is mediated by astrocyte ATP that likely acts on P2X receptors in the RTN, demonstrating how ATP neuromodulation has profound effects on RTN neuron function with clear behavioral outcomes in the form of respiratory responses to hypercapnia.

Systems and Mechanisms

Afferent sensations and signaling in the peripheral nervous system

Afferent nerves that carry information into the central nervous system can be excited at their peripheral ends either by direct mechanical distortion or by transmitters released from specialized sensory cells. Considerable evidence now supports the view that the transmitter released at some such sensory 'first synapses' is ATP, and that it activates P2X receptors on the primary afferent nerve endings.

Taste and chemosensation—In response to gustatory stimulation, Type II taste buds release ATP that acts on P2X2/3 heteromeric receptors to excite primary afferent nerves that run to the CNS in the facial or glossopharyngeal nerves (Finger et al., 2005). Indeed, the release of ATP is itself driven in part by positive feedback through P2X2 receptors on the taste buds themselves (Huang et al., 2011). A similar situation pertains with respect to sensory cells of the carotid body. In this case the glomus cells sense arterial oxygen levels. In response to hypoxia they release ATP, and this acts on P2X receptors to initiate impulses in the carotid sinus nerve (Rong et al., 2003).

Cough—Pulmonary vagal afferents can be directly excited by activation of P2X2/3 receptors (Kwong et al., 2008), and P2X3 receptor antagonists reduce cough in a commonly used guinea pig model (Kamei et al., 2005). ATP inhalation in humans induces cough and dyspnea, likely by direct activation of P2X receptors (Basoglu et al., 2005). There are now available highly selective antagonists for receptors containing P2X3 subunits (Gever et al., 2010) that may prove to have clinical utility.

Urinary bladder and intestine—As the bladder becomes distended, the stretch in its wall can lead to ATP release from urothelial cells (Ferguson et al., 1997). This excites the terminals of afferent fibers expressing heteromeric P2X2/3 receptors (Zhong et al., 2003), and mice with a disrupted P2X3 receptor gene exhibit bladder hyporeflexia (Cockayne et al., 2000). The central ends of these primary afferent fibers in the dorsal horn of the spinal cord also express P2X2/3 receptors, evidenced by the observation that intrathecal administration of a selective P2X2/3 receptor antagonist also strongly inhibits the micturition reflex (Kaan et al., 2010). A similar situation pertains in the intestine, where ATP and $\alpha\beta$ meATP directly excite mesenteric afferents running from the wall of the small intestine in the rat (Kirkup et al., 1999). Action potential discharge in pelvic nerve afferent fibers elicited by colon distension are much attenuated in P2X3 knock-out mice (Shinoda et al., 2009; Wynn et al., 2003). These observations all point to a role for ATP acting directly to excite primary afferent nerves in physiological and probably pathological circumstances.

Pain—Interest in P2X receptors and pain dates from early observations that ATP itself evokes pain when applied to blisters (Bleehen et al., 1976). This is likely mediated by activation of P2X3 subunits, which are restricted in their distribution to a subset of primary afferent neurons (Chen et al., 1995; Lewis et al., 1995) that also express receptors for capsaicin (TRPV1) and isolectin B4 (Guo et al., 1999; Vulchanova et al., 1998). The relevance of this observation to any role in chronic inflammatory or neuropathic pain remains less clear, however. P2X3 knockout mice do have an afferent phenotype, most notably reduced mechanical allodynia (Cockayne et al., 2000), and antagonists selective for receptors containing P2X3 subunits (Jarvis et al., 2002) reduce chronic neuropathic and inflammatory pain in the rat. Local administration of the P2X2/3 antagonist A-317491, or prior depletion of P2X3 receptor expression by intrathecal antisense oligonucleotides, also reduce the mechanical hyperalgesia evoked by carageenan (Oliveira et al., 2009) or complete Freund's adjuvant (Ballini et al., 2011). This is consistent with a role for ATP acting on peripherally situated P2X2/3 receptors, as a component of a local "inflammatory soup".

Selective P2X3 receptor antagonists can prevent the mechanical allodynia seen in neuropathic pain models, whether applied locally or intrathecally (Gum et al., 2012; McGaraughty et al., 2003). Locally, this is associated with sensitization (an increased in expression of membrane P2X3 receptors) rather than increased ATP release (Chen et al., 2005).

Cochlea—Several cells types in the cochlea express P2X receptors, and their functional roles have been comprehensively reviewed (Housley et al., 2009). The most compelling is that part played by ATP released from K llicker's organ. This organ (also known as the

greater epithelial ridge) consists of columnar supporting cells, and it is prominent in the neonatal cochlea before the onset of hearing. At this developmental stage, ATP is released from the supporting cells in bursts and depolarizes the adjacent inner hair cells, causing them to fire bursts of action potentials which, in turn, excites auditory nerves through release of glutamate (Tritsch and Bergles, 2010; Tritsch et al., 2007). The sporadic, spontaneous ATP release begins a few days after birth, and ceases with the onset of hearing. It provides the primary excitatory stimulus to the auditory nerve during this period, and this is responsible for the downstream survival and maturation of auditory neurons.

Retina—An extensive literature pertains to P2X receptor expression in the retina, both in neurons and in supporting cells (Housley et al., 2009). The speculation around their roles in normal function or disease currently lacks insight at the cellular level, but this is clearly an area worth exploring.

Disease and Mechanisms in the CNS

Pain

Recently, it has been proposed that central P2X4 receptors are involved in neuropathic pain (Trang and Salter, 2012). The key evidence here is that removal of P2X4 receptors strikingly prevents the development of mechanical allodynia following peripheral nerve injury (Tsuda et al., 2009; Tsuda et al., 2003; Ulmann et al., 2008). Peripheral nerve injury is followed by activation of spinal microglia. It is suggested that ATP acting on P2X4 receptors drives the release of BDNF from spinal microglia, and that this in turn is critical for the re-wiring that underlies the perception of mild tactile stimuli as noxious. The neuronal subtypes and specific microcircuitry involved remain to be elucidated.

