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Key words: P2X; dynamin; endocytosis; recycling; presynaptic; trafficking; heteromer; ATP; ionotropic receptor

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Department of Pharmacology, University of Cambridge, Cambridge CB2 1PD, United Kingdom

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Correspondence should be addressed to Ruth D. Murrell-Lagnado, Department of Pharmacology, Tennis Court Road, University of Cambridge, Cambridge CB2 1PD, UK. E-mail: rdm1003@cam.ac.uk.

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Initial plating density was 6.7 cm⁻² per OB. For the first 3 d of growth, the culture medium also included 10% horse serum. Serum-free medium containing cytosine arabinoside (ARA-C; 2 μM) was subsequently renewed every 3 d.

HEK293 cells were maintained in DMEM (NUT.MIX.F12) containing 10% fetal bovine serum (FBS) and 100 U/ml penicillin–streptomycin at 37°C and 5% CO₂.

DNA constructs. Initially, two P2X4 constructs were generated with GFP at either the N or C terminus. However, fusing GFP to the N terminus of P2X4 (GFP-P2X4) dramatically inhibited expression in Xenopus oocytes and in OB neurons, and therefore this construct was not studied further. To generate cDNAs encoding P2X2, P2X4, and P2X6 with GFP fused to the C terminus, the rat cDNAs (kind gifts from Prof. P. P. A. Humphrey, Cambridge, UK) were amplified by PCR using oligonucleotide primers designed to introduce a Kozak (1987) initiation sequence, nonpermeabilization of the stop codon, and an XbaI restriction site. AU5 (TDFYLK) was added to the 5' ends, respectively. Amplification products were then cloned into the pEGFP-N1 vector (Clontech, Palo Alto, CA).

For expression of either wild-type or epitope-tagged P2X receptors, the coding sequences of P2X2 and P2X4 were amplified to reintroduce the stop codon and an XbaI site. The subcloning of these fragments excluded the coding sequence of the first 60 amino acids, causing a frameshift that substituted into P2X4 at position 76 (TSQGLF) by two-step PCR. Epitope FLAG (DYDDDDK) was inserted into P2X2 between D78 and K79 (Stoop et al., 1998). The sequences of all amplified regions were verified by automated DNA sequencing. The sequences of all primers used are available on request.

Hemagglutinin (HA)-tagged human dynamin-1 and -dynamin-1(K44A) cDNAs in pcDNA3 expression vector were kind gifts from Dr. J. M. Edwardson (Cambridge, UK).

Cell transfection. OB neurons were transfected at 10–21 d in vitro (DIV) using a modified calcium phosphate method (Xia et al., 1996). Briefly, precipitates were formed by adding 6 μl of plasmid DNA to 60 μl CaCl₂ (250 mM) to which 60 μl of 2x HEPS-buffered saline (2x HEPS, 10 KCl, 1 K4HPO₄, 10 d-glucose, and 10 HEPES, pH 7.07) was added dropwise. Precipitate (55 μl/21-mm diameter well) was added to the cells whose culturing medium was replaced with transfection buffers (in mm: 10 MgCl₂, 10 HEPS, 0.5 L-glutamine, and 100 U/ml penicillin–streptomycin in Neurobasal medium, pH 7.2). Optimization of the volume of precipitate and pH of HEPS, removal of endotoxins from DNA solutions, and duration of the incubation (2 hr) resulted in low levels of toxicity and transfection efficiency of >5%. After transfection the neurons were washed with transfection buffer and then kept in the previous culturing medium for up to 2 d. For cotransfection experiments, equal amounts of DNA were used to a total 6 μg.

HEK293 cells were plated onto poly-d-lysine-treated coverslips (2.4 × 10⁻⁶ cells/cm²) and transfected 12 hr later using the same method with slight modifications. The amount of DNA used to form precipitate was 5 μg (in 100 μl of 2x HEPS) and 100 U/ml penicillin–streptomycin (21-mm diameter well) was added, and cells were kept with precipitate for 6 hr. For cotransfection experiments, equal amounts of DNA were used, and controls for lone expression had the total DNA. We included 0.5 μg of pEGFP-N1 vector for coexpression of GFP with nonfluorescent constructs.

Electrophysiological recordings. Standard whole-cell recordings (Hamill et al., 1981) were performed at room temperature (RT) using an Axopatch 200A amplifier (Axon Instruments, Foster City, CA). Patch pipettes (3–8 MΩ) were pulled from thick-walled borosilicate glass (GC150F-10; Harvard Apparatus, Holliston, MA) and filled with solution containing (in mm): 125 K-glucurate, 1 MgCl₂, and 10 HEPES, pH 7.3. The extracellular solution was composed of (in mm): 140 NaCl, 5 KCl, 2 CaCl₂, 1 MgCl₂, 10 d-glucose, and 10 HEPES, pH 7.3. ATP-evoked responses were measured at ~60 mV in neurons and at ~50 mV in HEK293 cells.

