Mutation of histidine 286 of the human P2X$_4$ purinoceptor removes extracellular pH sensitivity

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1. Effects of external pH on the human P2X$_4$ purinoceptor, an ATP-activated ion channel, were studied using the *Xenopus* oocyte expression system.

2. Changing the external pH from 7·4 to 6·5 significantly reduced, whilst an increase to pH 8 enhanced, maximum ATP-activated current amplitude, without changing the current–voltage relationship of the ATP-activated current.

3. Diethyl pyrocarbonate (DEPC; 10 mM) treatment of P2X$_4$-injected oocytes had no effect on the pH sensitivity of the ATP-activated current.

4. Site-directed mutagenesis of histidine 286 (H286) to alanine completely abolished the pH sensitivity of the P2X$_4$ receptor at all agonist concentrations. ATP potency showed a small (fourfold) leftward shift. Mutagenesis of the other three histidines present in the P2X$_4$ sequence had no effect on pH sensitivity.

5. The results show that pH modulation of P2X$_4$ in the pathophysiological range is mediated by protonation of H286. This provides direct confirmation that pH sensitivity resides in the P2X$_4$ channel protein rather than the agonist species.

P2X channels may mediate fast excitatory actions of ATP when ATP is co-released with other transmitters in both the central and peripheral nervous systems (see North & Barnard, 1997, for review). Modulators of ATP-gated channels may therefore act as important regulators of synaptic transmission. To date seven isoforms of the P2X receptor have been cloned. These share 35–50 % sequence homology and in *in vitro* expression systems are capable of forming homomeric channels (Surprenant, 1996). P2X receptors show widespread distribution in neuronal, secretory and muscle tissues with P2X$_4$ (Bo *et al.* 1995; Buell *et al.* 1996*b*; Seguela *et al.* 1996; Soto *et al.* 1996) and P2X$_6$ predominating in the CNS (Buell *et al.* 1996*a*; Collo *et al.* 1996) suggesting that these sub-types might have important physiological and pathophysiological roles in the brain.

Neuronal activity (Chesler & Kaila, 1992) as well as a number of pathophysiological conditions give rise to changes in extracellular pH across a range from pH 6·0 to pH 8·0 (Siesjö *et al.* 1996). In common with a number of other ion channels, ATP-gated channels are sensitive to extracellular pH. P2X receptors native to bullfrog dorsal root ganglion neurons and rat nodose ganglion neurons (Li *et al.* 1996, 1997), as well as rat P2X$_2$ (rP2X$_2$) receptors expressed in *Xenopus* oocytes (King *et al.* 1996), show enhanced ATP-activated currents when extracellular pH is reduced. In contrast, a reduction in extracellular pH decreased ATP-

induced currents in human P2X$_1$ (hP2X$_1$), rP2X$_3$ and rP2X$_4$ receptors (Stoop *et al.* 1997).

The detailed structure of P2X channels and how the structure determines channel function is poorly understood. Analysis of primary sequence data indicates that the channels are composed of intracellular N- and C-termini, and two transmembrane domains joined by a large extracellular loop region (Hansen *et al.* 1997). This has been supported by glycosylation (Newbolt *et al.* 1998) and cysteine scanning studies (Rassendren *et al.* 1997). There is little information on the molecular location of sites that modulate channel gating but this basic topography permits structure–function studies of modulation by extracellular ions to be focused on the single putative extracellular domain.

In this paper we have characterized the sensitivity of the hP2X$_4$ channel to extracellular pH and performed site-directed mutagenesis to identify the site of protonation. We chose to look at this member of the family because it is the most abundant P2X channel expressed centrally and because the rP2X$_4$ channel shows the greatest sensitivity to extracellular pH (Stoop *et al.* 1997). We show that ATP-gated currents in hP2X$_4$ channels are reduced by lowering extracellular pH in the physiological range. pH sensitivity is abolished by site-directed mutagenesis of histidine (H) 286 to alanine (A) indicating that protonation of this histidine

mediates the pH sensitivity of the $hP2X_4$ receptor. These data define the molecular basis of an important regulatory site on this channel. Brief reports of some of this work have appeared elsewhere in abstract form (Clarke *et al.* 1998).