P2X7 receptors can also fashion the behavioral responses to painful stimuli and, in sharp contrast to the situation with P2X4 receptors, there is now a wealth of pharmacological antagonists to be used as experimental tools, some of which are in clinical trials (Gum et al., 2012; Jarvis, 2010). The predominant expression of P2X7 receptors in the nervous system is on microglia, astrocytes and oligodendrocytes. However, some re-interpretation of experiments which used knock-out mice may be required. The mice produced by Pfizer (Masin et al., 2012) continue to express two shortened, alternatively spliced, forms of the P2X7 receptor (P2X7 13B and P2X7 13C). The mice produced by Glaxo have a functional P2X7 splice variant (P2X7(k)) that continues to be expressed in these mice: the corresponding protein is widely expressed, but it has a different N-terminus and TM1 (Nicke et al., 2009). It forms receptors which are more sensitive to ATP and which undergo a more rapid increase in permeability to organic cations (a measure of pore dilation).

The development of mechanical hypersensitivity in models of neuropathic pain is absent in the Glaxo P2X7 deletion mouse (Chessell et al., 2005), and a similar phenotype is observed in mice lacking both isoforms of IL-1 β (Honore et al., 2006b). Mechanical hypersensitivity is also prevented by intrathecal P2X7 antagonist A-438079 (Kobayashi et al., 2011) and Brilliant Blue G (He et al., 2012) or systemic administration of P2X7 receptor antagonists (Honore et al., 2006a). These effects in pain models appear to require release of IL-1 β , given that A-438079 blocks not only the ATP evoked release of IL-1 β but also the release evoked

by LPS (Clark et al., 2010). Indeed, LPS-induced release of IL-1 β is absent in lumbar spinal cord slices from P2X7 deletion mice. These results imply that the rewiring of microcircuitry in the spinal cord that occurs during prolonged inflammation or nerve injury involves ATP release, the activation of P2X7 receptors on spinal microglia, and the release of IL-1 β . Our understanding of the source of ATP has advanced little since Pamela Holton showed that sensory nerves release ATP when stimulated electrically (Holton, 1959). As for the target action of the released IL-1 β , one suggestion is that it leads to increased phosphorylation of the NR1 and NR2B subunits of the NMDA receptor and perhaps an LTP-like phenomenon (Zhang et al., 2008), which has been associated with behavioral hyperalgesia (Brenner et al., 2004).

Some independent support for considering P2X7 receptors as potential targets in pain therapies is also provided by genetic associations. Six of seven inbred mouse strains that showed less than average allodynia also expressed P2X7 receptors with a single nucleotide polymorphism resulting in the P451L mutation (Sorge et al., 2012). Mouse P2X7[P451L] receptors have impaired function as measured by ATP-evoked uptake of YO-PRO-1 (a commonly used optical measure of pore dilation) but not when measured as ATP-induced ionic current (Adriouch et al., 2002; Young et al., 2006). This polymorphism, as well as two others known to result in impaired P2X7 receptor function (H155Y, R270H) were found more often in subjects reporting a higher than average pain level following mastectomy, or in osteoarthritis (Sorge et al., 2012).

Demyelinating disease

Inflammation is a component of the degeneration that occurs in several nervous system disorders, and the involvement of P2X receptors has been investigated in certain animal models of disease. In mice, experimental autoimmune encephalitis (EAE) can be induced by immunization with myelin oligodendrocyte glycoprotein. Studies on P2X7 deletion mice appear to be contradictory. In the Pfizer mice, the clinical and pathological features of EAE are more marked in “knock-outs”, whereas in the Glaxo mice the deficiency of the P2X7 receptor suppressed the development of EAE (Sharp et al., 2008).

Spinal cord injury

Considerable interest was created by the report that blocking P2X7 receptors improves functional recovery after spinal cord injury in rats (Wang et al., 2004). The receptor antagonists used (oxoATP and PPADS) are non-selective, and their actions cannot be reliably attributed to P2X7 receptor blockade. The results were later extended by using Brilliant Blue G (Cotrina and Nedergaard, 2009; Peng et al., 2009), which is a more selective blocker of mouse and rat P2X7 receptors (Jiang et al., 2000). However, a recent study failed to replicate these results (Marcillo et al., 2012). Additional systematic approaches are clearly required, using the highly selective P2X7 receptor antagonists that are now available. A role for P2X4 receptors has also been suggested. In mice with deleted P2X4 receptors, there was a decrease in caspase-1 cleavage and IL-1 β production at the injury site, which is correlated both with diminished neutrophil and macrophage infiltration at the injury site, and improved functional outcome (de Rivero Vaccari et al., 2012).

Stroke

Large amounts of ATP are released from damaged cells as a result of ischemia, which may activate P2X7 receptors. This provides the logic for considering blockade of P2X7 receptors as a possible therapeutic regimen. Pretreatment with PPADS improves recovery from experimental stroke in rats with permanent middle cerebral artery occlusion (Lämmer et al., 2011). Treatment with brilliant Blue G beginning 30 minutes after middle cerebral artery occlusion caused a 60% reduction in brain damage measured three days later (Arbeloa et al., 2012). These studies need to be extended to determine if P2X7 receptors are valid targets in stroke and whether the relevant receptors are located on astrocytes or neurons.

Summary and future opportunities

In this review, we have highlighted some of the recent literature that sheds new light on how P2X receptors work and how they mediate neuromodulation in diverse systems.