To visualize cells expressing P2X receptors without a GFP tag, cells were cotransfected with GFP. Cells expressing GFP or GFP-tagged P2X subunits were observed under a microscope with an epifluorescence binocular (Nikon, Tokyo, Japan). Untransfected cells and cells expressing GFP alone were found to have no inward current in response to application of ATP. ATP was applied locally using a Picospritzer II (Parker Instrumentation, Fairfield, NJ). To ensure delivery of drug, 0.05% (w/v) fast green was used (local applications of 1% fast green evoked no response).

The effects of ATP on the frequency of miniature postsynaptic currents (mPSCs) were measured in neurons that were >14 DIV before transfection and that displayed spontaneous activity, indicating that they had formed synaptic connections. Whole-cell recordings were made from untransfected neurons that neighboring transfected neurons. Pairs of neurons were chosen on the basis that they were in close proximity but not immediately adjacent to one another. ATP was applied where their processes appeared to be in intimate contact. As a control, pairs of untransfected neurons were chosen, and ATP was applied to see if activation of endogenous receptors could elicit a change in mPSC frequency. In all cases, neurons were held at ~40 mV, so that miniature EPSCs (mEPSCs) and miniature IPSCs (mIPSCs) appeared as inward and outward currents, respectively. The extracellular solution contained tetrodotoxin (TTX; 1 μM). In some experiments either 6-cyano-7-nitroquinoline-2,3-dione (CNQX; 10 μM) or bicuculline (20 μM) were coapplied with ATP to block fast glutamatergic and GABAergic transmission, respectively.

Whole-cell currents were low-pass filtered at 2 kHz and digitized at 10 kHz. Acquisition was performed using Heka (Lambrecht, Germany) PulsP 8.30, and data were subsequently analyzed using IgorPRO 3.14. Concentration–effect curves were fitted to the Hill equation. The frequency of mEPSC was determined during 20 sec time interval before and during ATP application. Events were selected manually on the basis of their amplitude and duration.

Immunocytochemistry. Cells were fixed in 3% paraformaldehyde and 4% sucrose in PBS (in mm: 1.5 NaH₂PO₄, 8 Na₂HPO₄, and 145 NaCl, pH 7.3) for 10 min at 4°C. If required, permeabilization was done using 0.1% Triton X-100 in PBS for 10 min at 4°C. Nonspecific sites were blocked using PBS containing 4% normal goat serum and 3% bovine serum albumin (blocking buffer). Antibodies were diluted to their final concentration in blocking buffer. Primary antibodies were applied for 2 hr at RT.

Cells were rinsed once in blocking buffer and three times for 5 min with PBS, and then secondary antibodies were applied for 2 hr at RT. Finally, cells were washed five times for 5 min with PBS and mounted onto slides with Vectashield (Vector Laboratories, Burlingame, CA) as a mounting medium.

Representative images of immunocytochemical experiments were obtained using a Bio-Rad (Hercules, CA) MRC1024, Zeiss Axioplan LSM 510 or Perkin-Elmer UltraVIEW confocal microscope, with 60×, 63×, or 100× oil immersion objectives, respectively. Fluorescein isothiocyanate (FITC)–GFP, Cy3–tetramethylrhodamine isothiocyanate (TRITC), and Cy5 were excited at 488, 543, and 633 nm, respectively. When either two or three fluorophores were visualized, separate excitation and collection was used to minimize bleedthrough.

Live-labeling immunofluorescence protocols. The basic protocol for live-labeling was as follows: cells were incubated with the lysosomal protease inhibitor leupeptin (100 μg/ml), for 1 hr before incubation with anti-AU5, and leupeptin was included throughout all steps. Anti-AU5 was applied for 30 min at 37°C in serum- and supplement-free culture medium as a buffer. Cells were then washed five times and fixed in PFA, 1%, for 15 min. The surface of untransfected neurons were stained with an anti-mouse Cy3-conjugated secondary for 2 hr at RT. Cells were then washed five times in PBS, permeabilized with 0.1% Triton X-100, and stained with an anti-mouse FITC-conjugated secondary for 2 hr at RT, to visualize prelabeled internalized receptors.