METHODS

Preparation of $hP2X_4$ histidine mutants

The full-length clone of $hP2X_4$ (GenBank accession number A65875) was used in these studies. This sequence differs from that published by Garcia-Guzman *et al.* (1997) only in an alanine instead of a serine at position 6. Mutant H172A was introduced using the QuickChange polymerase chain reaction (PCR)-based mutagenesis kit (Stratagene) according to the manufacturer's protocol, using the expression vector, $pCDN-P2X_4$, as the template for mutagenesis. Mutants H140A, H241A and H286A were mutated using a PCR-based method (Li & Shapiro, 1993). The 5′ primers for each mutation (H140AFOR, H241AFOR and H286AFOR) were used with the 3′ primer, pCDNREV, and $pCDN-P2X_4$ as the template, to produce fragments of $P2X_4$ containing each mutation at the 5′ end. These products were then used as the 3′ primer in a second PCR reaction, with pCDNFOR as the 5′ primer and $pCDN-P2X_4$ as the template. The product of this second PCR reaction was a full-length $P2X_4$ gene containing the mutation. The PCR products were then digested with *Eco*RI and *Bam*HI, and ligated into *Eco*RI/*Bam*HI-digested pCDN.

All mutations were confirmed by sequencing.

The primer sequences are as follows.

H140AFOR:
    5′ CTCTGCCGGCACC**GCG**AGCAACGGAGTCTC 3′
H172AFOR:
    5′ GTGGAGGATGACACA**GCT**GTGCCACAACCTGC 3′
H241AFOR:
    5′ GTGGAGAACGCAGGA**GCG**AGTTTCCAGGACATG 3′
H286AFOR:
    5′ CACGGGACGTTGAG**GCG**AACGTATCTCCTGGC 3′
pCDNFOR:
    5′ CGCAAATGGGCGGTAGGCGTG 3′
pCDNREV:
    5′ GAGGGGCAAACAACAGATGGC 3′

Base changes introducing the mutations are in bold type and underlined.

Oocyte preparation and injection

Surgery was carried out in accordance with the United Kingdom Animals (Scientific Procedures) Act (1986) and conformed to SmithKline Beecham ethical standards. Oocytes were removed from each mature female *Xenopus laevis* on two separate occasions. For the first surgery the *Xenopus laevis* frog was completely anaesthetized by immersion in 0·2% MS-222. The level of anaesthesia was assessed firstly by checking if the frog could right itself when tipped over in the anaesthetic, it was then removed from the anaesthetic and reflexes tested by both pinching and pricking the toes with a pin. Once the frog was unresponsive a small incision was made in the abdomen and oocytes removed. The incision was closed in two layers and the frog was allowed to recover in a small freshwater tank whilst being observed before being returned to the main holding tank. At least 1 week after the first removal of oocytes the frog was completely anaesthetized by immersion in 0·4% MS-222 and then decapitated before the remaining oocytes were removed. Upon removal oocytes were

placed into a sterile modified Barth's solution (MBS) containing (mM): NaCl 88, KCl 1, $NaHCO_3$ 2·4, Hepes 1·5, $MgSO_4$ 0·82, $Ca(NO_3)_2$ 0·33, $CaCl_2$ 0·41; buffered to pH 7·4 with NaOH. The tissue was disaggregated into small clumps and agitated in calcium-free Barth's solution (direct replacement of calcium with magnesium) containing 1–2% collagenase A (Boehringer Mannheim) for 1–3 h at room temperature. Injections of cDNA for either human $P2X_4$ receptors or mutants (in pCDN vector) were made into the nuclei of defolliculated Stage 5–6 *Xenopus* oocytes (1·5 ng cDNA per oocyte or 5 ng cDNA per oocyte for H286A). The oocytes were then incubated in filtered MBS supplemented with 0·1 mg ml^{-1} gentamicin at 19–22 °C. Diethyl pyrocarbonate (DEPC) results were obtained by adding 10 mM DEPC to this solution and incubating for 24 h prior to recording.