Representing a novel structural class of ion channels with several unique functional properties, and mediating fascinating slow responses, the physiology of ATP P2X receptors has challenged our precepts of how a fast neurotransmitter-gated cation channel should look and behave. New biophysics and biology has been discovered, and many early biophysical and physiological insights have been supported with high resolution crystal structures, optical approaches and molecular genetics.

Based on the aforementioned latest breakthroughs, we propose that P2X receptors have evolved to fulfill unique biological functions and occupy signaling niches that are not readily met by other fast neurotransmitter systems. Given that glia constitute about half the cells in the brain, express multiple ATP receptors and release ATP through a variety of mechanisms, we suggest that a major facet of ATP and P2X receptor biology is related to glia and their slow neuromodulatory functions in the nervous system. Viewed from this neuromodulatory capacity and largely unexplored potential, ATP acting via P2X receptors is a physiologically important signal, particularly for linking slow glial communications with fast neural microcircuit computations.

For continued progress, it will be vital that we explore P2X receptor functions with the best available tools. Luckily, many P2X receptor knockout mice are now available and selective antagonists are being discovered (Table 1). With the publication of the P2X crystal structures, several classic biophysical questions have been answered and there can be little doubt the field has moved into an exciting new era. We now await the structure of a full length P2X receptor with its cytosolic domains, which will allow us to relate the findings to the wealth of studies on receptor function and gating. One would also like to know if the ATP bound state represents the only open state or an intermediate conformation of the permeation pathway. It will be important to continue to use simple systems to explore ion channel biophysics, but more nuanced cell biological issues such as trafficking, protein-protein interactions and receptor regulation should be performed in intact systems such as native cell lines, primary cultures and *in vivo* to render the findings meaningful in a physiological manner. Additionally, the generation of a battery of P2X reporter mice will allow for the study of live tissue specimens, thus ending our reliance on staining with antibodies of limited selectivity and on dead tissue. Use of such reporter mice is perhaps the

only immediately obvious way to address the issue of heterogeneous responses, because it would allow researchers to record from genetically specified cell populations. It also seems necessary to generate mouse lines expressing tamoxifen inducible Cre recombinase under the P2X4 and P2X7 receptor promoters because of their roles in pain, allowing mechanistic experiments *in vivo* to commence, thus ending correlative interpretations and aiding exploration of causation in the spinal pain microcircuitry. Our improved knowledge of ATP binding and gating in P2X receptors also presents itself as an opportunity to design/reengineer P2X receptors to perform specific tasks, such as chaperoning other proteins to the edges of dendritic spines or to nerve terminals in the treatment of synaptic diseases, or to provide known and desired fluxes in specific domains of neurons that need them. It seems necessary to evaluate if cross inhibition between P2X receptors and other ion channels has behavioral consequences, either physiologically or during disease states, and thus explore this neuromodulatory regulation. Similarly, we need to determine the proteome of key P2X receptors to understand both physiology and biophysics of these proteins: P2X4 and P2X7 receptors seem to be a good place to start given their roles in pain.

The field of P2X receptor signaling holds great promise to treat human diseases, which is an important remit and mandate of publicly funded biological research. Recent exciting breakthroughs offer ample chances for national research agencies to invest in an area with largely untapped clinical potential and thus deliver on their promise to advance human therapeutics. From this perspective, the entire field eagerly awaits the results of ongoing clinical trials on the first generation of drugs targeting P2X receptors to treat pain states, bladder dysfunction and cough (Ford, 2012; Gum et al., 2012). There is huge unexplored potential to exploit ATP signaling for treating CNS disease. We need to explore these possibilities from a grass roots level and up.

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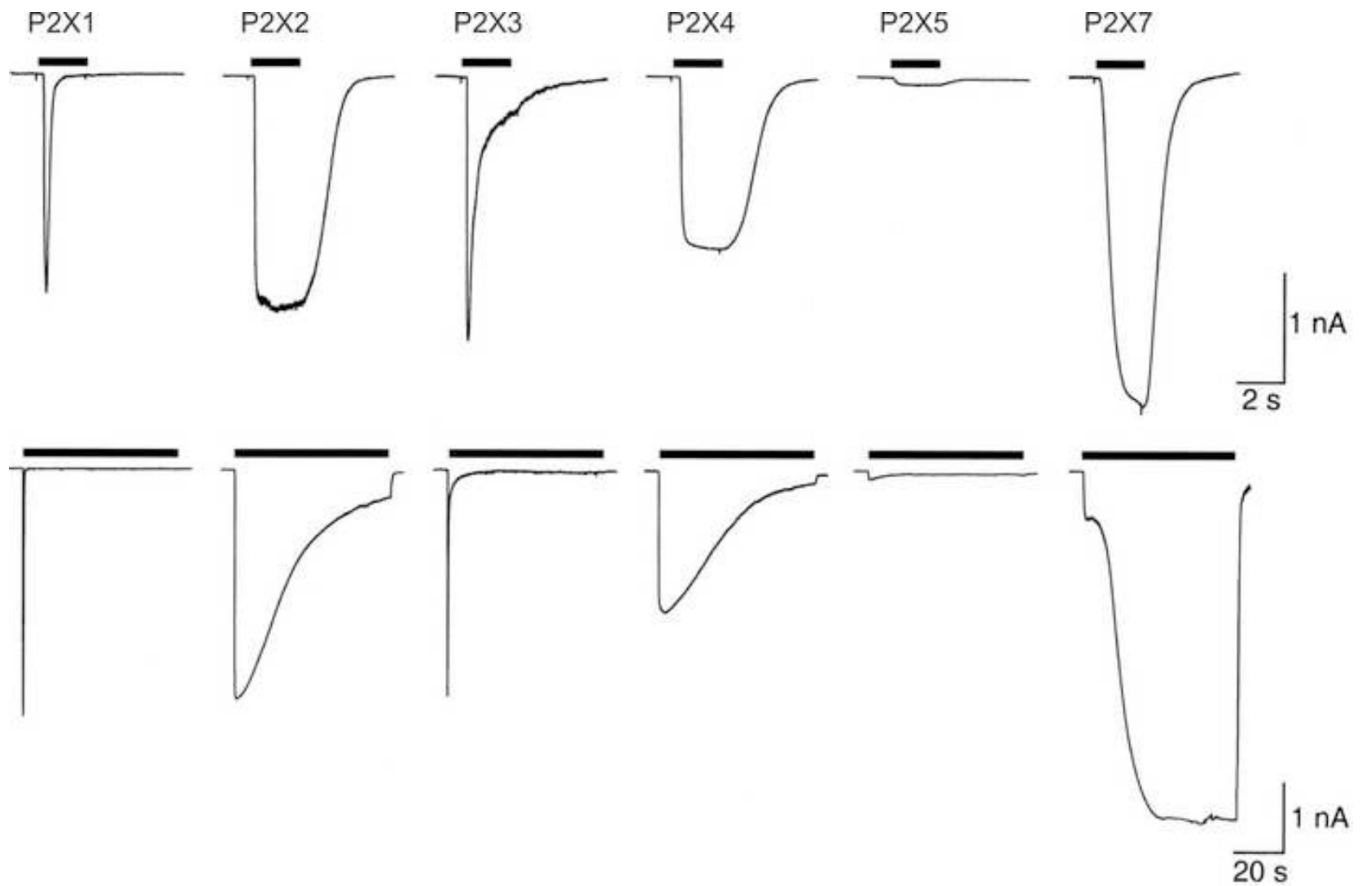


