

P2Y receptors on astrocytes and microglia mediate opposite effects in astroglial proliferation

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Abstract Nucleotides released upon brain injury signal to astrocytes and microglia playing an important role in astrogliosis, but the participation of microglia in the purinergic modulation of astrogliosis is still unclear. Highly enriched astroglial cultures and co-cultures of astrocytes and microglia were used to investigate the influence of microglia in the modulation of astroglial proliferation mediated by nucleotides. In highly enriched astroglial cultures, adenosine-5'-triphosphate (ATP), adenosine 5'-O-(3-thio)-triphosphate (ATP γ S), adenosine 5'-O-(3-thio)-diphosphate (ADP β S; 0.01–1 mM), and adenosine-5'-diphosphate (ADP; 0.1–1 mM) increased proliferation up to 382%, an effect abolished in co-cultures containing 8% of microglia. The loss of ATP proliferative effect in co-cultures is supported by its fast metabolism and reduced ADP accumulation, an agonist of P2Y_{1,12} receptors that mediate astroglial proliferation. No differences in ADP β S and ATP γ S metabolism or P2Y_{1,12} receptors expression were found in co-cultures that could explain the loss of their proliferative effect. However, conditioned medium from microglia cultures or co-cultures treated with ADP β S, when tested in highly enriched astroglial cultures, also prevented ADP β S proliferative effect. None of the uracil nucleotides tested had any effect

in proliferation of highly enriched astroglial cultures, but uridine-5'-triphosphate (UTP; 0.1–1 mM) inhibited proliferation up to 66% in co-cultures, an effect that was dependent on uridine-5'-diphosphate (UDP) accumulation, coincident with a co-localization of P2Y₆ receptors in microglia and due to cell apoptosis. The results indicate that microglia control astroglial proliferation by preventing the proliferative response to adenine nucleotides and favouring an inhibitory effect of UTP/UDP. Several microglial P2Y receptors may be involved by inducing the release of messengers that restrain astrogliosis, a beneficial effect for neuronal repair mechanisms following brain injury.

Keywords Astroglial proliferation · P2Y receptors · Nucleotide metabolism · P2Y_{1,6,12} expression · P2Y_{1,6,12} cell-type localization · Astrocyte–microglia communication

Introduction

Astrogliosis and microglia activation are common features of neurodegenerative diseases and acute pathological episodes of trauma, stroke, seizure or infection [1, 2]. ATP and other nucleotides are massively released under these conditions and by activation of P2Y receptors initiate astrogliosis, a response that is characterised by an increase in glial fibrillary acidic protein (GFAP) expression, cell stellation and astroglial proliferation [3, 4]. P2Y receptors also mediate astrocyte migration [5] and modulate the release of cytokines [6, 7] and prostaglandins [8], supporting the astroglial reactive phenotype observed during astrogliosis. In injury models, astroglial proliferation was found to be mediated by P2Y₁ receptors [9, 10], and recently, we have shown that P2Y₁₂ receptors may also be involved in this response [11]. Additionally, nucleotides

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activate microglia P2 receptors which induce chemotaxis [12, 13], phagocytosis [14], the release of trophic factors [15] and cytokines [16] that may have protective effects in cerebral injury [17, 18]. Several cytokines and growth factors released by microglia, such as interleukin-1 β , interleukin-6, interferon- γ , tumour necrosis factor- α and fibroblast growth factor-2 stimulate astrogliosis [19–21] whereas others, such as interleukin-10, attenuate astroglial reactivity through a decrease in microglia activation [22, 23]. Furthermore, the *in vitro* and *in vivo* demonstration that microglia activation precedes astrogliosis lead to the proposal that these cells are of major relevance in the modulation of this response [24].

Most of the known effects of nucleotides in astrogliosis are based on results obtained from studies in astroglial cultures which, regardless of the protocols used, contained microglia in different proportions, but the influence of microglia in the purinergic trophic effects was rarely addressed [25]. Microglia, even when present in small amounts, may regulate astroglial responses and may be responsible for some of the effects attributed to astrocytes [26].

In this study, we investigated the influence of microglia in the modulation of astroglial proliferation mediated by nucleotides using two types of primary astroglial cultures: highly enriched astroglial cultures and co-cultures of astrocytes and microglia. In a first approach to understand the differences observed in the effects of nucleotides in both types of cultures, several factors that could offer an immediate explanation were investigated: (1) the metabolism of nucleotides, (2) the expression and cellular localization of the P2Y receptors potentially involved in the modulation of astroglial proliferation and (3) the release of soluble messengers by microglia that could have influenced astroglial proliferation. With this experimental approach, we aimed to start disclosing the purinergic mechanisms that influence the astrocyte–microglia communication during astrogliosis, a hallmark of brain injury.

Materials and methods

Drugs and antibodies

The following antibodies and drugs were used: goat anti-mouse IgG conjugated to Alexa Fluor 488 from Invitrogen (Barcelona, Spain); rabbit polyclonal anti-P2Y₁ and anti-P2Y₆ from Alomone Laboratories (Jerusalem, Israel); mouse monoclonal anti-CD11b, rabbit polyclonal anti-actin and goat anti-rabbit IgG conjugated to horseradish peroxidase from Santa Cruz Biotechnology (Santa Cruz, CA, USA); rabbit polyclonal anti-P2Y₁₂, rabbit and mouse anti-glia fibrillary acidic protein (anti-GFAP), goat anti-rabbit IgG

conjugated to crystalline tetramethylrodamine isothiocyanate (TRITC), adenosine, adenosine-5'-monophosphate (AMP), adenosine-5'-diphosphate tetrasodium (ADP), adenosine 5'-O-(3-thio)-diphosphate tetralithium (ADP β S), adenosine-5'-triphosphate disodium (ATP), adenosine 5'-O-(3-thio)-triphosphate tetralithium (ATP γ S), cytosine β -D-arabino-furanoside (Ara-C), 2'-(4-hydroxyphenyl)-5-(4-methyl-1-piperazinyl)-2,5'-bi-1 H-benzimidazole trihydrochloride hydrate (Hoechst 33258), hypoxanthine, inosine, L-leucine methyl ester hydrochloride (LME), penicillin, streptomycin, uracil, uridine, uridine-5'-monophosphate disodium (UMP), uridine-5'-diphosphate sodium (UDP), uridine-5'-triphosphate trisodium (UTP) and uridine 5'-diphosphoglucose disodium (UDP-glucose) and methyl green from Sigma-Aldrich (Sintra, Portugal); methyl-[³H]-thymidine (specific activity 80–86 Ci.mmol⁻¹) and enhanced chemiluminescence Western blotting system from Amersham Biosciences (Lisbon, Portugal); Sulfo-NHS-SS-Biotin and Ultralink Immobilized Neutravidin from Pierce (Rockford, IL, USA). Stock solutions of drugs were prepared with dimethylsulphoxide or distilled water and kept at -20°C. Solutions of drugs were prepared from stock solutions diluted in culture medium immediately before use.