Recording

Oocytes were placed in a recording chamber 1–3 days after injection and continuously perfused (14 ml min^{-1}) with MBS solution (adjusted to the appropriate pH with 1 M NaOH or 5 M HCl). The solution was applied using large bore tubing (internal diameter 1·5 mm) which facilitated rapid solution exchange (half-time 350–1000 ms). Recordings were obtained using the two-electrode voltage clamp technique, holding potential −80 mV (unless otherwise stated). Electrodes used were low resistance (0·8–2 MΩ) when filled with 3 M KCl. Current responses, digitized at 2 kHz and filtered at 1 kHz, were stored for later analysis using chart (CED) software.

External solution pH was equilibrated for 1 min before applying ATP, buffered to the relevant pH, for 3–5 s. This length of application of ATP allowed the current to reach maximal amplitude before washout. Reproducible responses to 100 μM ATP at pH 7·4 were recorded at the start of each experiment. These could be obtained with a washout interval of 3–5 min. Concentration–response curves to ATP were obtained at extracellular pH values of 6·5, 7·4 and 8·0 over a range of concentrations from 30 nM to 3 mM. Data were normalized to the current evoked by 100 μM ATP at pH 7·4 and curves fitted to the logistic equation using Origin software. The concentration of ATP that gave 20% of the maximum response (EC_{20}) at each pH was estimated from these concentration–response curves for use in later experiments.

For the experiments investigating the effect of pH on the ATP current–voltage relationship, the oocyte was bathed in the test pH solution and currents activated by ATP EC_{20} whilst ramp tests from −100 to +50 mV were performed using pCLAMP software (Axon Instruments).

RESULTS

Rapid bath application of 300 μM ATP onto *Xenopus* oocytes expressing the $hP2X_4$ receptor, held under voltage clamp at −80 mV, resulted in rapidly rising inward currents. Maximal responses recorded reached up to 5 μA at 300 μM (pH 8) and showed rapid desensitization (Fig. 1*A*). With washout periods of 3–5 min, reproducible responses could be obtained. ATP-gated currents were extremely sensitive to extracellular pH. Figure 1*A* shows responses to 300 μM ATP from one oocyte over a range of extracellular pH. Peak amplitude of the inward current declined from 1·5 to 0·25 μA as pH was reduced from 8·0 to 6·5. Full concentration–response curves to ATP (30 nM to 3 mM) at pH 6·5, 7·4 and 8 show EC_{50} values of 14·5, 2·6 and 1·9 μM, respectively (Fig. 1*C*). Maximal responses at each of the pH

values tested were obtained at 300 μM ATP and relative to 100 μM ATP at pH 7·4 were: 0·53 ± 0·11 ($n = 7$) at pH 6·5, 1·02 ± 0·07 ($n = 6$) at pH 7·4 and 1·3 ± 0·12 ($n = 6$) at pH 8 indicating that the maximal amplitude of response to ATP decreased with decreasing pH as well as a rightward shift in the concentration–response curve. This differed to the findings for the rP2X$_4$ receptor in which pH 6·5 produced a rightward shift in the concentration–response curve to ATP but had no effect on the maximal response obtained (Stoop *et al.* 1997).

The best fitting pH titration curve to this data at 300 μM ATP yielded an apparent pK_a (negative log of the acid dissociation constant) of 6·8 and a slope of 1·65 (Fig. 1*B*). This pK_a suggests that an imidazole side chain on an extracellular histidine is most likely to be the proton binding site (Lehninger, 1975).

The rapid onset of pH-induced modulation suggests a direct effect of pH change by an extracellular site of action. To examine this further we analysed the effect of pH on the ATP-gated currents at a range of potentials. Figure 1*D* shows voltage ramp data generated at pH 8·0, 7·4 and 6·5 with EC$_{20}$ ATP concentrations and scaled to the current at −80 mV. The ramp currents do not differ significantly, showing that the pH sensitivity is not affected by membrane potential between −80 and +50 mV. This again indicates that the pH-sensitive site is outside the membrane field.