To estimate the time course of receptor internalization, neurons were incubated with anti-AU5 for 10, 20, 30, or 60 min at 37°C. After fixation, cells were double-labeled using two different methods. First, to determine the ratio between surface receptors and the total receptor population, nonpermeabilized cells were stained with an anti-mouse Cy5-conjugated secondary. After permeabilization, neurons were stained for total receptor using anti-P2X4 and an anti-rabbit Cy3-conjugated secondary. Second, to determine the ratio between AU5-labeled receptors (surface and internalized) and the total receptor population, neurons were permeabilized before staining with anti-mouse Cy5-conjugated secondary.

For recycling experiments, HEK293 cells expressing P2X4(AU5) were incubated with anti-AU5 as described above. Cells were washed five times in buffer before being gradually cooled to 4°C to inhibit trafficking. Cells were then incubated with nonfluorescent anti-mouse antibody for 1 hr at 4°C to block AU5-labeled receptors at the surface. Cells were either fixed at this point or returned to 37°C for 15 min to allow internalized receptors to recycle back to the surface. Cells were then fixed, and the surface and internalized receptors were detected with Cy3- and FITC-conjugated secondary antibodies, respectively.

To examine the effects of ATP and extracellular Ca²⁺ on receptor...
neurons were labeled as above, washed five times in buffer, and subsequently incubated for 15 min at 37°C with or without ATP (100 μM) in either a control solution or in a Ca²⁺-free solution. The control solution contained (in mM): 140 NaCl, 5 KCl, 2 CaCl₂, 1 MgCl₂, 10 d-glucose, and 10 HEPES, pH 7.3. For Ca²⁺-free, calcium was omitted, and 1 mM EGTA was added. Prelabelled receptors at the surface and internalized receptors were detected as above. The effect of agonist on P2X₂-GFP-expressing neurons was assessed by incubating transfected neurons in control solution for 15 min at RT with either ATP or 2-methylthioadenosine 5'-triphosphate (2meSATP; 100 μM). In some experiments, before ATP application, P2X₂-GFP-expressing neurons were preincubated for 10 min with pyridoxal-phosphate-6-azophenyl-2', 4'-disulfonic acid (PPADS; 10 μM) or suramin (100 μM). Cultures were subsequently fixed and stained for microtubule-associated protein-2 (MAP-2).

**Image analysis.** Fluorescence was visualized using a Zeiss Axiosvert LSM510 confocal microscope, using 63× oil immersion objective. For FITC–Cy3 anti-AU5 double labeling, FITC and Cy3 were excited at 7 and 60% of 488 and 543 nm laser power, respectively. For Cy3–Cy5 anti-AU5 and anti-P2X4 double labeling, Cy3 and Cy5 were excited at 4 and 70% of 483 and 633 nm laser power, respectively. For the P2X4(AU5)/dynamin-1 conditions, FITC and Cy3 were excited at 7 and 49% of 488 and 543 nm laser power, respectively. In P2X4(AU5)/dynamin-1(K44A) conditions, there was greater P2X4(AU5) labeling; accordingly, we reduced both laser powers to 1/4 (1.75 and 12.25%). A series of confocal sections (1 μm thickness) were taken of immunostained neurons at 0.14 μm/pixel. For individual experiments, images for all conditions were analyzed using identical acquisition parameters.

Tagged image file format (TIFF) images were imported into NIH Image 1.62, and the cells outlined and mean pixel values for each channel were obtained for a region over the cell soma. The images were selected so that confocal plane was focused on the middle of the cell to exclude signal from the top and bottom of the cell. Pixel values were on an 8-bit scale (2^8 = 256; 0–255). For time course experiments, ratios of either surface versus total expressed receptor (Cy5 vs Cy3) or labeled versus total expressed receptors (Cy5 vs Cy3) were obtained.

For recycling and internalization experiments, red (Cy5) or green (FITC) fluorescence intensities indicative of surface or internalized receptors were divided by total labeled (red plus green) fluorescence intensities, respectively. Units of surface or internalized receptor were measured as red per total or green per total fluorescence normalized to untreated controls. For dynamin experiments, the ratio of surface versus total expressed receptor (Cy3 vs FITC) was obtained. Experiments were repeated at least two times, and each time, data were analyzed from four to six cells from two separate coverslips. Within each experiment the data were normalized to control cells. The n value given refers to the number of cells analyzed. All data are presented as the mean ± SEM. Histograms and plots were done using IgorPRO 3.14 software.