Cell cultures

Animal handling and experiments were conducted according to the guidelines of the Directive 2010/63/EU of the European Parliament and the Council of the European Union. Primary cortical astroglial cultures were prepared from offspring of Wistar rats (Charles River, Barcelona, Spain) as previously described [27], with minor modifications. Briefly, the brains were placed in ice-cold Dulbecco's phosphate buffered calcium-free saline solution (DPBS) containing 0.2% glucose. The hemispheres were free of meninges and blood vessels, and after washing twice with ice-cold DPBS, they were cut into small pieces in culture medium, i.e., Dulbecco's modified Eagle medium containing 3.7 g/L NaHCO₃, 1.0 g/L D-glucose and stable glutamine, supplemented with 50 U/ml penicillin and 50 μ g/ml streptomycin. Tissue from two hemispheres was dissociated by triturating in 10 ml culture medium. The cell suspension obtained was passed through a 40- μ m pore nylon mesh and then centrifuged at 200 \times g for 5 min and the supernatant discharged. Centrifugation followed by cell suspension was repeated twice, and the pellet obtained was suspended in culture medium supplemented with 10% foetal bovine serum (FBS) and seeded at a density of 2 \times 10⁵ cells/ml. Cultures were incubated at 37°C in a humidified atmosphere of 95% air, 5% CO₂ and the medium was replaced 1 day after preparation and subsequently, twice a week.

Highly enriched astroglial cultures were obtained by treating confluent cultures, after 20 days *in vitro* (DIV),

with 8 μ M Ara-C for 4 days followed by treatment with 50 mM L-LME for 90 min [28]. At DIV28, two types of cultures were obtained: co-cultures of astrocytes and microglia, when no treatment was applied and highly enriched astroglial cultures, when cultures were treated with Ara-C plus LME. In both types of cultures, astrocytes were the main cell type, but the number of microglia present differed between the two types of cultures (see below). Cultures were synchronised to a quiescent phase of the cell cycle, by shifting serum concentration to 0.1% FBS for 48 h, being used in experiments at DIV30.

Cultures of microglia were obtained from confluent co-cultures that were shaken overnight at 200 rpm. Supernatants containing detached cells were centrifuged at $290\times g$ for 10 min. The pellet obtained was suspended in culture medium containing 10% FBS at a density of 3×10^4 cells/ml. Cells were seeded in 24-well plates, and the medium was changed 1 h later, allowing a selective attachment of microglia [29]. After cell synchronisation for 48 h, microglia cultures and co-cultures were treated with solvent or ADP β S (0.1 mM) for 8 h. After this period of incubation, the medium was discarded and replaced by fresh medium, which was collected 24 h later to be tested in highly enriched astroglial cultures. This medium was named microglia conditioned medium (MCM) or co-cultures conditioned medium (CCCM). Conditioned medium obtained from cells treated with solvent (MCM-S or CCCM-S) or with ADP β S (MCM-ADP β S or CCCM-ADP β S) was tested in proliferation assays of highly enriched astroglial cultures.

Immunocytochemistry

Cell cultures were fixed with a solution containing 4% formaldehyde and 4% sucrose in phosphate buffered saline (PBS; 100 mM NaH₂PO₄, 50 mM NaCl, pH adjusted to 7.3) and then treated with PBS containing 0.3% Triton X-100. For double-labelling astrocytes and microglia, cultures were incubated with the primary antibodies rabbit anti-GFAP (1:600) and mouse anti-CD11b (1:50), overnight at 4°C. For P2Y receptors localization, cultures were incubated with the primary antibodies mouse anti-GFAP (1:300) or mouse anti-CD11b (1:50) and rabbit anti-P2Y₁ (1:400), anti-P2Y₆ (1:200) or anti-P2Y₁₂ (1:400), overnight at 4°C. Visualisation of GFAP, CD11b and P2Y receptors positive cells was accomplished upon 1 h incubation, at room temperature, with the secondary antibodies anti-rabbit IgG conjugated to crystalline TRITC (1:100 and 1:400 for GFAP and P2Y receptors detection, respectively) and anti-mouse IgG conjugated to Alexa Fluor 488 (1:400). In negative controls, the primary antibody was omitted. Cell nuclei were labelled with Hoechst 33258 (5 μ g/ml) for 30 min at room temperature. To evaluate the percentage of microglia, the two types of

cultures were processed in parallel, and about 200 cells were counted in each culture. The number of CD11b-positive cells was expressed as percentage of the total number of cells counted. Images were captured with a Digital Sight DS-5Mc camera (Nikon, Japan) coupled to an Eclipse E400 fluorescence microscope (Nikon, Japan).

DNA synthesis

At DIV30, the cultures grown in 24-well plates were incubated with nucleotides or solvent for 48 h (tested in duplicate in each plate), and *methyl*-[³H]-thymidine was added in the last 24 h, at a concentration of 1 μ Ci/ml. When MCMs and CCCMs were tested in highly enriched astroglial cultures, they were added simultaneously with the nucleotides. Cells were then rinsed with PBS, fixed with 10% of trichloroacetic acid (TCA) for 30 min at 4°C, washed with ice-cold 5% TCA and rinsed again with PBS. Protein content and *methyl*-[³H]-thymidine incorporation were evaluated after cell lysis with 0.2 M NaOH. The effect of drugs in cell proliferation was determined by *methyl*-[³H]-thymidine incorporation which was quantified by liquid scintillation spectrometry (Beckman LS 6500, Beckman Instruments, Fullerton, USA) and normalised by the protein content determined by the Bradford method.

Metabolism of nucleotides

Cultures were rinsed three times with buffer at 37°C with the following composition (mM): 135 NaCl, 5 KCl, 0.8 MgSO₄, 1.8 CaCl₂, 10 HEPES, 10 glucose and 1 sodium pyruvate, with pH adjusted to 7.4 with NaOH (1 M). Nucleotides were added at zero time at concentration of 0.1 mM, and samples collected at 0, 1, 3, 8, 24 and 48 h were immediately stored at -20°C. Nucleotides and their metabolites were separated by ion-pair-reverse-phase high performance liquid chromatography with UV detection (HPLC-UV), as previously described [30]. Standards were analysed in the same conditions, and the retention time identified was (minutes): uracil (0.95), hypoxanthine (1.19), uridine (1.32), inosine (1.86), UMP (2.15), adenosine (3.93), UDP (4.40), AMP (4.76), UTP (6.40), ADP (6.63), ADP β S (7.70), ATP (7.87) and ATP γ S (8.10). The concentration of nucleotides and metabolites was calculated by peak area integration followed by interpolation in calibration curves obtained with standards.

Western blot analysis

Cells were rinsed with ice-cold PBS and total cell protein extracted in lysis buffer with protease inhibitors (1 mM Na₃VO₄, 1 mM NaF, 1 mM PMSF, 2 μ g/ml aprotinin and 2 μ g/ml leupeptin). After a brief sonication (10 s), the

lysate was incubated on ice for 1 h and then centrifuged at $20,000\times g$ for 45 min at 4°C . The protein concentration was determined in the supernatant, and equal amounts of protein (50 μg) were boiled at 95°C for 5 min in $6\times$ sample buffer (0.35 M Tris–HCl at pH 6.8, 10% sodium dodecyl sulfate (SDS), 30% glycerol, 9.3% dithiothreitol and 0.01% bromphenol blue) and subjected to 12% SDS-PAGE (SDS-polyacrylamide gel electrophoresis). Proteins were electrotransferred onto nitrocellulose membranes for 2 h at 40 V in a transfer buffer. Membranes were blocked overnight at 4°C with 5% of non-fat dry milk in PBS and then probed for 2 h at room temperature with appropriately diluted primary polyclonal antibodies: rabbit anti-P2Y₁, rabbit anti-P2Y₆ (both at 1:300) and rabbit anti-P2Y₁₂ (1:400) followed by secondary antibody goat anti-rabbit IgG conjugated to horseradish peroxidase (1:10,000). Immunoblots were then stripped by incubation in stripping buffer (62.5 mM Tris–HCl, 100 mM 2-mercaptoethanol and 2% SDS, pH adjusted to 6.8) for 15 min at 50°C and blocked overnight with 5% of non-fat dry milk in PBS. Subsequently, membranes were re-probed with the primary polyclonal antibody rabbit anti-actin (1:200) for 2 h at room temperature, followed by the secondary antibody. Immunocomplexes were detected by enhanced chemiluminescence system. Quantification of P2Y protein levels, obtained in arbitrary density units, was performed by densitometric analysis using Bio-Rad's Quantity One software (Basic version 4.6.5), and total P2Y receptors expression was normalised to actin.

