Discussion

In the present study, we first identified three motifs that are canonical consensus sequences of RAREs in the cloned 5′-flanking region of the Wistar rat *P2rx2* and found that 9-*cis*-RA, an endogenous vitamin A derivative, increases the expression of the $P2X_2$ receptor at the transcriptional level in the neuronal model PC-12 cells. The transcriptional effects of 9-*cis*-RA are primarily mediated by activating two families of nuclear receptors, RARs and RXRs (Chambon, 1996). RXRs can form as homodimers and as heterodimers with a number of other nuclear receptors such as thyroid hormone receptor, vitamin D receptor, and RAR (Aranda and Pascual, 2001). Among them, the RXR/RAR heterodimer is known to respond specifically to the RAR activator atRA (Kurokawa et al., 1994). The present study did not show direct binding of RAR and RXR with 5′-flanking region of *P2rx2* but did demonstrate that atRA-treated PC-12 cells also show an increase in the level of $P2X_2$ mRNA expression, suggesting the involvement of RAR in regulating the $P2X_2$ receptor expression in PC-12 cells. PA024 did not increase the level of $P2X_2$ mRNA. A slight, but not significant, increase in $P2X_2$ mRNA was seen. This finding corresponds with the fact that a low activity of PA024 alone was observed in an experiment of retinoid-induced HL-60 differentiation (Ishida et al., 2003). That PA024 scarcely increased the $P2X_2$ mRNA expression is consistent with the findings of previous studies showing that a single application of RXR-selective agonists does not induce gene transcription (Minucci et al., 1997; Ishida et al., 2003) and is supported by the notion that the RXR ligand induces homodimerization of RXR and inhibits heterodimerization without dimerization partner ligands; moreover, a partner ligand is sufficient for heterodimerization (Dong and Noy, 1998). The RXR/RAR heterodimer generally binds to the DR5 RARE (Kurokawa et al., 1994) and also binds to DR1 (Kurokawa et al., 1994). We determined the $P2X_2$ mRNA transcription start site by 5′ RACE, which is located near the site supposed by RefSeq entry (NM_053656). We also had the predictions for some transcription factor binding sites in the 5′-flanking region of the rat *P2rx2* cloned in the current study, which include the DR5 and DR1 sequences located at −2381/−2397 and −2292/−2294 from the transcription start site. It includes other factors such as simian virus 40 protein 1, activator protein-2, nuclear factor-κB, GATA-1, cAMP response element binding protein, GC-box, and initiator sequence as well. Consensus sequences for GC-box and initiator found in our cloned sequence imply that core promoter region would exist near the 5′ end of our cloned sequence. Although the factors we showed here were just the candidates estimated by the electrical search system, we confirmed that the cloned fragment has sensitivity to retinoid treatment and deletion of a fragment containing DR elements lead to abolishing the 9-*cis*-RA- and atRA-mediated and parts of basal transcriptional activities. On the other hand, the deleted fragment also contains DR4, but this is known as a binding site of RXR heterodimerized with nuclear receptors other than RARs (Aranda and Pascual, 2001). Because RAR/RAR homodimerization has not been reported, our series of results could suggest that retinoic acids activate RAR/RXR heterodimers that bind to RAREs (DR5 and/or to DR1-responsive elements) located at the distant place from transcription start site in the promoter region of the *P2rx2*, which in turn work as activators of basal transcription machinery and lead to an increase in the transcription of $P2X_2$ receptors in PC-12 cells.

The biochemical analysis in the present study indicated that the increase in $P2X_2$ transcription resulted in an increase in the level of $P2X_2$ protein. Furthermore, we found that the maximal responses of ATP-evoked currents were enhanced in 9-*cis*-RA-treated PC-12 cells. The inward currents evoked by ATP in PC-12 cells have been demonstrated to be inhibited by suramin, PPADS, and reactive blue 2 (Inoue et al., 1991a,b), a pharmacological profile that fits rat $P2X_2$ receptors, thus suggesting an increase in the level of functional $P2X_2$ protein. This view is strongly supported by the finding that the Ca^{2+} response evoked by ATP in 9-*cis*-RA-treated PC-12 cells was enhanced in the presence of a phospholipase Cβ inhibitor, which abolishes P2Y-mediated Ca^{2+} responses. It could be possible that ATP produces an inward current via activating another P2X subtype. Indeed, in addition to $P2X_2$ receptors $P2X_4$ transcript was also detected in PC-12 cells by our RT-PCR analysis (our unpublished observation). However, 20 μM PPADS almost completely blocked ATP-induced inward currents and $[Ca^{2+}]_i$ elevation, and 9-*cis*-RA did not alter the EC_{50} and Hill coefficient value of the ATP-evoked currents in the PC-12 cells. It is suggested that functional $P2X_4$ receptor is not expressed on the cell membrane. In addition, the mRNA level of the $P2X_4$ receptor in the PC-12 cells was not changed by treatment with 9-*cis*-RA (our unpublished observation). In human cervical epithelial cells, however, the expression of $P2X_4$ mRNA has been reported to be increased by atRA (Gorodeski, 2002). This discrepancy may be due to differences in the species, the basal expression levels of $P2X_4$ receptors, and the expression of RAR and RXR isoforms or the large numbers of coregulators.

In the nervous system, a key function of $P2X_2$ receptors is to increase release of neurotransmitters (Khakh et al., 2003). PC-12 cells are frequently used in studies investigating stimulus-induced vesicular transmitter release (Shafer and Atchison, 1991). We have observed that retinoid significantly enhanced the ATP-evoked release of DA from PC-12 cells. Because retinoid treatment might lead to the changes in many gene transcriptions involved in $[Ca^{2+}]_i$ elevation, exocytotic machinery, or packaging in vesicles, the enhancement of DA release seen in the present study might include multiple interpretations. However, we found that enhancement by 9-*cis*-RA of the $P2X_2$ receptor protein expression level and ATP-activated Ca^{2+} entry was almost identical to that of the ATP-evoked DA release. In addition, 9-*cis*-RA did not affect basal release or the total content of DA in PC-12 cells, suggesting the 9-*cis*-RA affects neither DA biosynthesis nor exocytotic machinery itself. Calcium is one of the most important factors to regulate exocytosis, and we previously showed that the ATP-evoked DA release from PC-12 cells is induced by Ca^{2+} influx directly via $P2X_2$ channels but not via VDCCs (Nakazawa and Inoue, 1992). Together with this, the most probable interpretation of the results could be that 9-*cis*-RA up-regulates $P2X_2$ receptor mRNAs and proteins, thereby leading to enhancement of $P2X_2$ receptor-mediated Ca^{2+} entry and DA release in PC-12 cells.

In native neurons, activating P2X receptors on the presynapses facilitates the release of neurotransmitters by directing Ca^{2+} influx through P2X receptors (Shigetomi and Kato,

2004). This raises the possibility that retinoids may increase the synaptic effects of ATP in modulating neurotransmitter release in native neurons by up-regulating $P2X_2$ receptors. In the adult brain, relatively high levels of retinoic acid are detected (Werner and Deluca, 2002). In particular, in the hippocampal region it has been shown that molecules required for retinoid signaling pathways are expressed (MacDonald et al., 1990; Werner and Deluca, 2002). These include cellular retinol binding proteins that facilitate retinol uptake into cells; retinal dehydrogenases, which are enzymes for the synthesis of retinoids; and cellular retinoic acid binding proteins, which are thought to deliver atRA to RAR in cell nuclei, as well as RARs and RXRs (Dong et al., 1999). The hippocampus is one of the areas where the roles of $P2X_2$ receptors in facilitating neurotransmitter release have been investigated (Khakh et al., 2003; Shigetomi and Kato, 2004). One can question that retinoid effect on the PC-12 cells is the consequence of the differentiation of PC-12 into neurons. However, morphological differentiation of PC-12 cells by retinoic acid requires a period of greater than 3 weeks, and retinoic acid treatment increased differentiation of nerve growth factor-stimulated PC-12 cells (Boniece and Wagner, 1995). Thus, retinoic acid-induced differentiation of PC-12 cells was suggested to be the consequence of complicated molecular modulations. In fact, we observed up-regulation of $P2X_2$ mRNA within 3 h after retinoids treatment. Hence, the effect of retinoids on $P2X_2$ expression could be a notable factor for the differentiation, but it might be distinguished from differentiation of PC-12 cells. The up-regulation of $P2X_2$ receptors by retinoids may be involved in some of the biological effects of retinoids in neuronal function and synaptic plasticity in the nervous system (Wang et al., 2004).