Figure 1. Fast (top) and slow (bottom) desensitization is compared for functional recombinant homomeric rat P2X receptors expressed in HEK-293 cells

Note the 10-fold difference in time scale between top and bottom panels. Fast desensitization is observed only with P2X1 and P2X3 during brief applications (2 s duration) of ATP (30 μ M, except 1 mM for P2X7 owing to its lower sensitivity – see Box 1). Slow desensitization is observed for P2X2 and P2X4 with more prolonged applications (60 s duration) of ATP (30 μ M, except 1 mM for P2X7). In all cases except P2X7, the response shown is that seen for the first application of ATP to that cell. For P2X7, one 2 min application of ATP had been made before the application shown. Note that the rising phase of the response recorded for P2X7 receptors is biphasic. This likely reflects pore dilation associated with the second slower phase (discussed in the text; see also Fig 4). The experiments were performed by Dr. A. Surprenant and the figure is reproduced here from North (2002) with the author's permission.

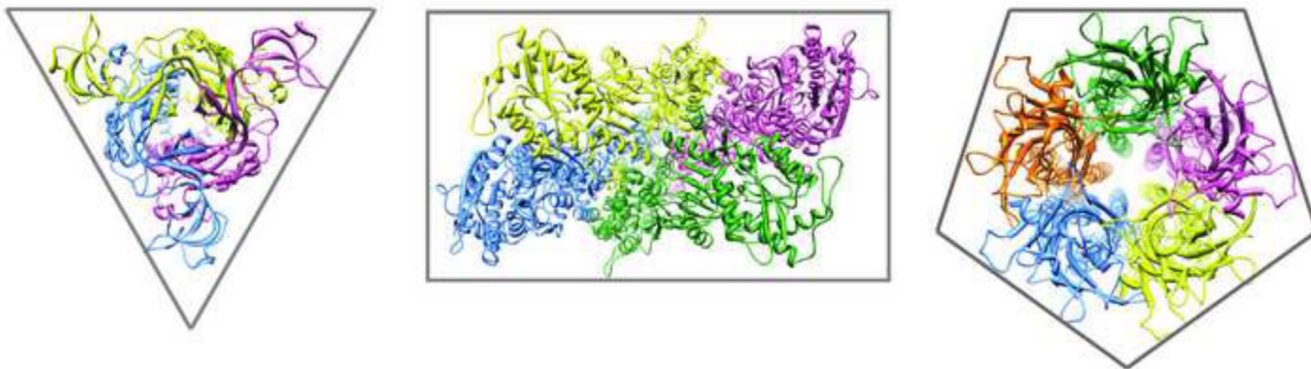


Figure 2. P2X receptors define a unique family of ligand-gated ion channels

Trimeric, tetrameric and pentameric ligand-gated ion channels, viewed from the outer surface. *Left*, P2X receptor family. ATP-gated P2X4 cation channel from zebrafish (*Danio rerio*) in an open state. Modified from 4DW1. *Middle*, AMPA receptor family. Glutamate-gated GluA2 cation channel from rat, in closed state (*Rattus norvegicus*). Modified from 3KG2. *Right*, Nicotinic acetylcholine receptor family. Glutamate-gated chloride channel from the nematode worm (*Caenorhabditis elegans*) in open state. Modified from 3RHW.