Diethyl pyrocarbonate (DEPC) carbethoxylates the imidazole ring of histidine and the side groups of arginine, cysteine and tyrosine (Leonard *et al.* 1970). DEPC treatment of the hP2X$_4$-injected oocytes was carried out in an attempt to alter the pH sensitivity of the hP2X$_4$ receptor. We tried prolonged exposure to high concentrations of DEPC as no
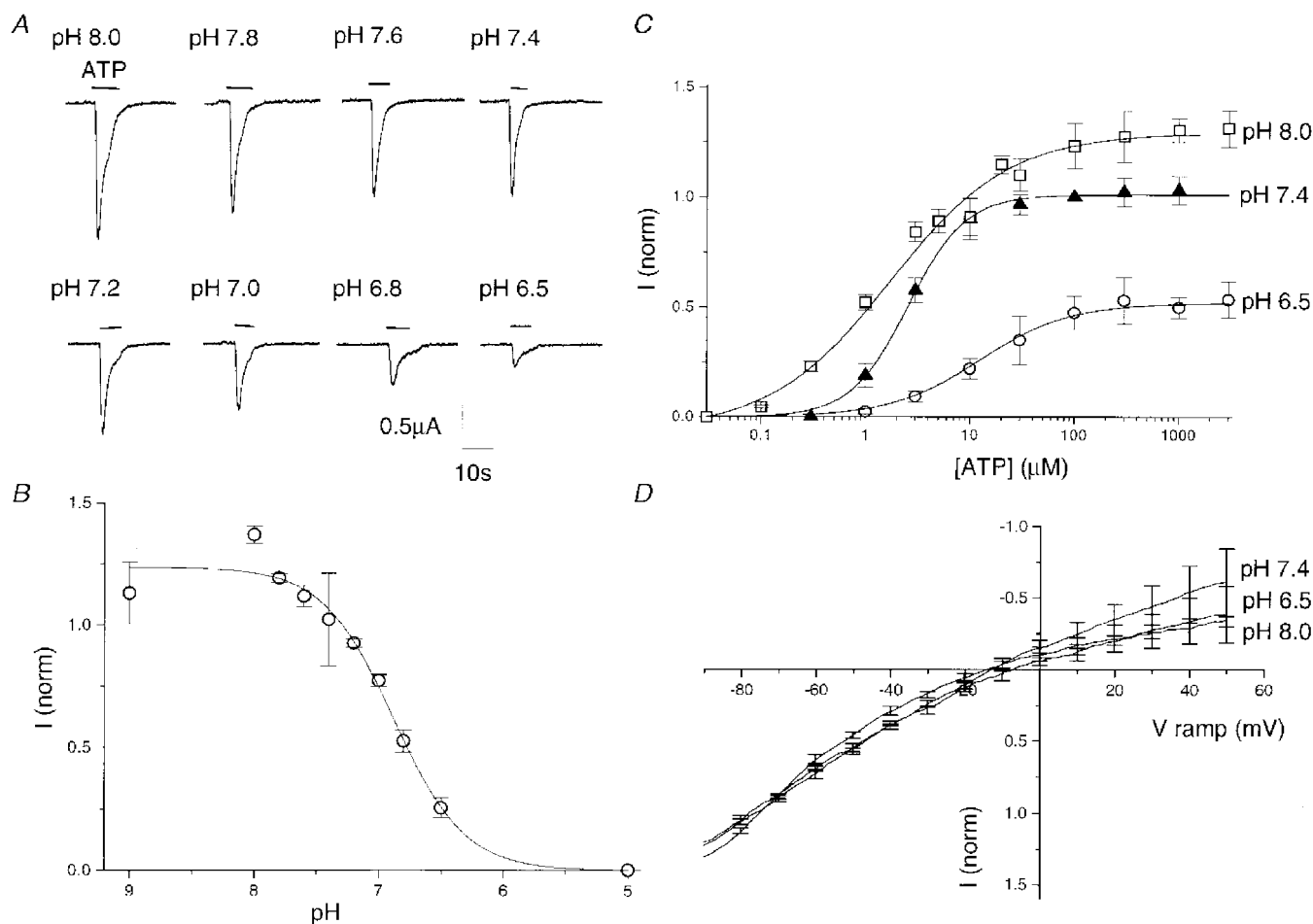