Cell surface biotinylation

Cell surface protein biotinylation was performed to determine membrane expression of P2Y₁ and P2Y₆ receptors. Briefly, cultures were rinsed twice with ice-cold PBS with 0.1 mM CaCl₂ and 1.0 mM MgCl₂ (PBS–Ca–Mg). The apical surface was then exposed to 1 mg/ml of Sulfo-NHS-SS-biotin in biotinylation buffer (10 mM triethanolamine, 2 mM CaCl₂, 150 mM NaCl at pH 8.0) for 50 min with horizontal motion at 4°C . After labelling, the cells were rinsed with quenching solution (PBS–Ca–Mg with 100 mM glycine) and then extracted in lysis buffer with protease inhibitors. Precipitation of biotinylated proteins was attained by adding Neutravidin-agarose beads to approximately 850 μg of total cell protein, with end-over-end rotation overnight at 4°C . Then, beads were centrifuged three times at $6,000\times g$ for 2 min at 4°C , washed with PBS and bound proteins solubilised with SDS sample buffer (0.0625 M Tris–HCl at pH 6.8, 2% SDS, 10% glycerol, 2.5% 2-mercaptoethanol and 0.01% bromphenol blue). Samples were subjected to SDS-PAGE and blotting as described in the previous section (see [Western blot analysis](#)).

Lactate dehydrogenase assays

Necrotic cell death was assessed by measuring the lactate dehydrogenase (LDH) release with an enzymatic assay according to the manufacturer's instructions (Sigma Aldrich). The assay was based in the oxidation of lactate to pyruvate by LDH with the formation of NADH, which reduces tetrazolium to coloured formazan that was measured at a wavelength of 490 nm. Following incubation with nucleotides or solvent for 48 h, culture supernatants were collected, and the respective extracts were obtained upon incubation of astrocytes with a lysis solution for 45 min at 37°C . Samples were then centrifuged at $250\times g$ for 4 min, and LDH activity was determined in the collected supernatants and respective extracts. The LDH released into the culture medium was expressed as percentage of total LDH.

Terminal transferase-mediated dUTP nick end-labelling assays

Apoptotic cell death was evaluated using the indirect terminal transferase-mediated dUTP-digoxigenin nick end-labelling (TUNEL) to detect DNA fragmentation using an ApopTag peroxidase detection kit (Millipore, Madrid, Spain). Cultures treated with nucleotides or solvent for 48 h were fixed in 4% paraformaldehyde in PBS pH 7.4, for 10 min at room temperature and subsequently post-fixed in pre-cooled ethanol/acetic acid (2:1, v/v) for 15 min at -20°C . The endogenous peroxidase activity was quenched with 3% hydrogen peroxide. Cultures were incubated with equilibration buffer and treated with terminal deoxynucleotidyltransferase plus digoxigenin-dNTPs for 1 h at 37°C . The anti-digoxigenin antibody conjugated to peroxidase was added for 30 min at room temperature, and colour was developed with 3,3'-diaminobenzidine substrate. Counterstaining of the nuclei was accomplished with 0.5% methyl green, and cells were visualised by bright field microscopy. The cell bodies were also labelled with Hoechst 33258 (5 $\mu\text{g}/\text{ml}$) to confirm the results obtained with the TUNEL assay. The number of TUNEL-positive cells was evaluated by analysing eight high-power fields ($\times 400$) in each culture and expressed as percentage of total cell number counted.

Statistical analysis

Data are expressed as means \pm standard errors of the mean (SEM) from n number of experiments, unless otherwise stated. Statistical analysis was carried out using the unpaired Student's t test or one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison test. The Western blot data analysis was performed by ANOVA repeated measures followed by Bonferroni's multiple

comparison test. *P* values lower than 0.05 were considered to indicate significant differences.

Results

Characterization of cell cultures

Cultures that grown without any treatment were named co-cultures and could be described as monolayers of astrocytes exhibiting a flattened, polygonal morphology, containing $8.0\pm 0.8\%$ ($n=5$) of microglia (Fig. 1a). Highly enriched astroglial cultures were obtained by applying a very effective treatment to eliminate microglia from astroglial cultures [28], which resulted in highly enriched astrocytes

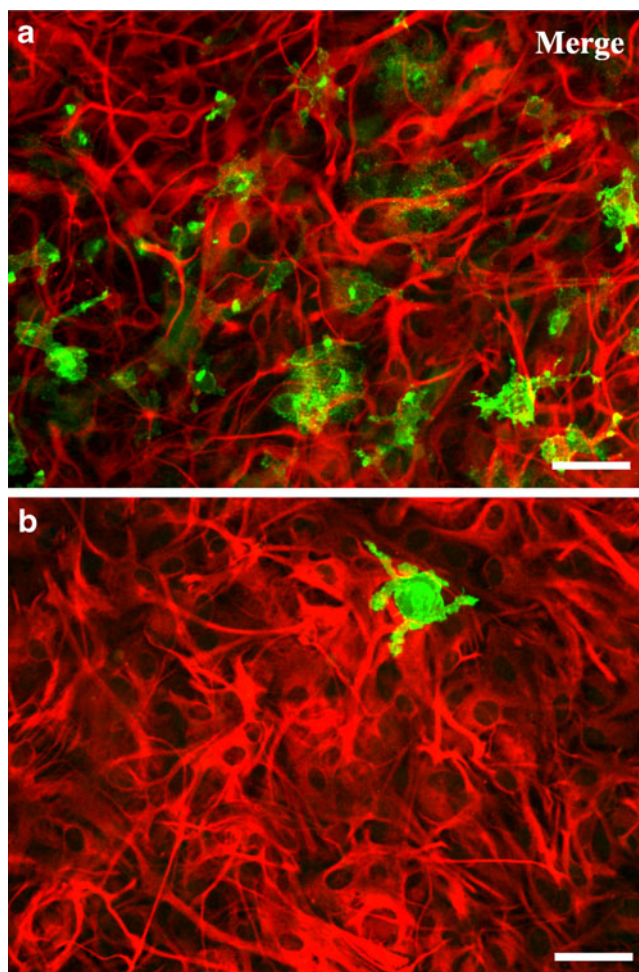


Fig. 1 Characterization of primary astroglial cultures containing different percentage of microglia. Astrocytes were labelled with anti-GFAP (TRITC, red) and microglia with anti-CD11b (Alexa Fluor 488, green). Representative immunofluorescent micrographs of the two types of cultures: **a** co-cultures and **b** highly enriched astroglial cultures, double-labelled for GFAP and CD11b. In co-cultures, the number of microglia present was $8.0\pm 0.8\%$ ($n=5$) and in highly enriched astroglial cultures was $1.0\pm 0.3\%$ ($n=5$). Scale bar: 50 μm

cultures containing only $1.0\pm 0.3\%$ ($n=5$) of microglia, which was considered negligible (Fig. 1b). The two types of primary cultures, highly enriched astroglial cultures and co-cultures were used in the experiments.