In the present study, we found that the $P2X_2$ receptor is up-regulated by retinoids as a result of increased transcription most likely mediated by the retinoid-activated RAR heterodimerized with RXR acting on RAREs (presumably DR5- and DR1-responsive elements) in the promoter region of *P2rx2* in neuronal cells. An increase in the expression of $P2X_2$ receptors in neuronal cells has recently been implicated in the development of several pathological states, such as brain ischemia (Cavaliere et al., 2003) and chronic pain (Xu and Huang, 2002), and $P2X_2$ receptor might thus be a target for their treatment. It is noteworthy that in an analysis of the human genomic sequence using TESS, we also found a putative DR5-responsive element in the 5′-flanking region of the human *P2X₂* gene. Together, the present results provide the molecular mechanism underlying the expression of $P2X_2$ receptors and may help in understanding the roles of $P2X_2$ receptors in the regulation of neuronal function, synaptic plasticity, and pathophysiology in the nervous system.

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Extracellular ATP Counteracts the ERK1/2-Mediated Death-Promoting Signaling Cascades in Astrocytes

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KEY WORDS
purinergic receptor; oxidative stress; ERK1/2; src family; protein tyrosine phosphatase

ABSTRACT
Oxidative stress is the main cause of neuronal death in pathological conditions. Hydrogen peroxide (H_2O_2), one of the reactive oxygen species, activates many intracellular signaling cascades including src family and mitogen-activated protein kinases (MAPKs), some of which are critically involved in the induction of cellular damage. We previously showed that H_2O_2-induced cell death in astrocytes and adenosine 5′-triphosphate (ATP), acting on $P2Y_1$ receptors, had a protective effect. Here, we examined the H_2O_2-induced changes in intracellular signaling cascades that promote cell death in astrocytes, showing the molecular mechanisms by which the activation of $P2Y_1$ receptors counteracts such signals. Although H_2O_2 activated three MAPKs including ERK1/2, p38, and JNK, only the activation of ERK1/2 participated in the H_2O_2-evoked cell death. H_2O_2 induced a sustained activation of ERK1/2 mainly in the nucleus region, which was well in accordance with the H_2O_2-induced cell death. H_2O_2 also activated the src tyrosine kinase family, which was an upstream signal for ERK1/2. Activation of $P2Y_1$ receptors by 2methylthio-ADP (2MeSADP) inhibited the H_2O_2-evoked activation of src tyrosine kinase, resulting in the inhibition of the phosphorylated-ERK1/2 accumulation in the nucleus. 2MeSADP enhanced the gene expression and activity of protein tyrosine phosphatase (PTP), which was responsible for the inhibition of src tyrosine kinase. Thioredoxin reductase, another cytoprotective gene we previously showed to be upregulated by 2MeSADP, also controlled the activity of PTP. Taken together, ATP, acting on $P2Y_1$ receptors, upregulates the PTP expression and its activity, which counteracts the H_2O_2-promoted death signaling cascades including ERK1/2 and its upstream signal src tyrosine kinase in astrocytes.

INTRODUCTION

Adenosine 5′-triphosphate (ATP) is an important signaling molecule that mediates gliotransmission and also glia-to-neuron communication in the CNS (Fields and Stevens-Graham, 2002; Hansson and Ronnback, 2003; Inoue, 2002). The $P2Y_1$ receptor has a central role in the ATP-mediated gliotransmission in astrocytes (Fam et al., 2000) and we previously demonstrated that ATP, acting on $P2Y_1$ receptors, protected astrocytes against oxidative stress, indicating the physiological importance of ATP-mediated gliotransmission in astrocytes (Shinozaki et al., 2005). However, details concerning the molecular mechanism(s) by which $P2Y_1$ receptor activation results in such protection are still lacking.

Hydrogen peroxide (H_2O_2), one of the reactive oxygen species (ROS), activates various intracellular signaling cascades including mitogen-activated protein kinases (MAPKs) (Fialkow et al., 1994; Konishi et al., 1999; Ushio-Fukai et al., 1999). The MAPK family includes extracellular signal-regulated kinase 1 and 2 (ERK1/2), p38 kinase, and c-Jun NH_2-terminal kinase (JNK). The latter two members are well known to be stress-responding MAPKs, which are activated by lipopolysaccharide, cytokines, and oxidative stress such as by H_2O_2 and induce cell death (Oppenheim, 1991). ERK1/2 is constitutively expressed in various regions including the CNS (Boulton et al., 1991). ERK1/2 is activated by various neurotransmitters, hormones and growth factors in physiological conditions, controls the transcription factor activity and induces various physiological responses, such as cell proliferation or differentiation (Boulton et al., 1991; Marshall, 1995; Segal and Greenberg, 1996). However, ERK1/2 is also activated by various types of stress such as oxidative stress or shear stress, and appears to control the survival of cells (Guyton et al., 1996; Takahashi and Berk, 1996; Wang et al., 1998; Xia et al., 1995). Concerning neuronal cells, recent reports have shown that activation of ERK1/2 even promotes cell death both in vivo and in vitro (Murray et al., 1998; Namura et al., 2001; Stanciu and DeFranco, 2002; Subramaniam et al., 2004). Thus, although ERK1/2 is an essential intracellular signaling molecule that mediates various physiological functions, it may also mediate the death of cells. The intensity of ERK1/2 activation and spatial and temporal differences in its activation would greatly affect physiological or pathophysiological events in cells.

The src family, a well-known protein tyrosine kinase (PTK), regulates a variety of cellular functions such as cell

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growth, proliferation, and differentiation. The src family is abundantly expressed in the CNS (Brugge et al., 1985; Sugrue et al., 1990), and is also involved in brain injury induced by oxidative stress. In fact, H_2O_2 or ischemia/reperfusion injury activates the src family in the hippocampus (Guo et al., 2003; Ohtsuki et al., 1996). Activation of the src family is prevented by protein tyrosine phosphatase (PTP). PTP is related to various events in the CNS such as inhibition of the NMDA receptor activation in neurons(Yu and Salter, 1999) and of microglial TNFa and nitric oxide generation by amyloid β (Tan et al., 2000). In addition, PTP upregulation is observed in kainic acid-treated (Boschert et al., 1997) and ischemia-injured neurons (Takano et al., 1996). Thus, the activity and expression of PTP is also important for the regulation of pathophysiological cellular functions as well as the regulation of src tyrosine kinase.

On the basis of these findings, we hypothesized that MAPKs and src family are key molecules for promoting the H_2O_2-evoked cell death in astrocytes and that ATP acting on $P2Y_1$ receptors may counteract these cell death-promoting signaling cascades.

In the present study, we demonstrate that activation of both src tyrosine kinase and the subsequent ERK1/2 are key events in the H_2O_2-evoked cell death in astrocytes. We also demonstrate that ATP/$P2Y_1$ receptor activation interferes with the H_2O_2-evoked src family—ERK1/2 cascades by increasing the expression and activity of PTP, thereby leading to protection against H_2O_2-induced cell death in astrocytes.