Statistical analyses were performed with Student’s unpaired t test or ANOVA with Bonferroni’s post hoc comparison using InStat software (version 2.01; GraphPad Software Inc., San Diego, CA). All results are given as mean ± SEM, and levels of statistical significance are *p < 0.05, **p < 0.01, and ***p < 0.001.

**Antibodies and reagents.** The following primary antibodies were used: affinity-purified anti-AU5 (5 μg/ml; Babco, Richmond, CA), anti-FLAG M2 (2 μg/ml; Sigma, St. Louis, MO), anti-synaptobrevin (1:750; a kind gift from Dr. J. M. Edwardson, Cambridge, UK), anti-EA1-1 (2.5 μg/ml; Transduction Laboratories, Lexington, KY), anti-MAP-2 (1:250; Roche Products, Hertfordshire, UK), and anti-hemagglutinin (0.8 μg/ml; Roche) all mouse monoclonal; anti-P2X4 (6 μg/ml; Alomone Labs, Jerusalem, Israel), anti-P2X2 (0.6 μg/ml; Alomone), anti-lgp110 (1:250; a kind gift from Dr. J. P. Luzio, Cambridge, UK), and anti-calciretinucin (1:200; a kind gift from Dr. P. Thorn, Cambridge, UK) all rabbit polyclonal, FITC-, Cy3-, TRITC-, and Cy5-conjugated goat anti-mouse, anti-rabbit (1:250), or non-fluorescent goat anti-mouse IgG (1:100) were used as secondary antibodies (Jackson Immunoresearch, West Grove, PA).

TTX (5 mM stock solution in ethanol), CNOX (50 mM stock solution in DMSO), bicuculline (80 mM stock solution in DMSO), PPADS, and leupeptin (10 mg/ml) were purchased from Tocris Cookson (Bristol, UK). Unless otherwise stated, all other reagents were obtained from Sigma or Invitrogen (San Diego, CA).

**RESULTS**

Subunit-dependent trafficking of P2X receptors expressed in HEK293 cells

To allow visualization of P2X receptor trafficking in live cells, we introduced AU5 and FLAG epitopes into the extracellular loops of P2X4 and P2X2, respectively, and we separately tagged both subunits with GFP at their C terminus. Wild-type and tagged receptors were expressed in HEK293 cells to compare their distributions and functional properties. The distributions of the tagged receptors were very similar to their wild-type counterparts (Fig. 1A–F). The P2X4 receptors all had a punctate distribution, and confocal images indicated that most of the clusters were intracellular (Fig. 1A). In contrast, P2X2 receptors were more uniformly distributed and predominantly at the periphery of the cell (Fig. 1B,D,F). Using epitope-tagged receptors, those at

Figure 1. Subcellular distribution and functional properties of P2X constructs in HEK293 cells. A–F, The subcellular distribution of P2X4 WT (A), P2X4–GFP (C), P2X4(AU5) (E), P2X2 WT (B), P2X2–GFP (D), and P2X2(FLAG) (F) receptors in HEK293 cells. Cells expressing untagged receptors were stained using anti-P2X4/Cy3 or anti-P2X2/Cy3. Cells expressing P2X4(AU5) were stained using anti-AU5/Cy3 before (E, left panel) and anti-P2X4/FITC after permeabilization. Cells expressing P2X2(FLAG) were stained using anti-FLAG/Cy3 before (F, left panel) and anti-P2X2/FITC after permeabilization. Scale bar, 10 μm. G, Concentration–effect curves for ATP and representative traces showing ATP-evoked whole-cell currents for P2X4 (left) and P2X2 (right) constructs used in this study. ATP-evoked peak currents shown are normalized to compare the time course of desensitization. Calculated EC₅₀ values were 3.8, 4.2, and 17 μM for P2X4 WT, P2X4–GFP, and P2X4(AU5), respectively. Calculated EC₅₀ values for P2X2 WT, P2X2–GFP, and P2X2(FLAG) were 3.9, 4.4, and 14.6 μM, respectively (n = 3–7 cells for each concentration).
the surface were labeled in addition to the total receptor population. For P2X2(FLAG), there was considerable overlap between the two labels, whereas for P2X4(AU5), the majority of puncta were not at the surface (Fig. 1E,F).