Effect of nucleotides in cell proliferation

Nucleotides that activate different P2Y receptor subtypes were tested in the cultures previously described: ATP and ATP γ S which are non-selective P2 agonists, ADP β S and ADP that have higher affinity for the P2Y_{1,12,13} subtypes, UTP that activates the P2Y_{2,4} subtypes, UDP that is selective for the P2Y₆ receptors and UDP-glucose, the selective agonist of the P2Y₁₄ receptors [31]. The nucleotides tested mediated opposite effects in cell proliferation, and their effects were influenced by the presence of microglia (Fig. 2).

In highly enriched astroglial cultures, the adenine nucleotides ADP (0.1–1 mM), ATP γ S and ADP β S (0.01–1 mM) increased cell proliferation in a concentration-dependent manner up to $382\pm 33\%$ (Fig. 2a). ATP increased cell proliferation up to a concentration of 0.1 mM and then the effect declined (Fig. 2a). In co-cultures, neither ATP nor ATP γ S or ADP β S had any effect in cell proliferation (Fig. 2b). ADP was not tested in co-cultures, since ADP β S, which is a more stable analogue [32], caused no effect. The absence of proliferative effects of adenine nucleotides in co-cultures was not due to the inability of cells to proliferate, since 10% FBS increased cell proliferation by $398\pm 54\%$ ($n=5$, $P<0.05$). The uracil nucleotides UDP and UDP-glucose (0.001–1 mM) had no effect in both types of cultures (Fig. 2c, d), but UTP (0.1–1 mM) inhibited cell proliferation up to $66\pm 6\%$ ($n=4$, $P<0.05$) in co-cultures (Fig. 2d). The results indicate that the presence of microglia in the cultures prevents the astroglial proliferative response to the endogenous nucleotide ATP and to the stable adenine nucleotide analogues ATP γ S and ADP β S [32] and favours the inhibition of proliferation caused by UTP.

Extracellular metabolism of nucleotides

In highly enriched astroglial cultures, ATP was rapidly metabolised into ADP, which was the main metabolite formed in the first hour of incubation and remained detectable in the following 7 h (Fig. 3a), whereas in co-cultures ATP was directly converted into AMP in the first hour, and ADP was hardly detected (Fig. 3b). ATP γ S was more stable than ATP, being slowly metabolised in both types of cultures. The half-life for ATP γ S was of 8.5 ± 0.3 h ($n=4$) in highly enriched astroglial cultures and 7.9 ± 0.4 h ($n=4$) in co-cultures. ADP β S was also metabolically stable in the first 8 h; then its concentration slowly declined without being completely metabolised after 48 h of incubation (Fig. 3c, d). Results indicate that in highly

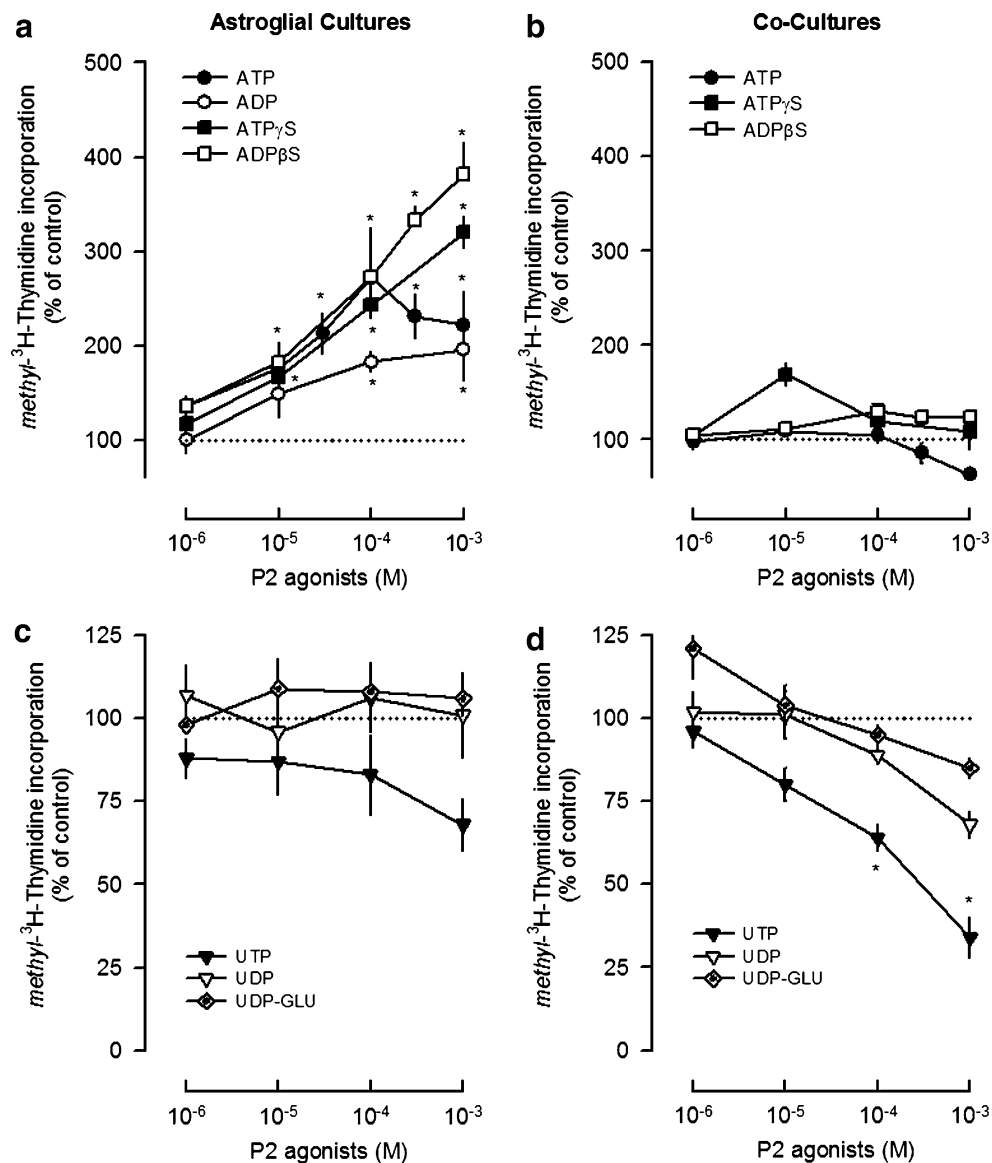


Fig. 2 Modulation of astroglial proliferation by nucleotides in highly enriched astroglial cultures (a, c) and co-cultures (b, d). Cultures were incubated with nucleotides or solvent for 48 h and *methyl*-³H-thymidine (1 μ Ci/ml) was added in the last 24 h. Cell proliferation

was estimated by *methyl*-³H-thymidine incorporation and expressed in percentage of control. Values are means \pm SEM from five to seven experiments. * P <0.05, significant differences from control (solvent)

enriched astroglial cultures the effect of ATP was predominantly mediated by ADP, whereas, in co-cultures, its effect in cell proliferation was lost (Fig. 2b) because the metabolism of ADP was faster. ADP β S and ATP γ S metabolism was much slower and similar in both types of cultures, indicating that other mechanisms need to be addressed to explain their loss of proliferative effect in co-cultures (Fig. 2b). In both types of cultures, UTP was rapidly metabolised into UDP, which attained higher concentrations in highly enriched astroglial cultures (Fig. 3e) than in co-cultures (Fig. 3f). Even though an inhibitory effect mediated by UTP was only observed in co-cultures (Fig. 2d). Therefore, besides the differences found

in the metabolism, other mechanisms come into play to explain the inhibitory effect of UTP in co-cultures.