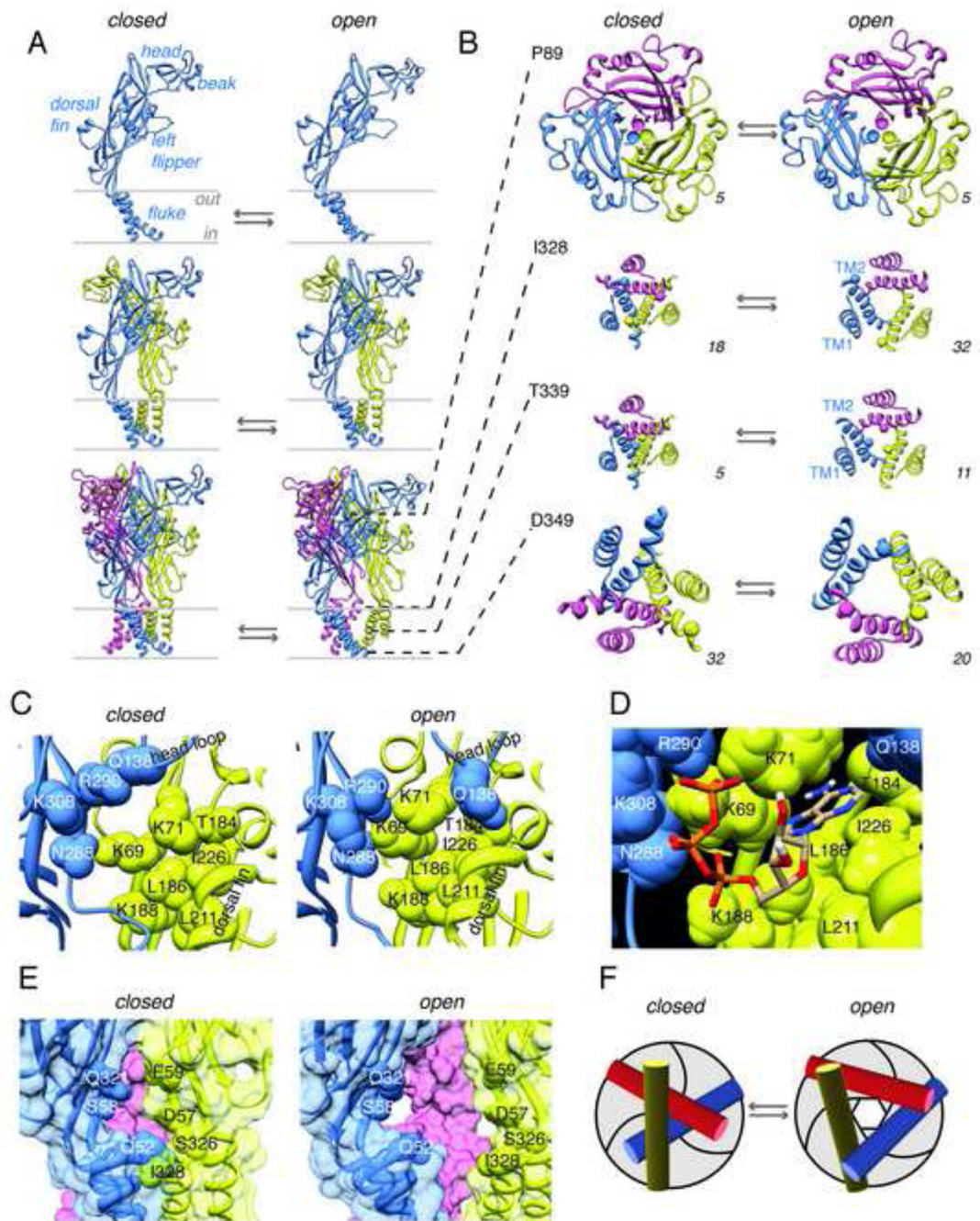


Figure 3. P2X receptor structure, the ATP binding site, and the permeation pathway

A. Assembly of P2X₂ receptors from three subunits, depicted in closed (*left*) and open (*right*) states. Upper, middle and lower panels show one, two and three subunits. **B.** Cross-sections of open and closed rat P2X₂ receptors, at the level of the residues indicated, the C β atoms of which are shown as spheres. Number in lower right of each panel is the diameter of the circle (in Å), passing through the three C β atoms. D349 viewed from below, others viewed from above. **C.** Key residues in the ATP binding pocket, in open and closed configurations. Two subunits depicted. **D.** ATP molecule positioned in binding pocket. The

U-shape of the triphosphate chain curls around the nitrogen atom of K69. Further explanation in text. **E.** The lateral portal between two subunits (blue and yellow; pink subunit visible through the portal. Key residues on the edge of the portal are indicated. **F.** Schematic to illustrate iris-like movement of TM2 domains. Molecular models created in Modeller 9v7 (Sali and Blundell, 1993) using 4DW0 (closed) and 4DW1 (open) as templates, energy minimized using MolProbity (Davis et al., 2007) and displayed in Chimera 1.6 (Pettersen et al., 2004).