Using whole-cell patch clamp we compared the functional responses to ATP for all six receptors (Fig. 1G). The time courses were similar for the tagged receptor currents and their wild-type counterparts. P2X2-GFP and P2X2(FLAG) receptor currents were slowly desensitizing, whereas P2X4-GFP, P2X4(AU5) and wild-type P2X4 receptor currents desensitized more rapidly. For each of the three P2X4 receptors, the decay was biphasic, and there was no significant difference in the values of the fast and slow time constants (τslow = 1.1 ± 0.3 sec, 1.4 ± 0.4 sec, and τfast = 9.5 ± 3.1 sec, 8.9 ± 0.5 sec, 11.1 ± 3.0 sec; n = 4 for P2X4, P2X4-EGFP, and P2X4(AU5), respectively). P2X4 and P2X4-GFP currents also had a similar EC50 values and peak current density. However, for P2X4(AU5) and P2X2(FLAG) receptors, the introduction of epitopes just downstream of the first transmembrane regions produced a shift in the concentration–effect relationship to higher concentrations of ATP. The peak current density for P2X2-GFP was also 1.4-fold higher than for the wild-type receptor. We conclude that tagging the receptors does not fundamentally disrupt their functional expression and that the trafficking of P2X2 and P2X4 receptors in HEK293 cells is different.

**Constitutive endocytosis of P2X4 receptors expressed in OB neurons**

We next compared the distribution of wild-type and tagged P2X2 and P2X4 receptors in dissociated OB neurons that were cultured for between 1 and 3 weeks. These neurons showed no response to ATP (n = 129) and were devoid of endogenous P2X2 or P2X4 immunoreactivity. Both P2X2 and P2X4 receptors were expressed throughout the cell (Fig. 2A,D). Similar to HEK293 cells, the principal difference between the two subtypes was that P2X4 receptors were clustered into discrete puncta, whereas P2X2 receptors were not (Fig. 2B,C,E,F). This difference was apparent as early as 4 hr after transfection when protein expression was far lower. We also transfected cultured hippocampal neurons from embryonic rats and obtained the same result: P2X4 was punctate and intracellular, whereas P2X2 was uniform and at the plasma membrane (results not shown). In P2X2-transfected OB neurons, two rails of fluorescence were apparent along large dendrites, suggestive of receptors at the plasma membrane (Fig. 2E,F). Whole-cell ATP-evoked peak current densities were 25.5 ± 5.2 pA/pF (n = 15) and 285.8 ± 51.4 pA/pF (n = 6) for OB neurons transfected with P2X4-GFP and P2X2-GFP, respectively.

Many of the P2X4-GFP fluorescent puncta within the processes colocalized with early endosomal antigen-1 (EEA-1), a marker for early endosomes (Fig. 3A). There was also correspondence between puncta in the soma and proximal dendrites and a marker for lysosomes (lgp110) (Fig. 3B). In contrast, there was no colocalization between P2X2 receptors and either EEA-1 or lgp110 (data not shown). Consistent with the presence of P2X4 receptors in early endosomes, we were able to directly visualize the internalization of surface-labeled P2X4(AU5) receptors in live neurons. After a 30 min incubation with anti-AU5 at 37°C, surface receptors were in discrete clusters around the soma and along the processes, and internalized receptors were present throughout the neuron (Fig. 3C). Similar experiments were performed with neurons transfected with P2X2(FLAG). Consistent with the lack of endosomal P2X2 receptors, the internalization of P2X2(FLAG) was negligible, whereas expression at the surface was high (Fig. 3D).

We estimated the time course of P2X4(AU5) receptor internalization in neurons, by incubating cells with anti-AU5 for different periods of time. After fixing neurons, either the total population of AU5-labeled receptors (labeled AU5) or just the surface receptors (surface AU5) were detected. In both cases the total receptor population was independently labeled using anti-P2X4 (total P2X4). As shown in Figure 4A, surface AU5 fluorescence did not change significantly between 10 and 60 min. Therefore, we assume that the increase in labeled AU5 fluorescence over time represents the rate at which AU5-labeled receptors were internalized. The time course was fitted by a single exponential curve with a time constant of 21 min. The binding of

**Figure 2. Subcellular distribution of heterologously expressed P2X receptors in neurons.** Confocal images of heterologously expressed P2X4 WT (A, B), P2X4–GFP (C), P2X2 WT (D, E), and P2X2–GFP (F) receptors in mature OB neurons. Insets, C, F. Example traces of whole-cell currents evoked by a 0.5 sec application of ATP (100 μM). Cells expressing untagged receptors were stained using anti-P2X4/Cy3 and anti-P2X2/Cy3, respectively. Scale bars: A, B, D, E, 10 μm; C, F, 5 μm.
the antibody did not change the surface expression of receptors in live cells. The ratio of surface AU5- to total P2X4-staining was comparable after a 60 min incubation with anti-AU5 at 37°C compared with applying anti-AU5 after fixing cells (0.27 ± 0.03; n = 5 vs 0.25 ± 0.08; n = 3, respectively).