Expression and cellular localization of P2Y₁, P2Y₁₂ and P2Y₆ receptors

The loss of ADP β S proliferative effect in co-cultures could not be explained by differences in its metabolic profile. Therefore, based on previous studies indicating that ADP β S mediates astroglial proliferation by activation of P2Y₁ and P2Y₁₂ receptors [11], the possibility that differences in the expression of these P2Y receptor subtypes could explain the attenuation of ADP β S proliferative effect

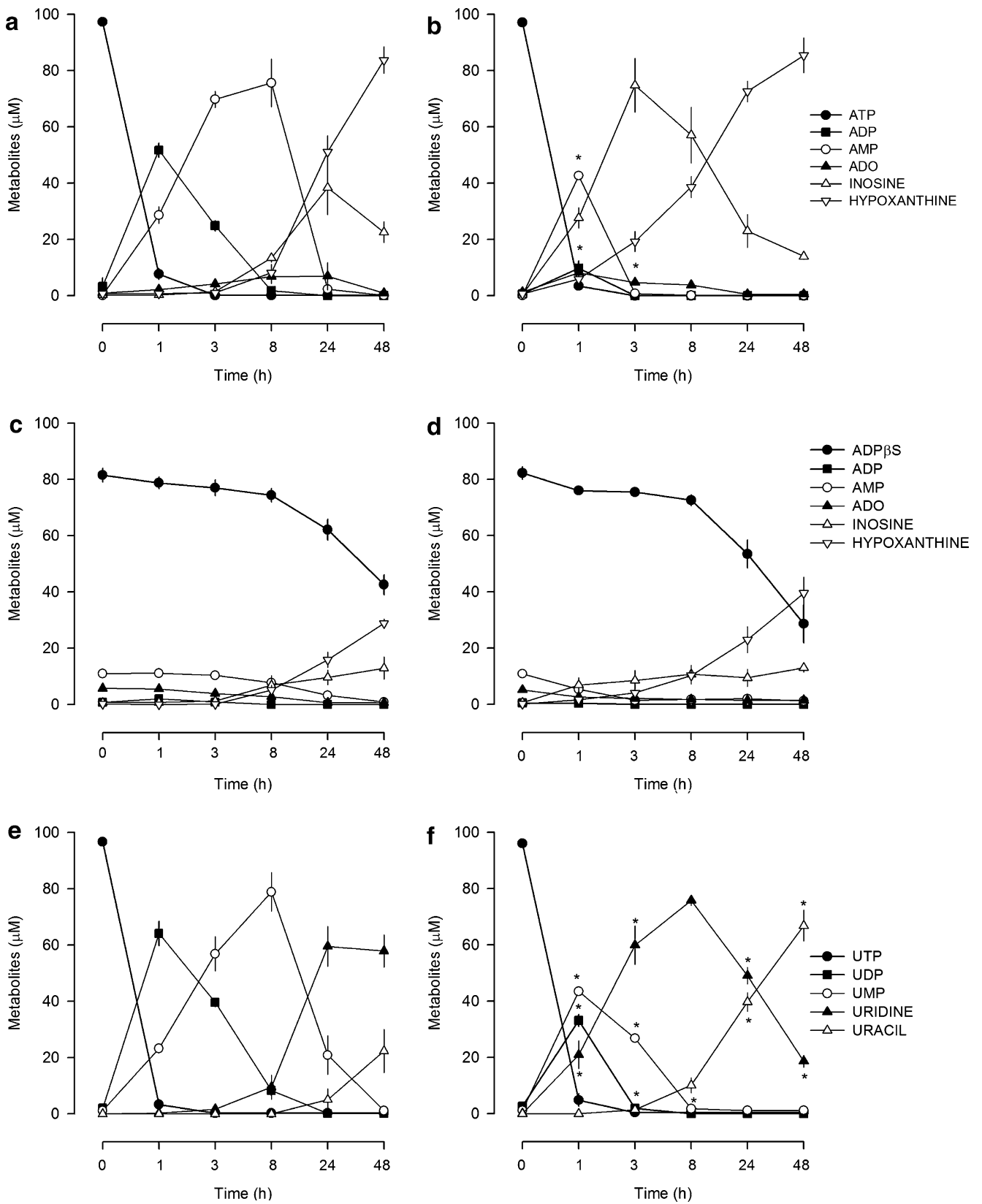


Fig. 3 Metabolism of nucleotides in highly enriched astroglial cultures (a, c, e) and co-cultures (b, d, f). Cultures were incubated with 0.1 mM of ATP, ADPβS or UTP, and samples were collected at 0, 1, 3, 8, 24 and 48 h. Nucleotides and their metabolites were

quantified by HPLC-UV as described in [Materials and methods](#). Values are means±SEM from four experiments. **P*<0.05, significant differences from highly enriched astroglial cultures

observed in co-cultures was investigated. Additionally, the rapid conversion of UTP into UDP, which is selective for P2Y₆ receptors [33], also raised the question of whether P2Y₆ receptor expression could be increased in cultures with microglia, thus favouring the inhibitory effect mediated by UTP/UDP in co-cultures.

These two hypotheses were tested by evaluating the relative expression of P2Y₁, P2Y₁₂ and P2Y₆ receptor subtypes in both types of cultures. The P2Y₁ receptor expression was represented by three immunoreactive bands of 42, 90 and 266 kDa that reacted with the anti-P2Y₁ antibody. These bands were absent in the presence of the P2Y₁ neutralising peptide indicating the P2Y₁ receptor

specificity (Fig. 4). Total P2Y₁ receptor expression (tP2Y₁) and the membrane fraction (mP2Y₁) were similar in the two types of cultures (Table 1). Total P2Y₁₂ receptor (tP2Y₁₂) expression was represented by a single band of 65 kDa (Fig. 4), with a similar expression in both types of cultures (Table 1), but, because the expression was low, the respective membrane fraction was not evaluated.