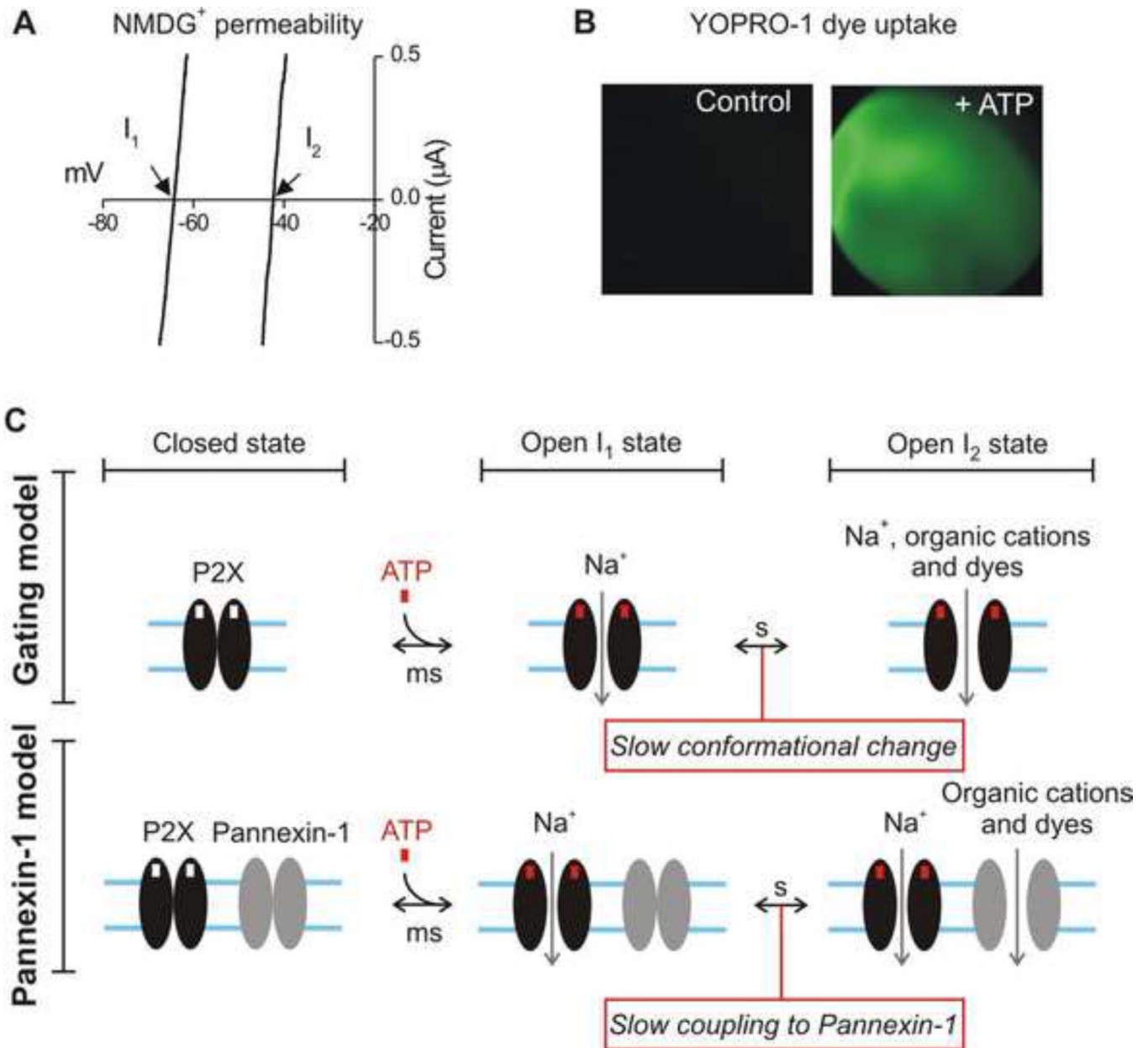


Figure 4. Examples and models of slow conformational changes in P2X receptors displaying pore dilation

A. Current-voltage plots for example recordings of the I₁ and I₂ states for P2X₂ receptors expressed in *Xenopus* oocytes are shown. The I₁ state has low permeability to the organic cation NMDG⁺ (hence a negative reversal potential). With time the permeability to NMDG⁺ increases, indicated as a shift in reversal potential by ~ +16 mV. **B.** Representative images of a *Xenopus* oocyte bathed in YOPRO-1 before and during 100 μM ATP (for 30s). The oocyte becomes fluorescent as YOPRO-1 enters the cell, presumably via the I₂ state. The data shown in A and B are from Supplementary information accompanying (Chaumont and Khakh, 2008). **C.** Diagrams illustrating the differences between the Panx-1 and gating models for the I₂ state. In the gating model, small cations, organic cations and dyes enter due

to slow conformational changes in the P2X pore. In the Panx-1 model, organic cations and dyes enter via an accessory ion channel protein called Panx-1. On balance the field is converging on the gating model, but there is also evidence for the Panx-1 model in the case of P2X7 receptors that are natively expressed (as discussed in the main text). In both models, the cytosolic domains of P2X receptors are important.