MATERIALS AND METHODS

Chemicals

Adenosine 5′-triphosphate (ATP), adenosine, 2-methylthio adenosine diphosphate (2MeSADP), bovine serum albumin (BSA), propidium iodide (PI), sodium orthovanadate (Na_3VO_4), MRS2179, and *N*-acetyl cystein were purchased from Sigma Chemical (St Louis, MO). The sources of other chemicals are shown in parentheses as follows; hydrogen peroxide (H_2O_2) (Wako Pure Chemicals, Osaka, Japan), 3-(4,5-dimethylthiazol-2-y1)-2,5-diphenyltetrazolium bromide (MTT) assay kit (CHEMICON International, Temecula, CA), PD98059, U0126, SB203580, SP600125, and PP3 (Calbiochem Biosciences, San Diego), PP1 and PP2 (Biosource, CA), auranofin (Alexis biochemicals, Lausen, Switzerland).

Abbreviations

ATP	adenosine 5′-triphosphate
2MeSADP	2-methylthio-adenosine 5′-diphosphate
ERK1/2	extracellular signal-regulated kinase 1 and 2
H_2O_2	hydrogen peroxide
JNK	c-Jun NH_2-terminal kinase
MAPK	mitogen-activated protein kinase
MAPKP	MAPK phosphatase
MEK1/2	MAPK kinase 1 and 2
Na_3VO_4	sodium orthovanadate
PI	propidium iodide
PTK	protein tyrosine kinase
PTP	protein tyrosine phosphatase
P-Tyr	phosphorylated tyrosine
ROS	reactive oxygen species
TrxR	thioredoxin reductase

Antibodies

Polyclonal antibodies against total ERK1/2, phosphorylated ERK1/2, phosphorylated p38, and phosphorylated JNK were purchased from Cell Signaling Technology (Beverly, MA). The monoclonal antibody against phosphorylated tyrosine was purchased from Sigma Chemical (St Louis, MO).

Cells and Cell Culture

Astrocytes were prepared from neonatal rat forebrain. The cells were cultured as previously reported (Shinozaki et al., 2005). For the cell viability assay, cells were seeded on 96-well plates (NUNC, Roskilde, Denmark) at a density of 1.25×10^4 cells/well.

Cell Viability Assay

For the cell viability assay, we used an MTT assay as previously reported (Shinozaki et al., 2005). A 1/10 volume of MTT solution (5 mg/mL in PBS) was added and incubated for 4 h under 10% CO_2/90% air at 37°C. Then, an equal volume of isopropanol (with 0.04 N HCl) was added to the cells. The absorbance was measured on an ELISA plate reader (ASYS Hitech, Eugendorf, Austria) with a test and reference wavelength of 570 and 630 nm, respectively.

Western Blotting

Astrocytes were prepared as described above. After H_2O_2-stimulation, cells were lysed and the lysates were resolved with 10% SDS-PAGE gels and transferred to PVDF membranes. The membranes were blocked for 1 h in Tris-buffered saline containing 0.1% Tween-20 (TBS/T) and 5% non-fat dry milk at room temperature. Then the membranes were incubated with primary antibody dilution buffer (1:1000 dilution into TBS/T containing 5% BSA) overnight at 4°C. After three washes with TBS/T, the membranes were incubated with horseradish peroxidase-conjugated anti-rabbit antibody (1:2000 dilution into TBS/T containing 5% non-fat dry milk) for 1 h at room temperature. The membranes were washed with TBS/T three times, and the proteins were visualized by chemiluminescence. The antibodies for anti-phospho-proteins used in the present study (anti-P-ERK1/2, P-p38, and P-JNK or P-Tyr antibodies) specifically detected only the activated and phosphorylated form of the proteins. To detect total ERK1/2, the aliquot of the same sample was resolved with 10% SDS-PAGE gels, transferred to PVDF membranes in the same conditions and exposed to anti-total ERK1/2 antibody.

Quantification of the Intensity of P-ERK1/2 Bands

To quantify the intensity of P-ERK1/2 bands, we used Image J (http://rsb.info.nih.gov/ij/). P-ERK1/2 bands were selected by rectangular selection. Then, we selected *Analyze-Gels-Select First Lane* from the menu bar. The

area corresponding to each band was measured using *Wand (tracing) tool* from the tool bar.

Immunocytochemistry

After each treatment, the cells were fixed for 30 min at room temperature in 3.7% paraformaldehyde. The fixed cells were permeabilized with PBS containing 0.1% Triton X-100 for 5 min at room temperature and then incubated with the polyclonal anti-ERK1/2 and anti-phospho-ERK1/2 antibodies for 24 h at 4°C. After washing, the cells were incubated with the appropriate secondary antibodies conjugated to Alexa 488 or 546, washed again, and mounted on glass coverslips (Matsunami Glass, Osaka, Japan). Astrocytes for immunocytochemistry were selected randomly. Images were collected in an MRC-1024 laser-scanning microscope (Bio-Rad) with 20× objective lenses. For the comparison of double-stained patterns, images were processed using Photoshop 5 (Adobe System, Mountain View, CA).

Tyrosine Phosphatase Assay

The tyrosine phosphatase activity was measured using a universal tyrosine phosphatase kit (Takara, Shiga, Japan). The measurement was done according to the manufacturer's instructions. Cells were lysed by lysis buffer and transferred into 96-well ELISA plates at a volume of 50 μL/well followed by incubation for 45 min at 37°C. After four times washing with tween-PBS (PBS containing 0.05% tween20), blocking buffer was added to the wells at a volume of 100 μL/well followed by 30 min incubation at 37°C. Then, the blocking buffer was discarded and coloring substrates were added to the wells (100 μL/well). After a 15 min incubation at room temperature, 1 N sulfuric acid was added to the wells (100 μL/well) to stop the reaction. The absorbance was measured by a plate reader (ASYS Hitech, Eugendorf, Austria) at a test wavelength of 450 nm.

Quantitative RT-PCR of PTP Genes

RT-PCR amplifications were performed using Taqman One-step RT-PCR Master Mix Reagents and 200 nM PTP specific primers as previously reported (Shinozaki et al., 2005). Using the computer software Primer Express (Applied Biosystems), clone-specific primers were designed to recognized rat PTP genes, i.e., rat PTP4a1 (Taqman probe, 5′-acacaatccaaccaatgcgaccttaaacaa-3′; forward, 5′-tgctcctgtggaagtcacataca-3′; reverse, 5′-gtcgtaagttgcttcgcatactctta-3′) and rat PTPro (Taqman probe, 5′-ccgctatacaaacatcctgccgtacgactt-3′; forward, 5′-ttccgctgaaccgatgtaaaa-3′; reverse, 5′-tgaggtgagttgtagccaggaata-3′). RT-PCR was performed by 30 min reverse transcription at 48°C, 10 min Amplitaq Gold activation at 95°C, then 15-s denaturation at 95°C, 1 min annealing and elongation at 60°C for 40 cycle in a PRISM7700 (Applied Biosystems). Each experiment was performed in triplicate.