Figure 1. Effect of pH on the hP2X$_4$ receptor

A, example of responses of the hP2X$_4$ receptor to 300 μM ATP over a pH range from 8·0 to 6·5 recorded from a single oocyte. *B*, pH concentration–response curve. Responses evoked at 300 μM ATP over a range of pH from 9 to 5 were normalized to the response recorded in the same oocyte at 100 μM ATP, pH 7·4. Data plotted are means ± s.e.m. ($n = 4$). *C*, ATP concentration–response curves for the hP2X$_4$ receptor at pH 6·5 (O), 7·4 (▲) and 8·0 (□). Responses were evoked at a range of ATP concentrations from 0·03 μM to 3 mM and normalized to the response recorded in the same oocyte at 100 μM ATP, pH 7·4. Values plotted are means ± s.e.m. ($n \geq 5$). *D*, current–voltage relationship of the hP2X$_4$ receptor at pH 8·0, 7·4 and 6·5, with ATP EC$_{20}$. Currents were obtained by voltage ramps from −90 to +50 mV and normalized to the response obtained at −80 mV for each test pH. Data are shown as means ± s.e.m. ($n = 4$).

effect of short pre-exposure to DEPC was found on rP2X$_4$ by Stoop *et al.* (1997). Responses to 300 μM ATP were recorded at pH 6·5, 7·4 and 8·0 in control oocytes and those incubated overnight in 10 mM DEPC. Data were normalized to 100 μM ATP, pH 7·4. No significant differences could be found between the two groups at any of the test pH values (*n* = 4 for each test pH, data not shown).

The results with DEPC are not inconsistent with pH sensitivity being mediated through histidine protonation. A histidine imidazole might be accessible to protons but not the much larger DEPC molecule. To test this hypothesis, we continued with a mutagenesis approach. As the hP2X$_4$ amino acid sequence only contains four histidines, all predicted to be extracellular, site-directed mutagenesis of each of the histidines to alanines was carried out to form

mutants H140A, H172A, H241A and H286A (Fig. 2*A*). Each of the mutants and wild-type P2X$_4$ were injected into *Xenopus* oocytes and 1–2 days later responses to a maximal concentration of 300 μM ATP were tested at pH 6·5, 7·4 and 8·0 (Fig. 2*B*). While data for H140A, H172A and H241A overlaid wild-type responses, wild-type pH sensitivity was completely lost in mutant H286A. Responses to 300 μM ATP at pH 6·5 were similar in amplitude to those at pH 8·0 for H286A. ATP-induced responses of H286A, like wild-type channel, were rapidly desensitizing (Fig. 3*A*), but maximal currents were much smaller than those seen in wild-type channel (0·45 ± 0·17 and 2·78 ± 0·38 μA, respectively, *n* = 4). We also noted that expression was less efficient with H286A. Expression rates for H286A were improved by injecting 5 ng of cDNA for H286A rather than 1·5 ng but were still only 20–30% compared to 80–90% of oocytes
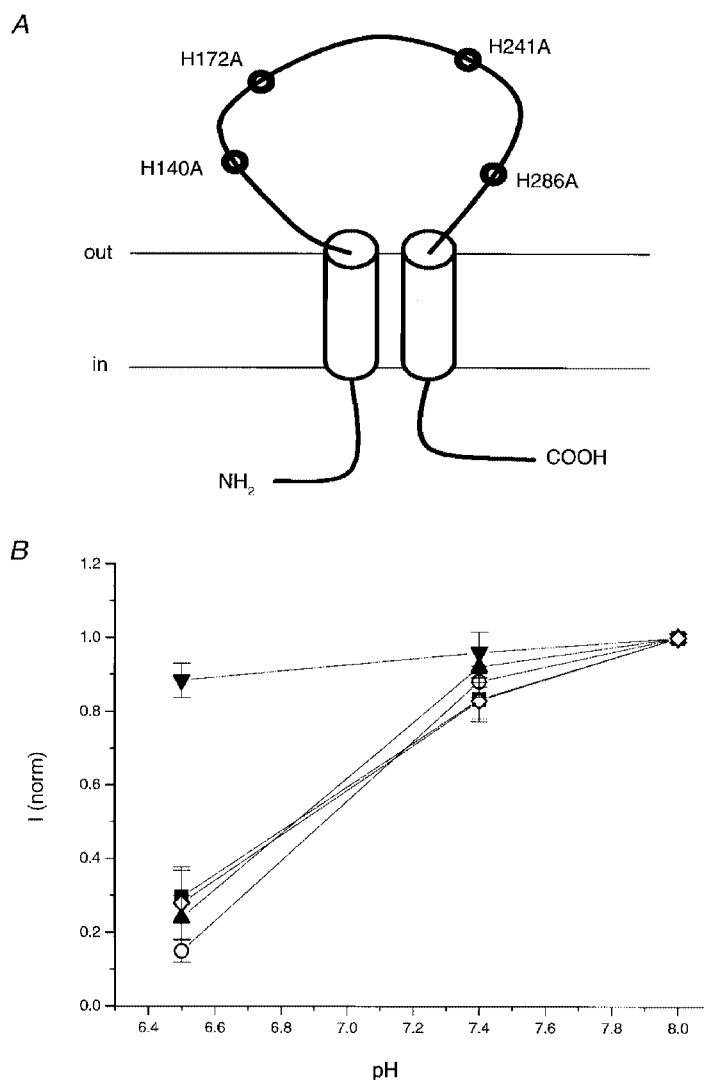