The P2Y₆ receptor comprised four bands, two of approximately 25, one of 36 and other of 86 kDa, which were all absent in the presence of the P2Y₆ neutralising peptide (Fig. 4), indicating they all represent the P2Y₆ receptor. When the four protein bands were quantified, total P2Y₆ receptor expression (tP2Y₆) was similar in both types of cultures

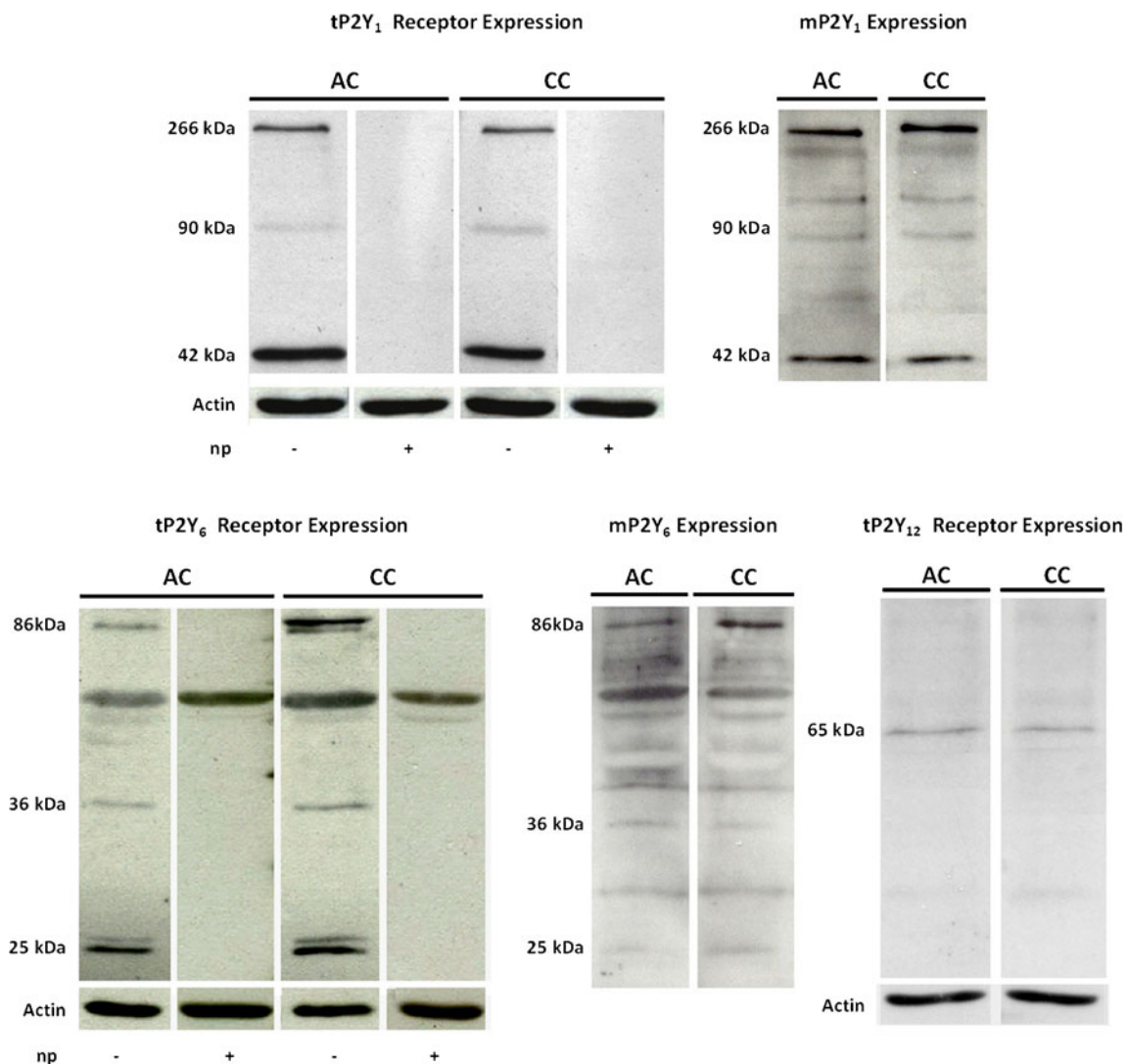


Fig. 4 Representative Western blots showing P2Y₁, P2Y₆ and P2Y₁₂ receptors expression in highly enriched astroglial cultures (AC) and co-cultures (CC). Total P2Y₁ (tP2Y₁), P2Y₆ (tP2Y₆) and P2Y₁₂ (tP2Y₁₂) receptors and actin (43 kDa) expression were obtained from whole cell lysates. Expression of membrane P2Y₁ (mP2Y₁) and P2Y₆ (mP2Y₆) receptors was obtained from surface biotinylated and

immunoprecipitated proteins. Three immunoreactive bands of 42, 90 and 266 kDa specifically reacted with rabbit anti-P2Y₁ antibody, and those of 25, 36 and 86 kDa specifically reacted with rabbit anti-P2Y₆ antibody. These bands were absent in the presence of the respective neutralising peptides. A single band of ~65 kDa was detected with the rabbit anti-P2Y₁₂ antibody

Table 1 Relative expression of P2Y₁, P2Y₆ and P2Y₁₂ receptors in astroglial cultures

% Highly enriched astroglial cultures							
	tP2Y ₁	mP2Y ₁	tP2Y ₆	tP2Y ₆ 86 kDa	mP2Y ₆	mP2Y ₆ 86 kDa	tP2Y ₁₂
Co-cultures	93±14 (5)	121±22 (3)	129±19 (4)	245±20* (4)	176±12* (3)	200±57* (3)	93±11 (3)

Quantification by densitometry of P2Y₁, P2Y₆ and P2Y₁₂ receptors total expression (tP2Y₁, tP2Y₆, tP2Y₁₂) and the membrane fractions of P2Y₁ and P2Y₆ receptors (mP2Y₁, mP2Y₆). The expression of the 86 kDa band of P2Y₆ receptors was also compared between cultures. Values are means±SEM from (*n*) cultures represented in percentage of the expression observed in highly enriched astroglial cultures. Statistical analysis was performed using one-way repeated measures analysis of variance followed by Bonferroni multiple comparison test

**P*<0.05, significant differences from highly enriched astroglial cultures

(Table 1). However, the analysis of individual bands revealed that expression of the 86 kDa band was higher in co-cultures than in highly enriched astroglial cultures (Table 1). Moreover, when the membrane fraction (mP2Y₆) was quantified, a higher expression was detected in co-cultures, considering either the four bands or only that of 86 kDa (Table 1).

The immunocytochemical studies designed to identify a preferential cellular localization of these receptors in co-cultures revealed that P2Y₁ and P2Y₁₂ receptors were localised in both astrocytes and microglia, without any predominant cell-type localization (Fig. 5). However, the pattern of P2Y₆ receptors distribution was different, revealing a higher co-localization with microglia (Fig. 5),

which is in agreement with the higher mP2Y₆ receptor expression observed in co-cultures.