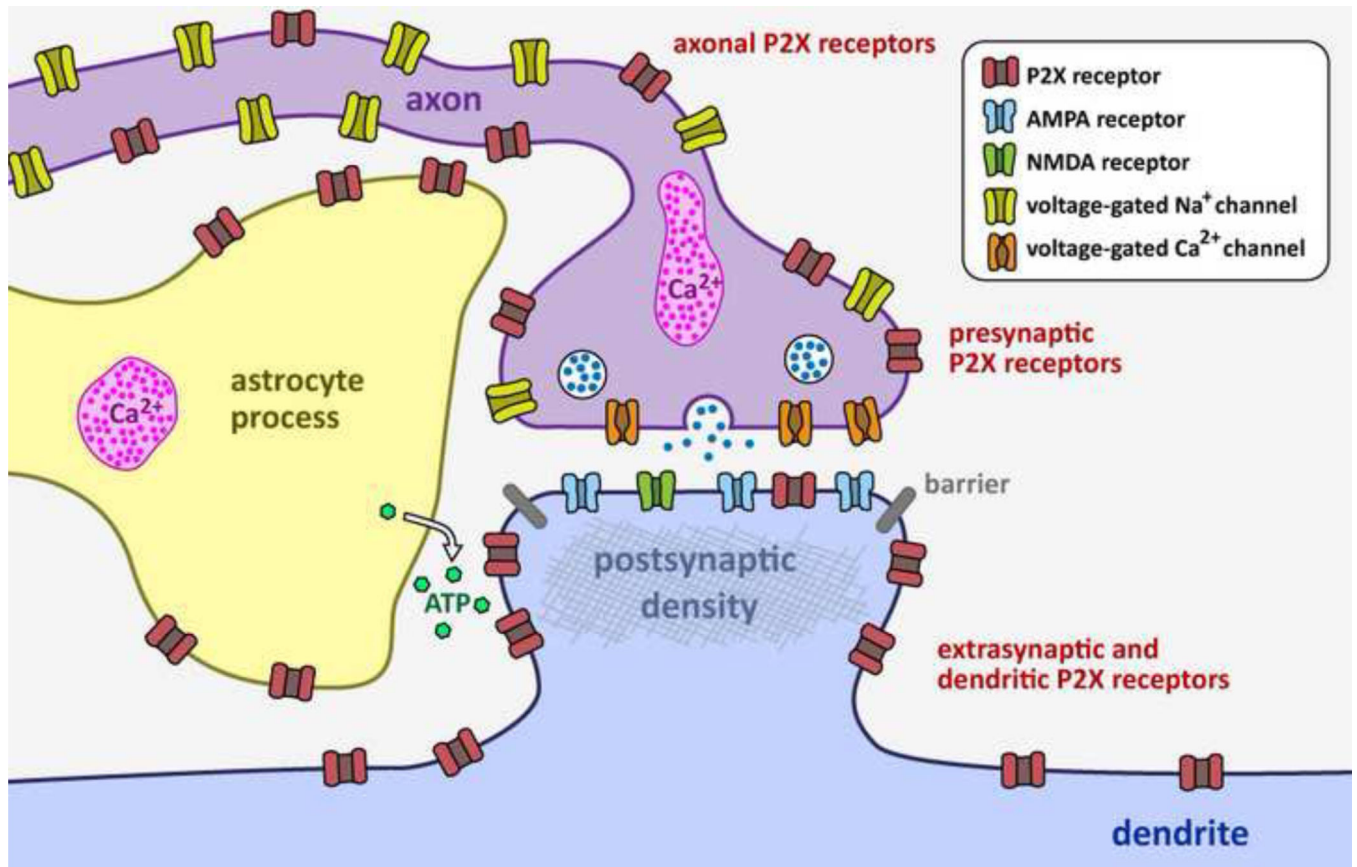


Figure 5. Summary showing locations of P2X receptors in synapses

Note: this is a cartoon summarizing twenty years of work on various synapses. As such we are not aware of any synapse that expresses P2X receptors in all the locations shown in this cartoon. ATP is released from astrocytes from lysosomes, vesicles and through hemichannels, as discussed in the text. All three of these release mechanisms need to be considered on a case-by-case basis, because it is likely that they operate in different settings and in different brain nuclei. Electron microscopy and single molecule imaging shows that P2X2 and P2X4 receptors do not enter synapses (Masin et al., 2006; Richler et al., 2011; Rubio and Soto, 2001), implying the existence of a restrictive barrier which is shown here as a gray bar at the edge of the postsynaptic density.

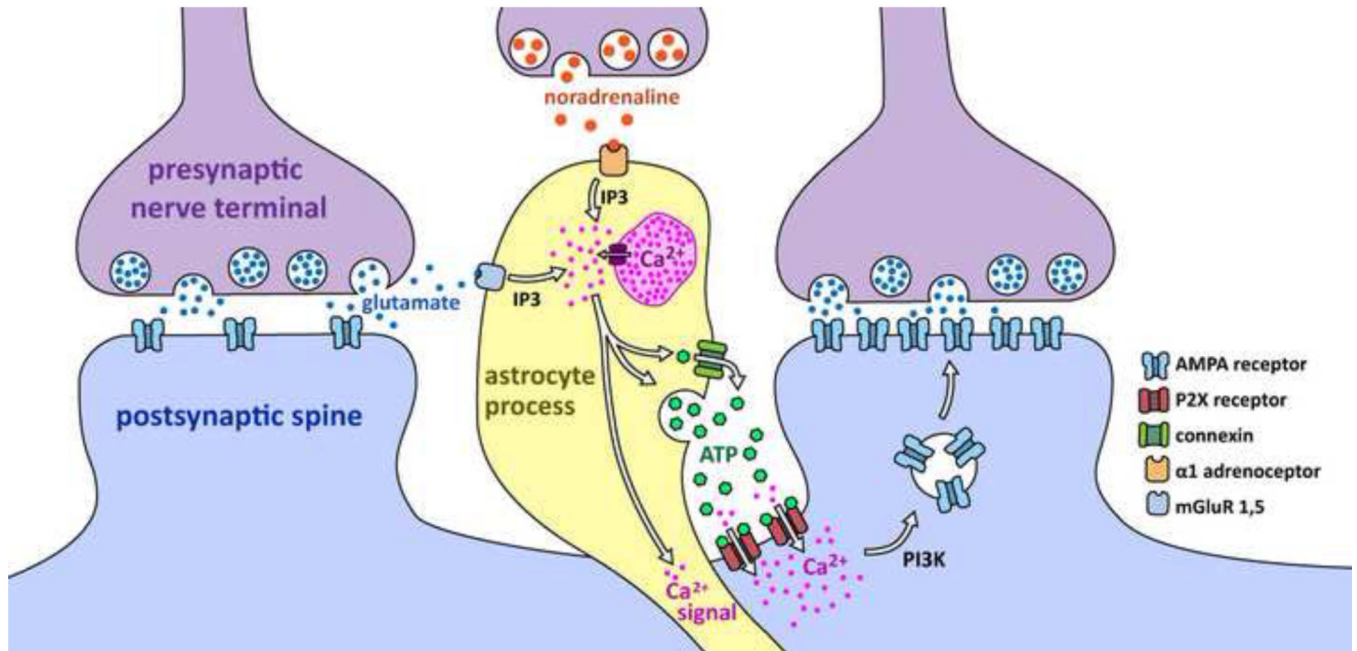


Figure 6. Cartoon summary of synaptic scaling by ATP acting on P2X receptors in magnocellular neurons (MCNs) of the hypothalamus

Based on work by the Bains lab (Gordon et al., 2005; Gordon et al., 2009). Glutamate and noradrenaline cause release of intracellular Ca^{2+} within astrocyte processes, leading to ATP release into the extracellular milieu. This activates extrasynaptic P2X receptors, which through a Ca^{2+} - and PI3K-dependent mechanism, leads to synaptic scaling by insertion of AMPA receptors into dendritic spines of MCNs. The astrocyte Ca^{2+} signals may spread within astrocytes leading to similar effects at other synapses. The effects of glutamate and noradrenaline in this context are both reduced when astrocyte processes withdraw from MCNs during lactation and dehydration (see text for details). In this context, the astrocyte process is not actively contributing to synaptic transmission, but rather via ATP it is exerting a strong activation-dependent neuromodulation. Similar responses at other brain synapses with astrocyte processes nearby ought to be explored.