RESULTS

MEK1/2 Inhibitors Protect Astrocytes Against H_2O_2-Evoked Cell Death

Figure 1A shows the effect of H_2O_2 on the activation of MAPKs in astrocytes. Western blotting analysis revealed that stimulation of astrocytes with 250 μM H_2O_2 for 2 h resulted in the activation of three MAPKs, i.e., ERK1/2, p38, and JNK (Fig. 1A). We then tested pharmacologically whether the activation of these MAPKs is involved in the H_2O_2-evoked cell death in astrocytes. The H_2O_2-induced decrease in the cell viability of the astrocytes obtained from the MTT assay was always accompanied by the activation of caspase-3, DNA damage, and nucleus condensation (data not shown). Thus, we defined the decrease in cell viability as cell death in astrocytes in the following experiments. MAPK kinase 1/2 (MEK1/2) activates ERK1/2. The MEK1/2 inhibitors PD98059 (10 μM) (Alessi et al., 1995) and U0126 (20 μM) (Favata et al., 1998) strongly inhibited the H_2O_2-evoked cell death (Fig. 1B). The inhibitory

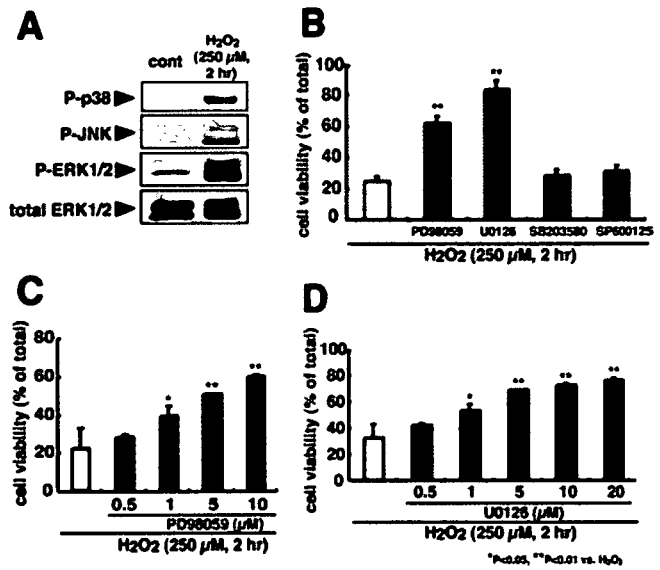

Fig. 1. The effect of MAPK inhibitors on the H_2O_2-evoked cell death of astrocytes. **A**: The effect of H_2O_2 on MAPK activation in astrocytes. H_2O_2 (250 μM, 2 h) activated p38, JNK, and ERK1/2. **B**: The effect of MAPK inhibitors on H_2O_2-evoked cell death. The MEK1/2 inhibitors PD98059 (10 μM) and U0126 (20 μM) strongly inhibited the H_2O_2-evoked cell death. Neither the JNK inhibitor SP600125 (20 μM) nor the p38 inhibitor SB203580 (20 μM) affected the H_2O_2-evoked cell death. **C**: The concentration-dependent effect of PD98059 against H_2O_2-evoked cell death. The protective effect of PD98059 was dose-dependent in a concentration range from 0.5 to 10 μM. **D**: The concentration-dependent effect of U0126 against H_2O_2-evoked cell death. The protective effect of U0126 was dose-dependent in a concentration range from 0.5 to 20 μM. Neither inhibitor alone affected the cell viability of the astrocytes. Inhibitors added to the cells 1 h before H_2O_2 treatment. Asterisks show significant difference from the response evoked by H_2O_2 (*$P < 0.05$, **$P < 0.01$ vs. H_2O_2 alone, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$).

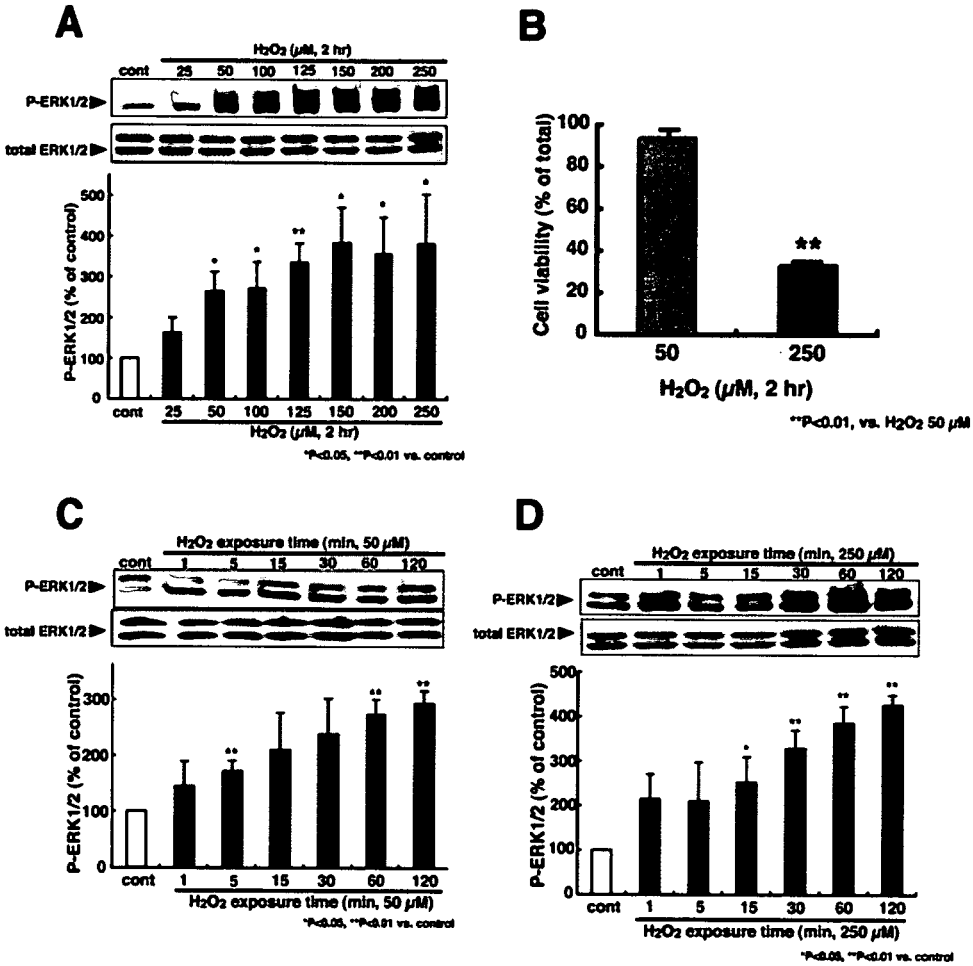

Fig. 2. The concentration-dependency and time course of the H_2O_2-evoked ERK1/2 activation in the astrocytes. **A**: The concentration-dependency of the H_2O_2-evoked ERK1/2 activation. H_2O_2 activated ERK1/2 in a concentration-dependent fashion at the whole cell level (0–250 μM, 2 h). H_2O_2 did not affect the amount of total ERK1/2. Asterisks show significant difference from control (*$P < 0.05$, **$P < 0.01$ vs. control, Student's t-test). **B**: The degree of the decrease of the cell viability of H_2O_2-treated astrocytes markedly decreased the cell viability in the astrocytes 250 μM but not at 50 μM H_2O_2. Asterisks show significant difference from the response evoked by H_2O_2 (**$P < 0.01$ vs. 50 μM H_2O_2, Student's t-test). **C** and **D**: A temporal analysis of the H_2O_2-evoked ERK1/2 activation in the astrocytes. At the whole cell level, ERK1/2 was activated by H_2O_2 treatment time dependently (0–120 min) despite the concentration of H_2O_2 (50 and 250 μM). H_2O_2 did not affect the total ERK1/2. Asterisks show significant difference from control (*$P < 0.05$, **$P < 0.01$ vs. control, Student's t-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$).

effects by PD98059 and U0126 were dose-dependent in a concentration-range from 0.5 to 10 μM (Fig. 1C) and 0.5 to 20 μM (Fig. 1D), respectively. In contrast, neither the p38 inhibitor SB203580 (Alessandrini et al., 1999; McLaughlin et al., 1996) nor the JNK inhibitor SP600125 (20 μM) (Bennett et al., 2001) had any effect on the H_2O_2-evoked cell death.

The Concentration-Dependency and Time Course of H_2O_2-Evoked ERK1/2 Activation

Among the H_2O_2-activated MAPKs tested, only ERK1/2 was involved in the H_2O_2-evoked cell death (Fig. 1B). H_2O_2 activated ERK1/2 in a concentration- and exposure time-dependent fashion at the whole cell level (Figs. 2A,C,D). Although a lower H_2O_2 concentration (50 μM) activated ERK1/2, H_2O_2 at this concentration did not cause cell death (Fig. 2B). At a higher concentration, H_2O_2 (250 μM) evoked ERK1/2 phosphorylation and cell death in astrocytes (Figs. 2A,B). The phosphorylation of ERK1/2 evoked by 250 μM H_2O_2 was stronger than that evoked by 50 μM H_2O_2. H_2O_2 at either concentration did not affect the total ERK1/2 at the whole cell level.