Figure 2. Effects of histidine mutagenesis on pH sensitivity of maximal ATP responses

A, diagram showing the four extracellular histidines present in the hP2X$_4$ amino acid sequence that were mutated to alanines. *B*, graph showing the responses of H140A (■), H172A (○), H241A (▲), H286A (▼) and wild-type hP2X$_4$ (◇) to 300 μM ATP at pH 6·5, 7·4 and 8·0. Responses have been normalized to the response obtained at pH 8·0. The actual values at pH 8 were: H140A, 3·05 ± 0·46 μA; H172A, 2·42 ± 1·03 μA; H241A, 3·3 ± 0·42 μA; H286A, 0·45 ± 0·17 μA and wild-type, 2·78 ± 0·38 μA. Values plotted and actual values are means ± S.E.M. (*n* = 4).

responding when injected with all other mutant types and
wild-type receptor. To further characterize the change in pH
sensitivity in the H286A mutant, full concentration–
response curves to ATP were carried out at pH 6·5, 7·4
and 8·0 (Fig. 3B). In contrast to wild-type (c.f. Fig. 1C)
concentration–response curves for H286A superimpose
showing that the pH sensitivity of the hP2X₄ receptor at all
ATP concentrations had been lost.

Comparison of the ATP concentration–response curves for
both wild-type and mutant H286A at pH 7·4 (Fig. 3C)
revealed that the mutation caused a fourfold decrease in
EC_{50} from 2·6 to 9·5 μM. The Hill slope for the H286A
concentration–response curve at pH 7·4 was reduced from
1·58 to 0·71. These data suggest that in addition to being
responsible for pH sensitivity H286 has a minor effect on
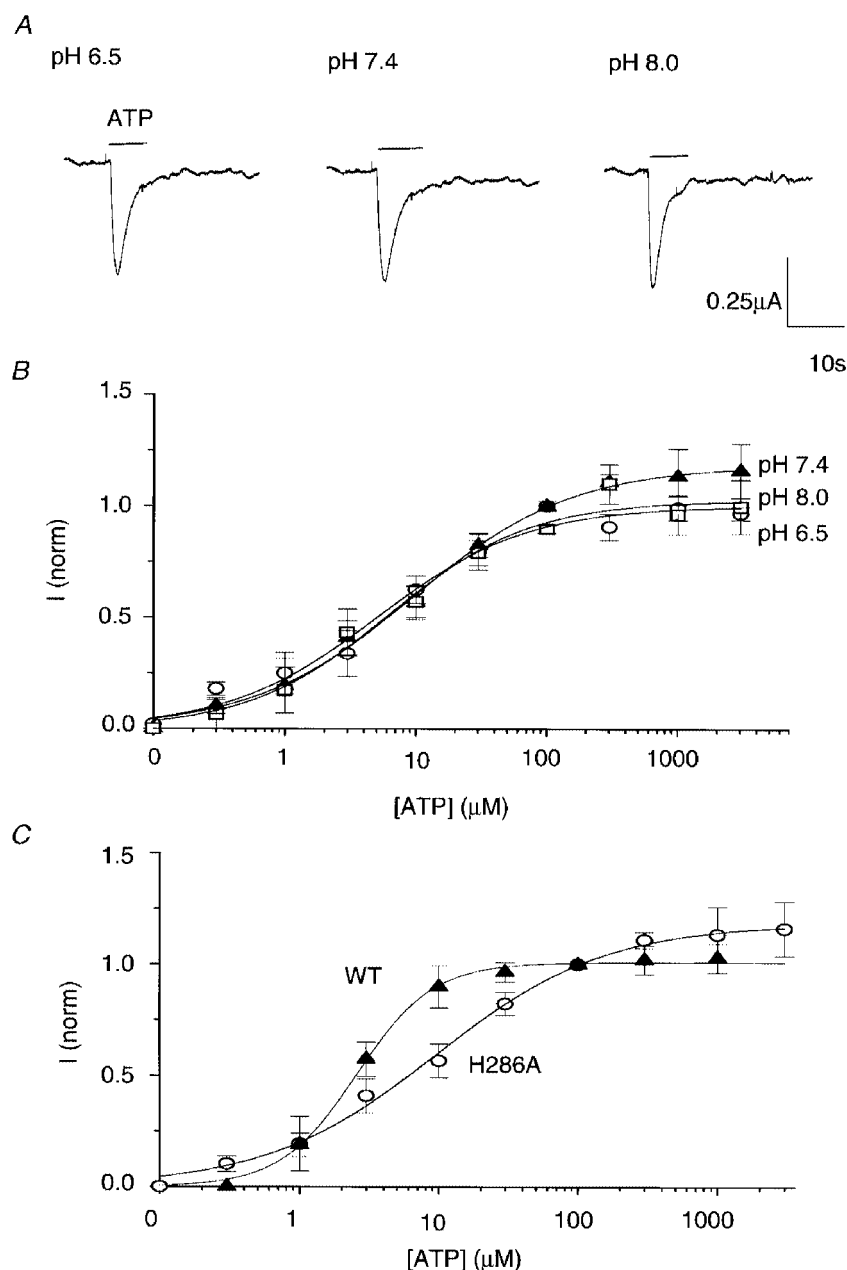ligand binding and/or gating.

Figure 3. The H286A mutant is pH insensitive