**Recycling of P2X4 receptors to the surface**

In neurons, the rate at which P2X4 receptors were labeled with anti-AU5 approached a plateau, suggesting that internalized receptors recycled back to the surface. We tested this directly in P2X4(AU5)-transfected HEK293 cells by blocking AU5-labeled surface receptors with a nonfluorescent secondary antibody at 4°C and returning cells to 37°C to allow receptor trafficking to continue. There was a 1.5-fold increase in the cell surface fluorescence after 30 min, indicating that internalized receptors had recycled back to the plasma membrane (Fig. 4B,C).

**Agonist-regulated internalization of P2X4 receptors**

The constitutive cycling of P2X4 receptors into and out of the membrane suggests a mechanism for rapidly upregulating or downregulating the functional expression of receptors. Previously it was shown that application of ATP increased the internaliza-
We therefore tested the effects of ATP on P2X4 trafficking of smooth muscle P2X1 receptors (Ennion and Evans, 2001). We investigated the effects of ATP on P2X4 trafficking in neurons by live labeling AU5-tagged receptors followed by incubation with or without ATP. A 15 min incubation with 100 μM ATP caused an ~60% increase in the fraction of P2X4(AU5) receptors that were internalized. Removing external Ca2+ had no effect on basal or ATP-induced internalization, indicating that the effects of ATP binding were not mediated by calcium influx (Fig. 5A,B). We used the same approach to look at the effects of ATP on the internalization of surface P2X2(FLAG) receptors. After a 15 min incubation with ATP, there was no change in the proportion of P2X2(FLAG) receptors that were internalized (Fig. 5C). Thus, ATP binding promotes the internalization of P2X4 but not P2X2 receptors.

In a previous study, P2X2-GFP receptors expressed in hippocampal neurons were reported to redistribute within the membrane in response to ATP (Khakh et al., 2001a). To examine the redistribution of receptors, we imaged P2X2-GFP-transfected neurons in real time before and after application of ATP. Within <1 min after application of either 10 or 100 μM ATP, we observed a beading effect in which clusters of P2X2-GFP fluorescence appeared to form. This effect was not dependent on external Ca2+, and beading also occurred with the P2X2 receptor agonist, 2meSATP. Preincubation with the antagonists, PPADS or suramin, completely abolished the beading effect (results not shown). Previously, application of glutamate to cortical neurons has been shown to cause varicosity formation in dendrites as a result of an influx of Na+ (Hasbani et al., 1998). To determine whether ATP caused a change in the morphology of the dendrites themselves, we stained with an antibody to the dendritic protein, MAP-2. In control cells, MAP-2 staining was uniform along the length of the processes, whereas in cells exposed to agonist, it occurred in beads that corresponded exactly with the beads of P2X2-GFP fluorescence (Fig. 5D,E). Thus, our results suggest that the redistribution of P2X2–GFP reflects a change in the morphology of the dendrites and that P2X2–GFP is simply reporting this change.

**Dynamin-dependent endocytosis of P2X4 receptors**

To investigate the mechanism of P2X4 receptor internalization, we tested the effects of a dominant-negative mutant form of dynamin-1 [dynamin-1(K44A)] on the distribution of the receptor. The effect of inhibiting dynamin-dependent endocytosis with this mutant was essentially to cause the distribution of P2X4 receptors to become similar to that of P2X2 receptors. Coexpression of dynamin-1(K44A), but not wild-type dynamin-1, changed the distribution from punctate to uniform, suggesting that internalization of receptors into endosomal compartments was inhibited (Fig. 6A,B). It also caused a dramatic increase in the expression of P2X4(AU5) receptors at the cell surface, as shown by labeling with extracellular anti-AU5 (Fig. 6C–F). Consistent with the increase in surface immunofluorescence, the ATP-evoked whole-cell currents recorded from P2X4-GFP-transfected neurons were also significantly increased (p < 0.01) by coexpression of dynamin-1(K44A) (Fig. 6G). Nevertheless, the current densities of P2X4-transfected neurons after coexpression with dynamin-1(K44A) were still approximately fourfold less than for P2X2-transfected neurons. One reason for this is that the single-channel conductance is more than twofold higher for P2X2 than for P2X4 receptors (9 vs 21 pS; Evans, 1996). Not surprisingly, coexpressing dynamin-1(K44A) with P2X2–GFP had no obvious effect on either the distribution of the receptor or on the amplitude of the ATP-evoked whole-cell currents (Fig. 6H). Thus, similar to other ligand-gated ion channels, the internalization of P2X4 receptors involves a dynamin-dependent mechanism.