Table 1

Key facts and the useful pharmacopeia for exploring P2X receptors.

Gene name	P2X1 <i>P2RX1</i>	P2X2 <i>P2RX2</i>	P2X3 <i>P2RX3</i>	P2X4 <i>P2RX4</i>	P2X5 <i>P2RX5</i>	P2X6 <i>P2RX6</i>	P2X7 <i>P2RX7</i>
Desensitization (complete in)	Fast (<1s)	Slow (>20s)	Fast (<1s)	Slow (>20s)	Slow (>20s)	-	Slow (>20s)
Pore "dilatation"	No	Yes	No	Yes	-	-	Yes
P_{Ca}/P_{Na}	4.8	2.8	1.2	4.2	1.5	-	-
Fractional Ca^{2+} current (%)	12.4	5.7	2.7	11	4.5	-	4.6
KO made	Yes	Yes	Yes	Yes	-	-	Yes
Agonist EC₅₀ values (μM)							
ATP	0.07	1.2	0.5	10	10	12	100
2MeSATP	0.07	1.2	0.3	10	10	9	100
αβmeATP	0.3	> 300	0.8	> 300	> 300	> 100	> 300
BzATP	0.003	0.75	0.08	7	> 500	-	20
Antagonist IC₅₀ values (μM)							
Suramin	1	10	3	>500	4	>100	500
PPADS	1	1	1	>500	3	>100	50
TNP-ATP	0.006	1	0.001	15	-	-	> 30
IP ₅ I	0.003	> 300	2.8	Potentiation	-	-	-
A-317491	> 10	> 100	0.10	> 100	-	> 100	> 100
RO-3	> 100	> 100	0.10	> 100	> 100	> 100	> 100
A-740003	> 100	> 100	> 100	> 100	-	> 100	0.05
AF-353	> 10	> 10	0.01	> 10	> 10	-	> 10
A-438079	> 100	> 100	> 100	> 100	-	> 100	0.06
A-804598	> 100	> 100	> 100	> 100	-	> 100	0.01
MRS2179	80	> 100	> 100	-	-	-	> 100
NF279	9	30	50	> 100	-	-	20
NF449	0.7	> 100	> 100	> 100	-	-	> 100
Modulator EC₅₀ values (μM)							

	P2X1	P2X2	P2X3	P2X4	P2X5	P2X6	P2X7
Ivermectin	-	> 30	> 30	0.25	-	-	> 30
Zn ²⁺	-	Increase EC ₅₀ = 7μM	-	Increase 2μM	-	-	-
H ⁺	Decrease pKa 6.3	Increase pKa 7.3	Decrease pKa 6.0	Decrease pKa 6.8	-	-	Decrease pKa 6.1

Notes for table 1: -Indicates that this information is not yet available. The Pf(%) values are for the rat subunits, but the pCa^{2+}/pNa^+ values are also shown for human and rat isoforms as appropriate for the available data. In the case of suramin and PPADS IC50 values the data for P2X4 are highly species-dependent.

Abbreviations used in the table: **ATP:** adenosine 5'-triphosphate, **2meSATP:** 2-methylthioadenosine 5'-triphosphate, **α,βmeATP:** alpha beta methylene adenosine 5'-triphosphate, **BzATP:** 2,3-O-(4-benzoylbenzoyl)-ATP, **PPADS:** pyridoxal-5'-phosphate-6-azophenyl-2',4'-disulphonic acid, **TNP-ATP:** 2',3'-O-(2,4,6-Trinitrophenyl) adenosine 5'-triphosphate, **IP5I:** Diinosine pentaphosphate, **A-317491:** 5-((3-phenoxybenzyl)((1S)-1,2,3,4-tetrahydro-1-naphthalenyl)amino)carbonyl)-1,2,4-benzenetricarboxylic acid **RO-3:** 5-(2-isopropyl-4,5-dimethoxybenzyl)pyrimidine-2,4-diamine, **A-740003:** N-[1-(N"-cyano-N'-quinolin-5-ylcarbamimidamido)-2,2-dimethylpropyl]-2-(3,4-dimethoxyphenyl)acetamide, **AF-353:** 5-(5-iodo-2-isopropyl-4-methoxyphenoxy)-pyrimidine-2,4-diamine, **A-438079:** 3-(5-(2,3-dichlorophenyl)-1H-tetrazol-1-yl)methyl pyridine, **A-804598:** 2-Cyano-1-[(1S)-1-phenylethyl]-3-quinolin-5-ylguanidine, **MRS2179:** 2'-deoxy-N⁶-methyl adenosine 3',5' diphosphate, **NF279:** 8,8'-(carbonylbis(imino-4-1-phenylenecarbonylimino-4,1-phenylenecarbonylimino))bis(naphthalene)-1,3,5-trisulfonic acid, **NF449:** 4-(((3-(((3,5,5-bis(2,4-disulfophenyl)carbamoyl)phenyl)carbonyl)amino]benzene-1,3-disulfonic acid.