The Temporal and Spatial Aspect of P-ERK1/2 Induced by H_2O_2

We investigated the temporal and spatial distribution of P-ERK1/2 using immunocytochemical techniques. H_2O_2-evoked ERK1/2 activation was observed in GFAP-positive astrocytes (data not shown). It was reported that ERK1/2 is translocated into the nucleus from the cytoplasm when it is activated (Chen et al., 1992; Gonzalez et al., 1993). We thus analyzed the distribution of P-ERK1/2 after H_2O_2 stimulation. When the cells were stimulated with 250 μM H_2O_2 for 2 h, P-ERK1/2 signals were observed in the center part of individual astrocytes and were colocalized with the signals of PI, a DNA binding dye, suggesting that P-ERK1/2 had translocated into the nucleus (Fig. 3A). In contrast, when stimulated with 50 μM H_2O_2 for 2 h which activated ERK1/2 but did not induce cell death, the P-ERK1/2 signals were observed but were not colocalized with the PI signals (Fig. 3A). The total amount of ERK1/2 signals, however, was not affected by H_2O_2 (50 and 250 μM), and they were not colocalized with the PI signals (Fig. 3B). Without H_2O_2 stimulation, the P-ERK1/2 signals were too low (Figs. 2A,D and 3E, cont.) to detect. To quantify the degree of the colocalization of the P-ERK1/2 and PI signals, we

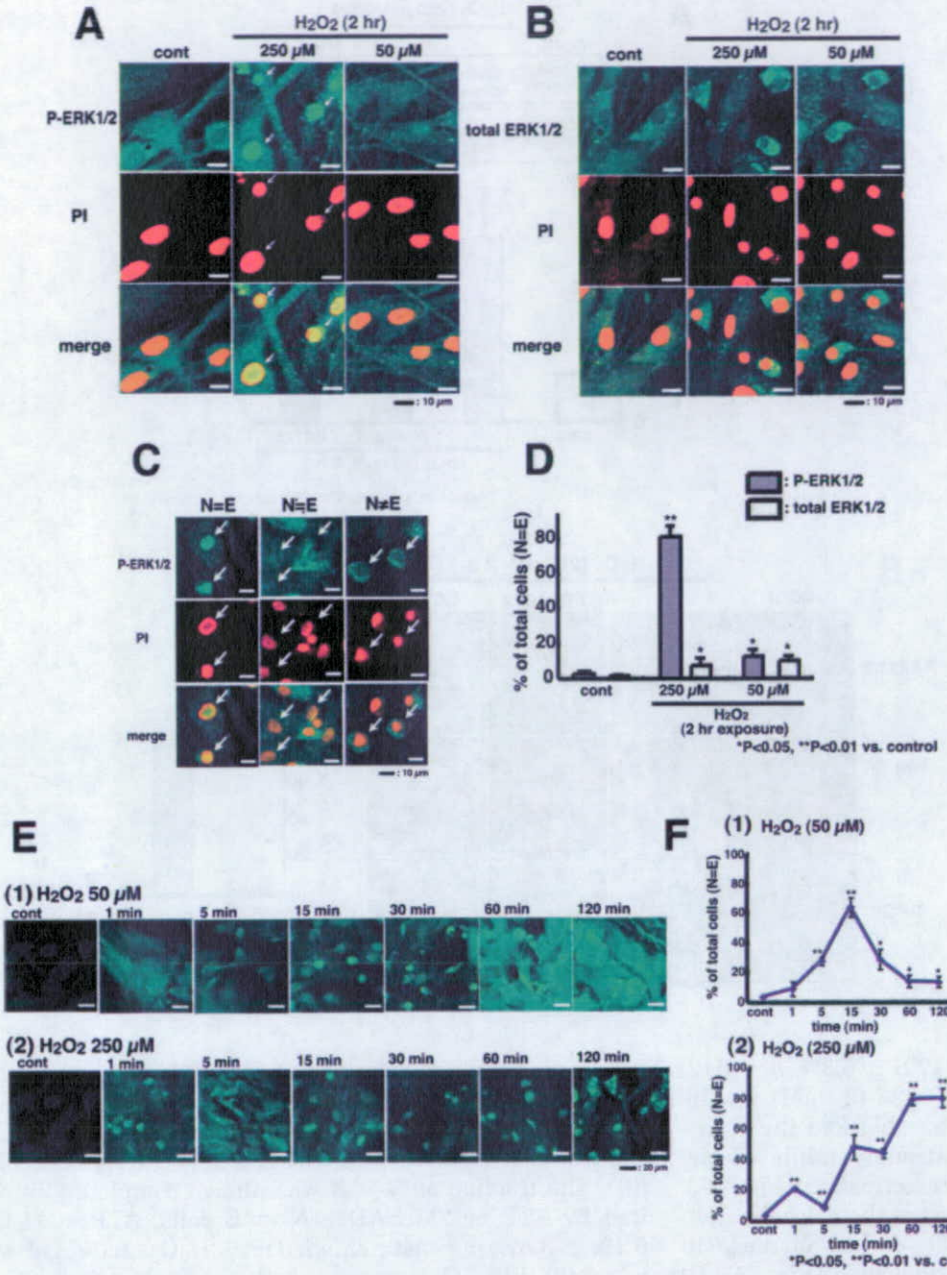

Fig. 3. The effect of H_2O_2 on the intracellular localization of ERK1/2 in astrocytes. **A**: The effect of H_2O_2 on the intracellular localization of P-ERK1/2. In control, the P-ERK1/2 and PI signals did not colocalize (left). When the cells were treated with 250 μM H_2O_2 (2 h), a large number of P-ERK1/2 and PI signals colocalized (center, arrow). At 50 μM H_2O_2, the P-ERK1/2 and PI signals did not colocalize (right). **B**: The effect of H_2O_2 on the intracellular localization of total ERK1/2. Total ERK1/2 did not colocalize with the PI signals irrespective of the H_2O_2 treatment (control, 50 and 250 μM). The signals of P-ERK1/2 were enhanced by photoshop to clarify their intracellular localization. **C**: Classification of the H_2O_2-treated cells into three groups. Astrocytes in which P-ERK1/2 signals were colocalized with PI signals were defined as "N = E" (left), those in which P-ERK1/2 signals were partly colocalized with PI signals were defined as "N =E" (center), and those in which the P-ERK1/2 signals were not colocalized with the PI signals were defined as "N ≠ E" (right). **D**: Quantification of P-ERK1/2 localization into the nucleus. When cells were stimulated by 250 μM H_2O_2, the colocalization of P-ERK1/2 and PI was observed in most cells but not when they were stimulated by 50 μM H_2O_2. Asterisks show significant difference from control ($^*P < 0.05$, $^{**}P < 0.01$ vs. control, Student's *t*-test). **E** and **F**: The temporal analysis of the colocalization of the P-ERK1/2 and PI signals. At 50 μM H_2O_2, N = E cells increased transiently (5–30 min after H_2O_2 stimulation) (E(1), F(1)). At 250 μM H_2O_2, N = E cells increased time-dependently (E(2), F(2)). The DNA binding dye PI was used for identifying the nuclear region. Asterisks show significant difference from control ($^*P < 0.05$, $^{**}P < 0.01$ vs. control, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements.