Microglia soluble factors and cell death

Attenuation of the proliferative response to adenine nucleotides in co-cultures could not be explained by differences in the P2Y₁ and P2Y₁₂ receptors expression or by a preferential cellular localization of these receptors in astrocytes or microglia. However, P2Y receptors localised in microglia may still play a role in these effects. Since ADPβS activates P2Y_{1,12,13} receptors subtypes, which are expressed by microglia [34, 35], it was investigated if ADPβS mediates

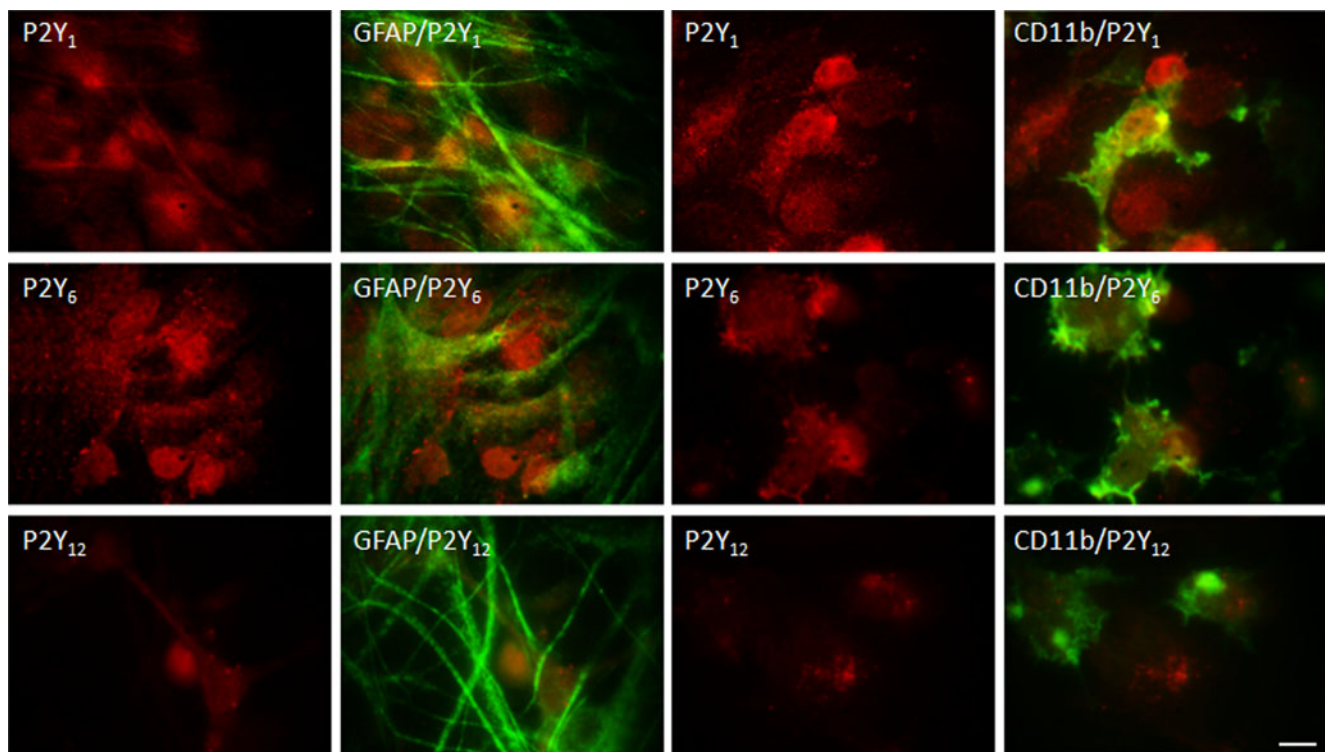


Fig. 5 Representative immunofluorescent micrographs of co-cultures double-labelled for P2Y receptors and GFAP or CD11b. P2Y receptors were labelled with anti-P2Y₁, anti-P2Y₆ and anti-P2Y₁₂ (TRITC, red), astrocytes and microglia with anti-GFAP and with anti-

CD11b (Alexa Fluor 488, green), respectively. P2Y₁, P2Y₆ and P2Y₁₂ receptors are expressed both by astrocytes (first and second panels) and microglia (third and fourth panels). A stronger immunoreactivity for P2Y₆ receptors co-localised with microglia. Scale bar=10 μm

the release of messengers from microglia that might interfere with its proliferative effect. This hypothesis was tested by comparing the effect of ADP β S (0.1 mM) in highly enriched astroglial cultures, in the absence and in the presence of conditioned medium obtained from microglia cultures or co-cultures that were previously treated with solvent (MCM-S or CCCM-S, respectively) or with ADP β S (MCM-ADP β S or CCCM-ADP β S, respectively).

The proliferative effect of ADP β S (0.1 mM) was not modified by MCM-S but was abolished in the presence of MCM-ADP β S (Table 2). Furthermore, in the presence of CCCM-S the proliferative effect of ADP β S (0.1 mM) was attenuated, an effect even more evident when the cultures were treated with CCCM-ADP β S (Table 2). In none of the conditions tested was astroglial proliferation near its maximum, as previously demonstrated the proliferative effect mediated by ADP β S (0.3–1 mM) in highly enriched astroglial cultures (see Fig. 2a). These results suggest that ADP β S activates microglia P2Y_{1,12} and/or P2Y₁₃ receptors, which induce the release of diffusible messengers that attenuate its proliferative effect in astrocytes. They further suggest that the co-presence of astrocytes and microglia, and eventually their physical contact, facilitates the release of these inhibitory messengers since CCCM-S, but not MCM-S, attenuated ADP β S proliferative effect (Table 2). The same experimental approach was adopted for UTP, but no effects could be detected (not shown), possibly because of its higher metabolic instability, but, since UTP caused an inhibition of proliferation in co-cultures, its contribution to cell death was investigated. UTP (1 mM) caused no change in LDH release (not shown), but it increased the number of cells presenting DNA fragmentation, detected by the TUNEL assay, from 1.25 \pm 0.13% ($n=3$) in solvent treated cultures to 6.21 \pm 0.72% ($n=3$, $P<0.05$). Results suggest that, in co-cultures, the presence of microglia may favour UTP/UDP-induced cell apoptosis, contributing to the inhibition of cell proliferation.

Discussion

Astrogliosis and microglia activation are two closely related responses involved in the brain repairing mechanisms to injury, and research in the field has been focusing the attention on how astrocytes and microglia influence each other activity during this process [24]. Astroglial proliferation induced by nucleotides was previously demonstrated both *in vitro* [3, 4] and *in vivo* [9]. However, whereas the *in vivo* studies did not discriminate the influence of microglia, the *in vitro* studies were performed in astroglial cultures, with the special concern of working without microglia interference. When the influence of microglia was to be explored, the mostly used strategy consisted in the evaluation of microglia conditioned medium influence in astroglial cultures [36, 37]. Although being an interesting and valid approach, it only evaluates astroglial response to soluble factors released by microglia, considering a unidirectional communication between both types of cells, which excludes the possibility that astrocytes may also release soluble factors that regulate and/or influence microglia responses [38, 39]. In the present study, the influence of microglia in the astroglial proliferative response to nucleotides was investigated by comparing their effects in two types of cultures: highly enriched astroglial cultures with a negligible presence of microglia and co-cultures of astrocytes containing 8% of microglia. This approach has the advantage of preserving the physical contact between astrocytes and microglia, facilitating the action of soluble messengers and a bidirectional communication between both cell types.