**P2X receptors act presynaptically to regulate glutamatergic transmission**

Other ligand-gated ion channels, such as the AMPA and GABA<sub>A</sub> receptors, cycle into and out of the postsynaptic membrane. Within the adult rat OB, P2X4 immunoreactivity was shown to be associated with both dendrites and axon terminals, suggesting a presynaptic and postsynaptic role. In transfected mitral cells, where the dendrites and axon were clearly distinguishable, both P2X2 and P2X4 receptors were present throughout the cell, and there was no indication of specific targeting to either only somatodendritic or axonal compartments. OB neurons in situ have dendrodendritic reciprocal synapses, and therefore both presynaptic and postsynaptic structures are present within the dendrites. Some of the puncta of P2X4–GFP receptors colocalized with synaptobrevin-immunopositive structures, suggesting that receptor clusters were in the vicinity of synapses (Fig. 7A). We looked...
ATP (100 μM/H9262) tested) (Fig. 7) evoked an increase in the frequency of mEPSCs (9 of 25 cells), indicating that they were mediated by AMPA receptors (Fig. 7C), whereas application of bicuculline had no effect. Previous studies have shown that OB neurons of adult rat also express P2X2 receptors (Kanjhan et al., 1999). However, in both freshly plated and cultured OB neurons of rat neonates, P2X2 receptor expression was below the level of detection, suggesting that the expression of P2X2 is upregulated during development. Similarly, cultured hippocampal neurons from embryonic rats show no ATP-evoked responses, whereas hippocampal neurons in brain slices from adult rat have robust P2X-mediated currents (Khakh et al., 1999).

**P2X4/6 heteromeric receptors in HEK293 cells have a distribution similar to that of P2X4 homomeric receptors**

The endogenous receptors may be heteromeric or homomeric complexes. P2X4 forms heteromeric receptors with P2X6 but not with P2X2, and in the CNS, P2X4 and P2X6 have an overlapping distribution (Khakh et al., 2001b; Rubio and Soto, 2001). To test whether the distribution of the heteromer is determined by the P2X4 subunit, we expressed P2X6-AU5 receptors alone and together with P2X4-AU5 in HEK293 cells. P2X6 does not form functional homomeric receptors (Khakh et al., 1998), but expression of P2X6–GFP receptors was at the periphery of the cell, whereas P2X6–GFP fluorescence was diffusely distributed within the cell and colocalized with calreticulin, a marker for the endoplasmic reticulum, suggesting that the majority of receptors were not delivered to the cell surface (Fig. 9A). In contrast, P2X6–GFP coexpressed with P2X4 had a punctate distribution that resembled that of homomeric P2X4 receptors. It colocalized with anti-EEA1 and anti-P2X4 immunoreactivity (Fig. 9B). We also coexpressed P2X2–GFP with P2X4 as a control, to test whether the coexpression of subunits might influence the distribution of each other, even without their coassembly. In fact we saw no change in the distribution of either receptor (Fig. 9C). P2X2–GFP receptors were at the periphery of the cell, whereas P2X4 receptors were predominantly in intracellular puncta, and...
there was no indication that P2X2 receptors were endocytosed along with P2X4 receptors, suggesting that they might be excluded from sites of internalization. This result also confirms that the differences in the trafficking of P2X2 and P2X4 receptors cannot be explained by differences in their levels of expression.

**DISCUSSION**

These results demonstrate that two subtypes of purinoceptor, P2X2 and P2X4, are trafficked in distinct ways in cultured neurons. Whereas P2X2 receptors in the plasma membrane were...
relatively stable, P2X4 receptors were rapidly retrieved by a dynamin-dependent mechanism and subsequently cycled back to the plasma membrane. Binding of ATP to the P2X4 receptor stimulated its internalization, but ATP did not affect retrieval of P2X2 receptors. Within P2X4/6 heteromeric complexes, P2X4 played a dominant role in determining the subcellular distribution of the receptor. Our observation that P2X receptors can act presynaptically to enhance the release of glutamate suggests that the regulated cycling of P2X4-containing receptors might provide a mechanism for modulation of receptor density at presynaptic sites.