classified the cells into three groups, i.e., cells in which P-ERK1/2 signals were colocalized with PI signals (defined as "N = E"), those in which the P-ERK1/2 signals were not colocalized with the PI signals (defined as "N ≠ E"), and P-ERK1/2 signals were partly colocalized with IP signals (defined as "N = E") (Fig. 3C). When stimulated with 50 and 250 μM H_2O_2 for 2 h, the fraction of N = E was 12.7% ± 3.8% (n = 304) and 81.0% ± 6.3% (n = 612), respectively. Without H_2O_2 stimulation, no colocalization of P-ERK1/2 and PI was observed in almost any of the cells (N = E cells, 3% ± 0.2%, n = 346) (Fig. 3D). In contrast to P-ERK1/2, most of the total ERK1/2 signals did not colocalize with PI irrespective of H_2O_2 stimulation(N = E cells, control, 1.3% ± 0.8%, n = 225; 250 μM H_2O_2, 6.5% ± 4.0%, n = 545; 50 μM H_2O_2, 10.7% ± 1.2%, n = 229). Furthermore, we analyzed the time course of the fraction of "N = E" cells after H_2O_2 stimulation. When stimulated with 50 μM H_2O_2, the N = E fraction peaked at 15 min after the stimulation, and the fraction decreased to the prestimulated level after 120 min (N = E cells, 1 min, 7.8% ± 5.4%, n = 201; 5 min, 27.8% ± 2.9%, n = 213; 15 min, 66.6% ± 7.1%, n = 216; 30 min, 29.1% ± 7.4%, n = 226; 60 min, 13.1% ± 4.5%, n = 254; 120 min, 12.7% ± 3.8%, n = 405). In contrast, when stimulated with 250 μM H_2O_2, the N = E fraction gradually increased, reached the maximal level at 60 min, and remained even 120 min after the stimulation (N = E cells, 1 min, 20.7% ± 7%, n = 354; 5 min, 8.1% ± 0.7%, n = 213; 15 min, 48.4% ± 8.3%, n = 220; 30 min, 44.2% ± 3.3%, n = 214; 60 min,

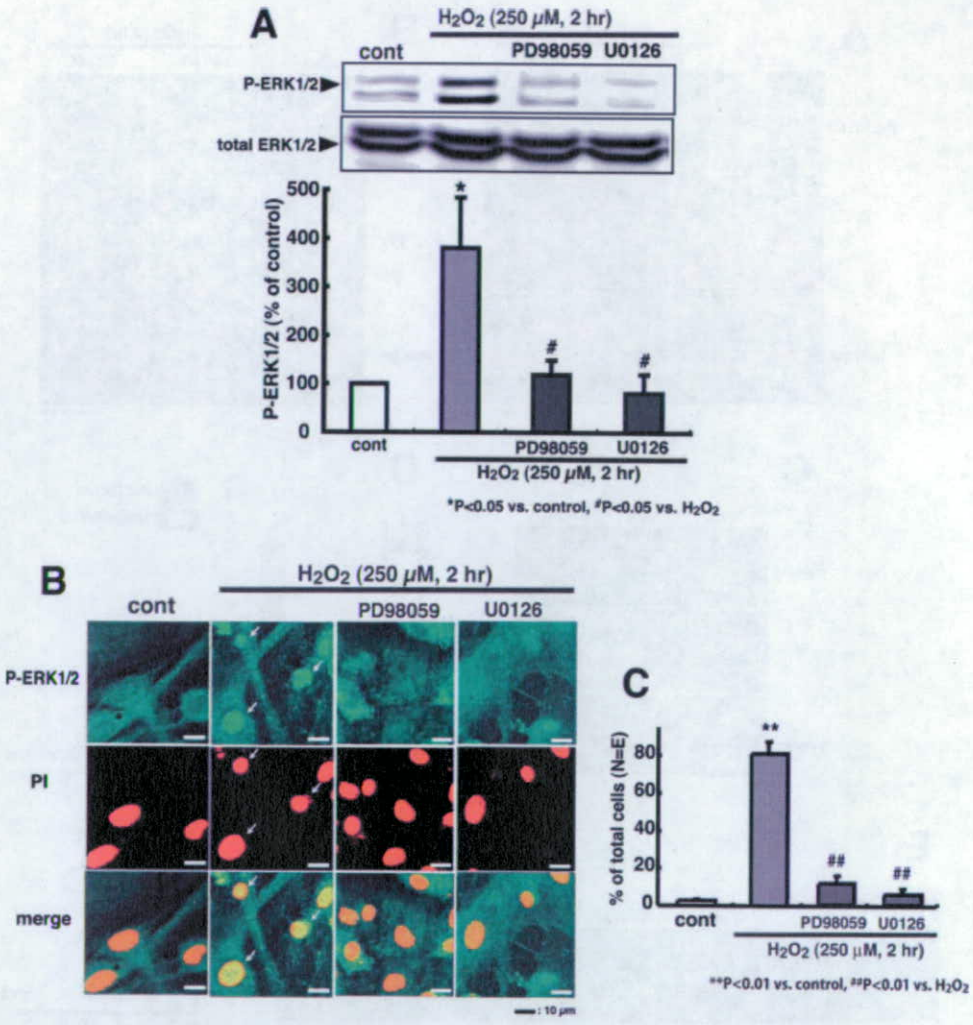

Fig. 4. The effect of MEK1/2 inhibitors on H_2O_2-evoked ERK1/2 activation and P-ERK1/2 translocation. **A**: The effect of MEK1/2 inhibitors on H_2O_2-evoked ERK1/2 activation. PD98059 (10 μM) and U0126 (20 μM) strongly inhibited the H_2O_2-evoked ERK1/2 activation. Asterisks show significant difference from control ($^*P < 0.05$ vs. control, Student's *t*-test). Sharps show significant difference from H_2O_2 ($^\#P < 0.05$ vs. H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$). **B** and **C**: The effect of the MEK1/2 inhibitors on the H_2O_2-evoked P-ERK1/2 translocation. PD98059 (10 μM) and U0126 (20 μM) blocked the colocalization of the P-ERK1/2 and PI signals evoked by H_2O_2 (250 μM, 2 h). In the immunocytochemical analysis, the signals of P-ERK1/2 were enhanced by photoshop to clarify their intracellular localization. The DNA binding dye PI was used for identifying the nuclear region. The MEK1/2 inhibitors were applied to the cells 1 h before and during H_2O_2 treatment. Asterisks show significant difference from control ($^{**}P < 0.01$ vs. control, Student's *t*-test). Sharps show significant difference from H_2O_2 ($^{\#\#}P < 0.01$ vs. H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements.

80.2% ±3.8%, $n = 205$; 120 min, 81%S ± 6.3%, $n = 612$) (Fig. 3F). PD98059 (10 μM) and U0126 (20 μM), at the concentrations that the two inhibitors blocked the H_2O_2-induced cell death (Figs. 1C,D), strongly inhibited the H_2O_2 (250 μM, 2 h)-evoked ERK1/2 activation (Fig. 4A). In addition, these inhibitors prevented the colocalization of P-ERK1/2 and PI, i.e., the H_2O_2-evoked increase in the fraction of N = E was almost abolished (Figs. 4B,C) (N = E cells, PD98059 + H_2O_2, 11.7% ± 3.8%, $n = 524$; U0126 + H_2O_2, 5.9% ± 2.9%, $n = 400$).

ATP Inhibits the H_2O_2-Evoked Activation of ERK1/2 and Its Localization of P-ERK1/2 in the Nucleus

Our previous report by Shinozaki et al., demonstrated that ATP and 2MeSADP inhibited the H_2O_2-induced cell death in astrocytes (Shinozaki et al., 2005). Thus, we examined the effect of ATP and 2MeSADP on the H_2O_2-evoked ERK1/2 activation. In astrocytes pretreated with ATP (100 μM) or 2MeSADP (1 μM) for 24 h, the H_2O_2-induced ERK1/2 activation was markedly inhibited (Fig. 5A). We also analyzed the effect of ATP/2MeSADP on the spatiotemporal behavior of P-ERK1/2 in astrocytes. Immunocytochemical studies showed that pretreatment of the cells with ATP/2MeSADP prevented the H_2O_2-evoked colocalization of P-ERK1/2 and PI signals (Fig. 5B). The fraction of N = E was almost completely inhibited by ATP or 2MeSADP (N = E cells, ATP + H_2O_2, 5.1% ± 3.3%, $n = 340$; 2MeSADP + H_2O_2, 1.1% ± 1.4%, $n = 342$) (Fig. 5C).