A, example of responses of mutant H286A at pH 6·5, 7·4 and 8·0. Recordings were taken from a single
oocyte held at −80 mV. *B*, ATP concentration–response curves for the H286A mutant at pH 6·5 (O), 7·4
(▲) and 8·0 (□). Responses were evoked at a range of ATP concentrations from 0·03 μM to 3 mM and
normalized to the response recorded in the same oocyte at 100 μM ATP, pH 7·4. Data plotted are
means ± s.e.m. ($n \geqslant 5$). *C*, plots of the concentration–response curves for ATP at pH 7·4 for wild-type (▲)
and mutant H286A (O). Values were normalized to responses at 100 μM ATP, pH 7·4. The curve for
mutant H286A is shifted approximately fourfold to the right compared with wild-type hP2X₄. ATP EC_{50}
values are 2·56 and 9·5 μM, respectively. Data are means ± s.e.m. ($n \geqslant 4$).

DISCUSSION

Our results show that $hP2X_4$ receptor ATP-activated currents are reduced by reducing extracellular pH and that this pH sensitivity is completely abolished by mutation of histidine 286 to alanine. Thus, protonation of a single histidine seems to confer pH sensitivity on this channel. The $rP2X_4$ is also pH sensitive but unlike the findings of Stoop *et al.* (1997) we found that the maximal response to ATP in $hP2X_4$ was also affected by pH changes. Recently published experiments on $rP2X_4$ extending to lower pH show a reduction in maximal ATP response indicating that there is not a major difference in the behaviour of the two isoforms (Wildman *et al.* 1999). Our mutagenesis data provide direct support for the hypothesis that pH sensitivity is due to protonation of the channel rather than protonation of the agonist ATP (Wildman *et al.* 1999).