The P2X4 subunit determines the distribution of heteromeric receptors

The distribution of P2X4/6 heteromers resembled that of the P2X4 homomeric receptors, suggesting that the P2X4 subunit is dominant in determining the trafficking of the heteromer. Similarly, for AMPA receptors, there is diversity of trafficking that is subunit-dependent, and for heteromeric complexes it appears that one subunit will play a dominant role in determining the insertion and retrieval of receptors from the postsynaptic membrane (Sheng and Lee, 2001; Shi et al., 2001). There are several regions of the CNS where P2X2, P2X4, and P2X6 have been shown to be coexpressed, and therefore within neurons the endogenous P2X receptor population may consist of homomers and heteromers (Norenberg and Illes, 2000). However, even in regions where P2X4 mRNA is abundant and P2X4 protein expression was detected (e.g., in hippocampal neurons), neuronal P2X-mediated currents do not resemble homomorphic P2X4 receptor currents or even P2X4/6 heteromers with regards to their pharmacological properties (Khakh et al., 2001b; Rubio and Soto, 2001). One possible explanation is that there are no P2X4 homomers or P2X4/6 heteromers within the receptor population. However, our results offer a different possible explanation. The P2X4-containing receptors are predominantly in intracellular compartments and are upregulated at the cell surface only in response to certain physiological or pathological stimuli, whereas receptors with P2X2 subunits are more stable at the surface, and thus mediate the majority of the ATP-evoked response. Key questions to be addressed in future studies are how the cycling of P2X4 into and out of the plasma membrane is regulated and what are the important physiological regulators other than ATP?

Regulation of receptor trafficking by ATP

Application of ATP enhanced the rate of P2X4 receptor internalization. Despite the fact that P2X4 receptors are highly permeable to calcium (Soto et al., 1996), this effect of ATP occurred in the absence of extracellular calcium, suggesting that the increase might be caused by an agonist-induced conformational change in the receptor promoting an interaction with components of the endocytic machinery. However, the molecular interactions between P2X4 and endocytic proteins have yet to be characterized. Depending on the rate at which internalized receptors are subsequently reinserted into the plasma membrane, increased internalization might lead to a reduction in the size of the response to multiple applications of ATP. Ennion and Evans (2001) showed that for P2X1 receptors in smooth muscle, the ATP evoked increase in internalization was matched by an increase in the rate of reinsertion, and as a consequence the number of receptors at the surface remained fairly constant. They therefore concluded that after an application of ATP to P2X1 recep-
tors, the rate of recovery from desensitization was not limited by the rate at which receptors reinserted into the plasma membrane.

Application of ATP to neurons transfected with P2X2–GFP caused varicosities to appear along dendrites. Application of glutamate to cortical neurons leads to a similar change in the morphology, believed to be a pathological response caused by excessive cation influx (Faddis et al., 1997; Hasbani et al., 1998). It has been suggested that the “beading” of P2X2–GFP fluorescence in response to ATP might additionally reflect a redistribution of P2X2–GFP receptors within the membrane (Khakh et al., 2001a). A change in the surface distribution of receptors is difficult to quantify when the morphology of dendrites are changing, but the beading of P2X2–GFP fluorescence in response to ATP that we observed was exactly mimicked by changes in MAP-2 immunofluorescence (Fig. 5E). Thus, the simplest explanation for the change in the pattern of P2X2–GFP fluorescence is the obvious change in dendritic morphology rather than movements of receptors within the plasma membrane.

**Cycling of synaptic ionotropic receptors**

The cycling of receptors into and out of the plasma membrane provides a means of modulating receptor density by regulation of internalization or insertion rates (Burrone and Murthy, 2001). AMPA and GABA<sub>A</sub> receptors undergo both constitutive and regulated cycling, and this has been shown to be important in modulating excitatory and inhibitory transmission (Carroll et al., 2001; Moss and Smart, 2001). In contrast, NMDA receptors are more stable at the postsynaptic membrane, not because they lack a suitable endocytotic motif, but because they are stabilized via an interaction with PDZ domain proteins (Roche et al., 2001). The relative stability of P2X2 receptors might similarly reflect an interaction with some other protein or the lack of a determinant for rapid internalization present in P2X4. However, no suitable P2X2 receptor binding partners have been identified as yet.

The physiological role of P2X receptors in central and peripheral neurons is not well established. However, there are several examples of synapses where P2X receptors act presynaptically to increase neurotransmitter release. These include the first sensory synapse in the spinal cord (Gu and MacDermott, 1997; Nakatsuka and Gu, 2001), the synapse between brainstem and trigeminal neurons (Khakh and Henderson, 1998), and synapses in the nucleus tractus solitarii (Kato and Shigetomi, 2001). Our results suggest that heterologously expressed P2X receptors can also act presynaptically to regulate the release of glutamate. The regulation of P2X receptor density at presynaptic sites might therefore represent a mechanism for the modulation of synaptic transmission.

**REFERENCES**