ATP itself is known to activate ERK1/2 in some cells including astrocytes (Neary et al., 1999, 2003). Using Western blotting analysis, we found that both ATP (100 μM) and 2MeSADP (1 μM) activated ERK1/2 but the activation was only transient (lasting 1–15 min after stimulation) and returned to the prestimulated level within 120 min [Figs. 6A(i,ii)]. Thus, after the initial phosphorylation of ERK1/2 evoked by ATP/2MeSADP the activation should have returned to the prestimulated level when the astrocytes were stimulated with H_2O_2 24 h after ATP-treatment. Interestingly, when 2MeSADP was pretreated with PD98059 (10 μM) or U0126 (20 μM), the 2MeSADP-induced cytoprotective effects against H_2O_2 disappeared (Fig. 6B). Furthermore, using quantitative RT-PCR, we found that U0126 inhibited the 2MeSADP (1 μM, 2 h)-induced upregula-

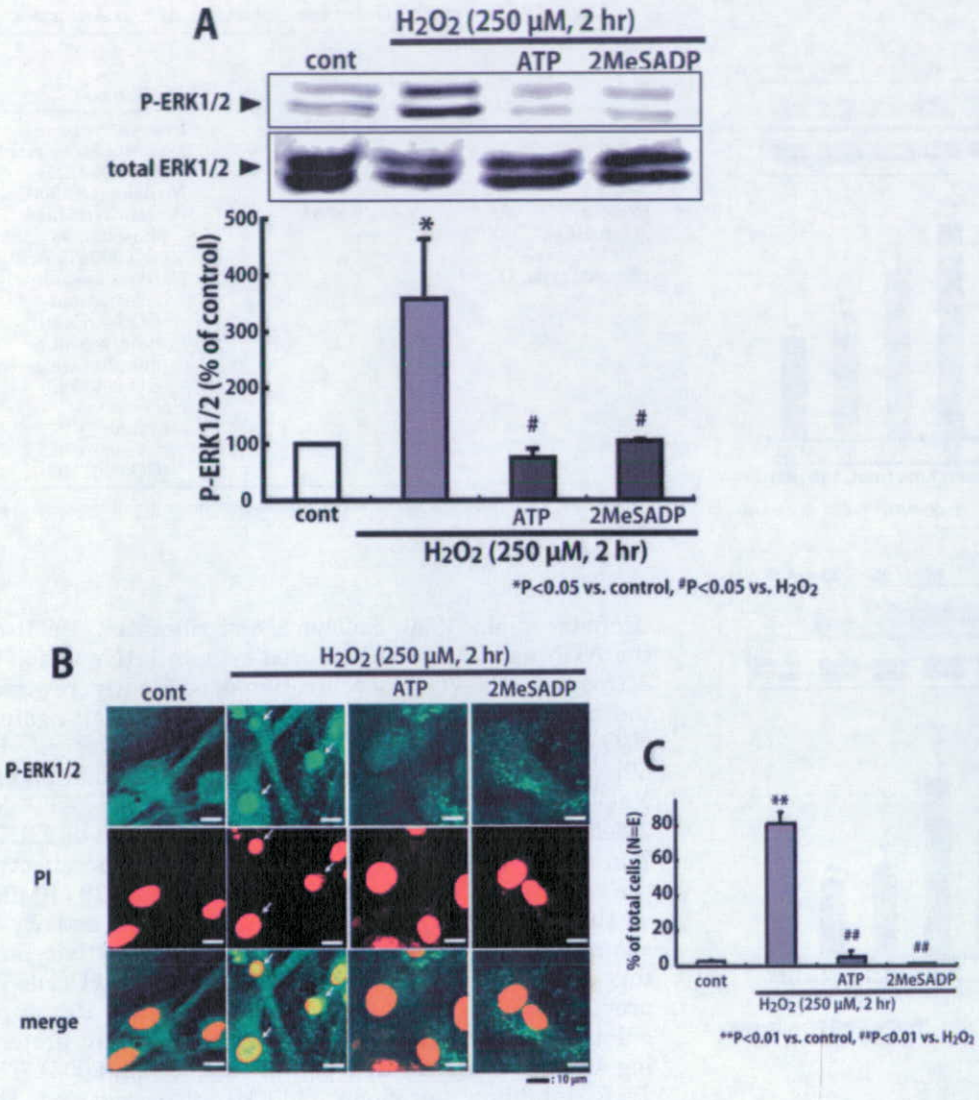

Fig. 5. The effect of ATP and 2MeSADP on H_2O_2-evoked ERK1/2 and P-ERK1/2 translocation. **A**: The effect of ATP and 2MeSADP on H_2O_2-evoked ERK1/2 activation. ATP (100 μM) and 2MeSADP (1 μM) strongly inhibited the H_2O_2-evoked ERK1/2 activation at the whole cell level. Asterisks show significant difference from control ($^*P < 0.05$ vs. control, Student's *t*-test). Sharps show significant difference from H_2O_2 ($^\#P < 0.05$ vs. H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$). **B** and **C**: The effect of ATP and 2MeSADP on H_2O_2-evoked P-ERK1/2 translocation. ATP (100 μM) and 2MeSADP (1 μM) abolished the colocalization of the P-ERK1/2 and PI signals evoked by H_2O_2 (250 μM, 2 h). In the immunocytochemical analysis, the signals of P-ERK1/2 were enhanced by photoshop to clarify their intracellular localization. The DNA binding dye PI was used for identifying the nuclear region. ATP and 2MeSADP was applied to the cells 24 h before and during H_2O_2 treatment. Asterisks show significant difference from control ($^{**}P < 0.01$ vs. control, Student's *t*-test). Sharps show significant difference from H_2O_2 ($^{\#\#}P < 0.01$ vs. H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements.

tion of thioredoxin reductase (TrxR) (2MeSADP, 232.2% ± 54.4% of control, $P < 0.01$ vs. control; 2MeSADP + U0126, 114.9% ± 9.4% of control, $P < 0.05$ vs. 2MeSADP alone; $n = 3$) and PTPs (PTP4a1: 2MeSADP, 291.1% ± 78.5% of control, $P < 0.05$ vs. control, 2MeSADP + U0126, 102.8% ± 12.2% of control, $P < 0.05$ vs. 2MeSADP alone; $n = 3$; PTPro: 2MeSADP, 608.0% ± 153.8% of control, $P < 0.01$ vs. control; 2MeSADP + U0126, 264.7% ± 47.7% of control, $P < 0.05$ vs. 2MeDADP alone; $n = 3$). U0126 alone did not affect the expression level of TrxR (94.5% ± 30.8% of control; $n = 4$) and PTP genes (PTP4a1, 107.5% ± 21.7% of control; PTPro, 113.6% ± 14.3% of control, $n = 4$). The MEK1/2 inhibitors were added to the cells 1 h before and during 2MeSADP-treatment, and was washed out before the H_2O_2 stimulation. Thus, the phosphorylation of ERK1/2 induced by H_2O_2, represented by the sustained and intense responses seen mainly in the nucleus appeared to cause cell death in the astrocytes, while the ATP-induced transient phosphorylation of ERK1/2 appeared to have a cytoprotective action.