Comparison of the sequences of rat and human $P2X_4$ (Rassendren *et al.* 1997) shows that H286 is conserved suggesting that H286 might also mediate the pH sensitivity in the rat isoform. The observation that the $rP2X_4$ shows different pH sensitivity permitting full agonist responses to ATP at pH 6·3, might be explained by the subtly altered consequences for gating, dependent on other non-conserved amino acids. $hP2X_1$ and $rP2X_3$ channels also show similar but less dramatic pH sensitivity and have ten and two extracellular histidines, respectively. Pairwise comparisons show that none of these are precisely homologous to H286 on $hP2X_4$. Interestingly, $rP2X_5$ does have an analogous histidine (Collo *et al.* 1996), but the pH sensitivity of this channel has not been explored.

Knowledge of the structure of the P2X family of channels is to date limited. Hydrophobicity plots suggest that each subunit has two transmembrane (TM) regions and the lack of a signal peptide sequence suggests a structure with intracellular N- and C-terminals and a large extracellular loop between the two TM regions. This basic motif is supported by studies of N-glycosylation sites (Newbolt *et al.* 1998; Torres *et al.* 1998*a*,*b*). Cysteine mutagenesis of the second TM domain of $P2X_2$ suggests that hydrophilic residues on this helix form part of the pore lining and that the N-terminal end of this region (positions 328–336) define part of the outer vestibule (Rassendren *et al.* 1997; Egan *et al.* 1998). The large extracellular loop codes for the ATP binding site and by analogy to other ion channels may contain a pore-forming loop that provides further structural elements of the pore (Brake *et al.* 1994). Thus the location of H286 relative to the pore cannot be defined at present. The absence of voltage dependence of pH sensitivity (Fig. 1*D*) indicates a site extracellular to the membrane field, consistent with the location of H286 in the extracellular loop. The lack of voltage sensitivity is also clear evidence that the pH effect is not mediated by proton block of the permeation pathway. The lack of effect of DEPC might be explained by lack of accessibility to the H286 protonation site suggesting that

the H286 imidazole projects into a restricted steric space on the extracellular surface. Based on this limited data it seems most likely that the pH modulation is mediated by an allosteric conformational change that affects gating rather than by a direct interaction with the channel gate itself.

Many ion channels are modulated by pH but for relatively few has the molecular basis been determined. The H452Q mutant of the Kv1.5 channel shows about 50% reduction in sensitivity to low extracellular pH (Steidl & Yool, 1999). Extracellular histidines also affect the pH sensitivity of the inward rectifier potassium channel, HIR (Coulter *et al.* 1995). However, for other channels regulated by pH, glutamate residues have been shown to be important such as the L-type voltage-gated Ca^{2+} channel where several glutamate residues that line the pore affect pH sensitivity (Chen *et al.* 1996), and CNG (cyclic nucleotide gated) channels (Rho & Park, 1998).

Human $P2X_4$ mRNA shows widespread distribution; it was detected in many tissues analysed by Garcia-Guzman *et al.* (1997), and it is the predominant P2X receptor, along with $P2X_6$ receptor in the CNS (Buell *et al.* 1996*a*). The CNS is exposed to both transient shifts in pH with normal neuronal activity (Chesler & Kaila, 1992), and more sustained acidosis with ischaemia and hypoxia in the brain (Siesjö *et al.* 1996). ATP may be co-released as a transmitter during normal CNS function and in greater amounts during ischaemia. The pH sensitivity of $P2X_4$ channels may naturally limit signalling and toxicity mediated by this pathway during acidosis and might, together with the pH sensitivity of NMDA-gated channels (Traynelis & Cull-Candy, 1990), explain why increases in extracellular alkalinity have been shown to exacerbate injury of cultured cortical neurons subjected to both glutamate neuronal toxicity and oxygen–glucose deprivation (Giffard *et al.* 1992). In this paper we have shown that ATP-gated $hP2X_4$ channels are acutely sensitive to extracellular pH and that the basis for this sensitivity is protonation of histidine 286.

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