As expected, in highly enriched astroglial cultures, the adenine nucleotides increased astroglial proliferation in a concentration-dependent manner [3, 4]. The concentration–response curve to ATP was biphasic; it caused an increase in astroglial proliferation, mainly by activation of P2Y₁, A_{2A} and A_{2B} receptors [11], but at the highest concentrations tested (0.3–1 mM), the proliferative response

Table 2 Influence of conditioned medium from microglia cultures or co-cultures in the ADP β S-induced astroglial proliferation in highly enriched astroglial cultures

Methyl-[³ H]-thymidine incorporation (% of control)					
Medium/drugs	No treatment	MCM-S	MCM-ADP β S	CCCM-S	CCCM-ADP β S
ADP β S 0.1 mM	222 \pm 10* (16)	193 \pm 17* (6)	114 \pm 18*** (6)	157 \pm 7* (8)	128 \pm 5* (8)

Conditioned medium from microglia cultures or co-cultures treated with solvent (MCM-S and CCCM-S, respectively) or with 0.1 mM ADP β S (MCM-ADP β S and CCCM-ADP β S, respectively) was tested in highly enriched astroglial cultures in combination with 0.1 mM ADP β S for 48 h and methyl-[³H]-thymidine (1 μ Ci/ml) was added in the last 24 h. The effects in cell proliferation were estimated by methyl-[³H]-thymidine incorporation and expressed in percentage of respective control (solvent, MCM-S or CCCM-S). Values are means \pm SEM from (n) experiments

* $P<0.05$, significant differences from respective control

** $P<0.05$, from the effect of MCM-S or CCCM-S

*** $P<0.05$, from the effect of ADP β S alone

declined, which may be explained by activation of inhibitory adenosine A₁ receptors [3] or by activation of P2X₇ receptors that have opposite effects to those of P2Y receptors in cell proliferation [40]. Several uracil nucleotides were also tested but did not change astroglial proliferation. Even the P2Y₁₄ receptors that are highly expressed in glial cells [41] were not involved in the modulation of astroglial proliferation since the selective agonist at these receptors, UDP-glucose, had no effect. Previous studies have shown that UTP may contribute to astroglial proliferation but only in synergism with the fibroblast growth factor-2 [40].

In co-cultures of astrocytes and microglia, the proliferative effect of adenine nucleotides was abolished, whereas UTP or its metabolite UDP inhibited astroglial proliferation.

This study demonstrates that microglia modify the astroglial response to nucleotides. In order to explain the differences observed in the proliferative responses to nucleotides in the two types of cultures, several hypotheses were considered: (1) the nucleotides metabolic profile was different in the presence of microglia, (2) the P2Y receptors expression could be different in the presence of microglia, (3) P2Y receptors could have a preferential cellular localization or (4) microglia P2Y receptors induced the release of messengers that modified the astroglial response to nucleotides and/or reduced cell viability.

The time course of ATP and UTP metabolism was similar in both types of cultures, but the accumulation of ADP and UDP differed. The presence of microglia in cultures accelerated the metabolism of the intermediates, having a higher influence on ADP than on UDP metabolism. Therefore, in co-cultures, ADP was hardly detected, which can explain the loss of ATP proliferative effect, whereas in highly enriched astroglial cultures the accumulation of ADP favoured the activation of P2Y₁ receptors, which mediate astroglial proliferation [11]. However, the differences observed in the metabolism of UTP are insufficient to explain why it only caused inhibition of proliferation in co-cultures. Additionally, the metabolic profile of the stable nucleotides ADPβS and ATPγS failed to explain the loss of their proliferative effects in co-cultures.

The nucleotide ADPβS induced astroglial proliferation through the activation of P2Y₁ and P2Y₁₂ receptors, and this effect was lost in co-cultures, therefore the relative expression of these receptors in both types of cultures and their cell-type localization were evaluated. P2Y₁ receptor expression presented a multiple band pattern: a band of 42 kDa that corresponds to the P2Y₁ receptor monomer [42] and additional bands with molecular weights of 90 and 266 kDa. Similar bands were previously described and may represent homomeric forms of the P2Y₁ receptor, or heteromers with A₁ receptors [43–45]. P2Y₁ and P2Y₁₂ receptors expression was similar in both types of cultures

(Table 1), as well as the pattern of distribution in astrocytes (not shown), arguing against these two factors as relevant contributors to the loss of proliferative effects mediated by the stable nucleotides in co-cultures. The most feasible explanation for the differences in proliferation found in both types of cultures resides in the influence mediated by the microglia P2Y receptors. Our findings show that microglia exert an inhibitory influence in the P2Y receptor-mediated proliferative effects in astrocytes. Furthermore, this inhibitory influence of microglia is more pronounced when in contact with astrocytes, since CCCM-S, but not MCM-S, attenuated the proliferative effect of ADPβS in highly enriched astroglial cultures. In agreement with this observation, previous studies have demonstrated that in co-cultures the spontaneous release of ATP by astrocytes may enhance the release of messengers by microglia, which may be involved in the regulation/control of astroglial proliferation [38]. The CCCM-ADPβS and MCM-ADPβS prevented ADPβS-induced astroglial proliferation in highly enriched astroglial cultures. Therefore, besides the inhibitory influence provided by the presence of microglia, activation of microglia P2Y receptors sensitive to ADPβS seem to contribute to the release of messengers that interfere with the proliferative effects mediated by astroglial P2Y receptors. Microglia express P2Y₁, P2Y₁₂ and P2Y₁₃ receptors [34, 35], which are activated by ADPβS and may regulate the release of messengers, such as interleukin-10 [46]. These messengers may regulate astroglial proliferation mediated by P2Y receptors through a functional interaction occurring at the receptor level and/or at the intracellular signal transduction pathways. For example, interleukin-1β was shown to decrease the activity of P2Y₁ receptors by a mechanism that involves an interaction with connexin 43 [47, 48], but interleukin-1β and tumour necrosis factor-α may also activate intracellular signalling pathways shared by P2Y receptors [49] and consequently regulate astroglial response. Other messengers such as transforming growth factor-β or interleukin-4 mediate opposite effects to adenine nucleotides [50, 51] and may prevent astroglial proliferation. Therefore, several candidates exist to mediate this interaction between astrocytes and microglia in the modulation of astroglial proliferation and it is likely that several mediators participate in this process.

A comparative study of the P2Y₆ receptors expression and their cell-type localization helped to have clearer picture of the differences found in the UTP effect between the two types of cultures. P2Y₆ receptor expression also revealed a multiple band pattern; besides the band of 36 kDa predicted for this receptor [52], additional bands of lower molecular weight were also detected, which may correspond to degradation products, and a band of 86 kDa that was previously reported to represent a homomeric association of P2Y₆ receptors or a heterodimeric association between P2Y₆ and P2Y₄ receptors [53]. The tP2Y₆

expression was similar in both types of cultures, but the 86 kDa band and the mP2Y₆ had a higher expression in co-cultures. Additionally, P2Y₆ receptors were highly localised in microglia with a more diffuse and less intense distribution in the astrocyte net. These results indicate that UTP metabolism with UDP formation and a higher expression of P2Y₆ receptors by microglia may have favoured activation of microglia P2Y₆ receptors, mediating UTP/UDP-induced apoptosis through the release of inhibitory messengers. These messengers may include several cytokines described to inhibit proliferation (see above); a short-lived messenger such as nitric oxide [54] or other messengers not yet identified that may be released by microglia P2Y₆ receptor-stimulation.

Our results indicate that microglia present in these co-cultures are sufficient to influence the effects of modulators of astroglial proliferation and underline the importance of studying the contribution of microglia P2Y receptors to the modulation of astroglial proliferation induced by nucleotides. This modulation is mediated through the release of messengers, not yet identified, but whose identity is currently under investigation. Activation of P2Y₁, P2Y₁₂ and/or P2Y₁₃ receptors attenuates the proliferative effect of adenine nucleotides, and activation of P2Y₆ receptors mediates cell apoptosis triggered by uracil nucleotides. In addition to the roles previously described for P2Y₆ receptors, i.e. secretion of cytokines and phagocytosis [55], these receptors also modulate astroglial proliferation, a mechanism that is important to prevent excessive astroglial proliferation that may compromise neuronal repair.

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