ATP Increases PTP Expression and Its Activity, Leading to Protection Against the H_2O_2-Evoked Cell Death in Astrocytes

We comprehensively studied whether ATP induces the expression of genes that could regulate ERK1/2 activity using a GeneChip microarray. We expected that ATP might upregulate the expression of genes that dephosphorylate ERK1/2 such as MAPK phosphatase (MAPKP), which dephosphorylates ERK1/2, thereby leading to the inactivation of ERK1/2 and the cytoprotective action by ATP. However, no upregulation of any MAPKPs in the GeneChip microarray was observed. Instead, we found the ATP (100 μM, 2 h)-induced upregulation of PTP genes containing PTP4a1 and PTP, receptor type O (PTPro) (Table I).

In the PTP activity assay, both ATP (100 μM, 24 h) and 2MeSADP (1 μM, 24 h) significantly increased the PTP activity (ATP, 148.3% ± 8.8%; 2MeSADP, 168.8% ± 4.8%) (Fig. 7A). Subsequently, we analyzed the effect of the PTP inhibitor sodium orthovanadate (Na_3VO_4)

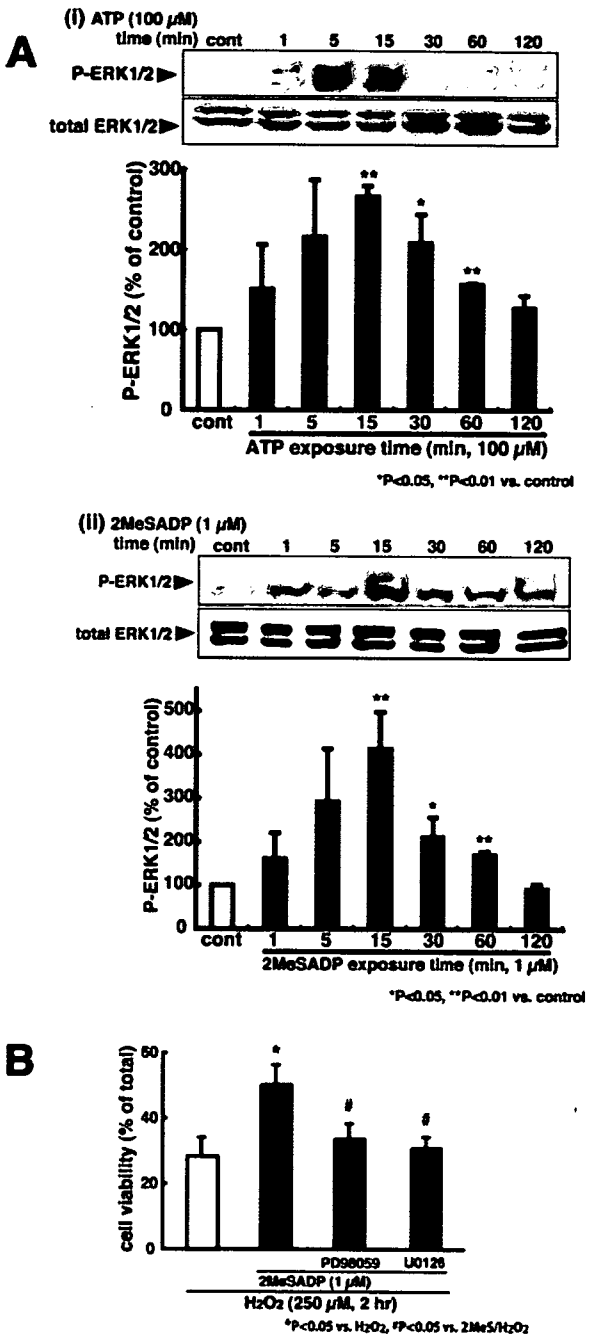

Fig. 6. Analysis of ATP-stimulated ERK1/2 activation in astrocytes. **A**: The time course of ATP-induced ERK1/2 activation. ATP (100 μM) activated ERK1/2 transiently (1–15 min after stimulation) (i). 2MeSADP (1 μM) also activated ERK1/2 transiently (1–15 min after stimulation) (ii). ATP and 2MeSADP did not affect the total amount of ERK1/2. Asterisks show significant difference from control ($^*P < 0.05$, $^{**}P < 0.01$ vs. control, Student's *t*-test). **B**: The effect of 2MeSADP-stimulated ERK1/2 on the $P2Y_1$ receptor-mediated cytoprotective effect. The MEK1/2 inhibitors PD98059 (10 μM) and U0126 (20 μM) strongly inhibited the protective effect induced by 2MeSADP (1 μM, 24 h). The MEK1/2 inhibitors were added to the cells 1 h before the 2MeSADP treatment and were washed out before the H_2O_2 treatment. Asterisks show significant difference from the response evoked by H_2O_2 ($^*P < 0.05$ vs. 250 μM H_2O_2, Student's *t*-test). Sharps show significant difference from 2MeSADP/H_2O_2 ($^\#P < 0.05$ vs. 2MeSADP/H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$).

TABLE 1. List of PTP Genes Upregulated by ATP in Astrocytes

Title	Fold increase (RT-PCR)	Gene ontology[a]
Protein tyrosine phosphatase 4a1	1.7 (5.7)	Protein tyrosine phosphatase activity (GO:0004725) Nucleus (GO:0005634)
Protein tyrosine phosphatase, receptor type, O	1.8 (4.4)	Protein tyrosine phosphatase activity (GO:0004725) Nervous system development (GO:0007399)
	1.6	Protein tyrosine phosphatase activity (GO:0004725) Nervous system development (GO:0007399)

[a]GO ontology defined by Gene Ontology Consortium (www.godatabase.org/cgi-bin/amigo/go.cgi).

(Heffetz et al., 1990; Shisheva and Shechter, 1993) on the ATP- and 2MeSADP-induced cytoprotective action in astrocytes. Na_3VO_4 concentration-dependently reversed the cytoprotective effect by ATP and 2MeSADP against H_2O_2 (Fig. 7B). Na_3VO_4 alone did not affect the cell viability of the astrocytes (light gray column). Additionally, Na_3VO_4 also reversed the inhibition by ATP or 2MeSADP of the H_2O_2-evoked phosphorylation of ERK1/2 in astrocytes (Fig. 7C). Then, we studied the effect of the selective $P2Y_1$ receptor antagonist MRS2179 (10 μM) on the PTP activity. H_2O_2 decreased the PTP activity to about one half. 2MeSADP restored the PTP activity and this effect was reversed by MRS2179 [Fig. 7D(1)]. As previously reported, $P2Y_1$ receptor activation also upregulates oxidoreductases such as TrxR, thereby protecting astrocytes against H_2O_2 (Shinozaki et al., 2005). The TrxR inhibitor auranofin (1 μM) also reversed the 2MeSADP-restored PTP activity. Furthermore, the thiol-containing antioxidant *N*-acetyl cystein (NAC, 10 mM) restored the H_2O_2-decreased PTP activity. In addition, we studied the effect of 2MeSADP, MRS2179 (10 μM), auranofin (1 μM), and NAC (10 mM) on the H_2O_2-evoked ERK1/2 activation [Fig. 7D(2)]. The inhibition of P-ERK1/2 activation by 2MeSADP (1 μM) was reversed by MRS2179 (10 μM) and auranofin (1 μM). In contrast, the H_2O_2-evoked ERK1/2 activation was prevented by NAC (10 mM). MRS2179 was added to the cells 1 h before the 2MeSADP treatment. Auranofin and NAC were added to the cells 1 h before H_2O_2 stimulation.

Involvement of src Tyrosine Kinase Family on H_2O_2-Evoked Cell Death and ERK1/2 Activation

Using Western blotting analysis, we studied whether H_2O_2 induces protein tyrosine phosphorylation. H_2O_2 evoked protein tyrosine phosphorylation, which was inhibited by pretreatment with ATP (100 μM, 24 h) or 2MeSADP (1 μM, 24 h) (Fig. 8A). This ATP- and

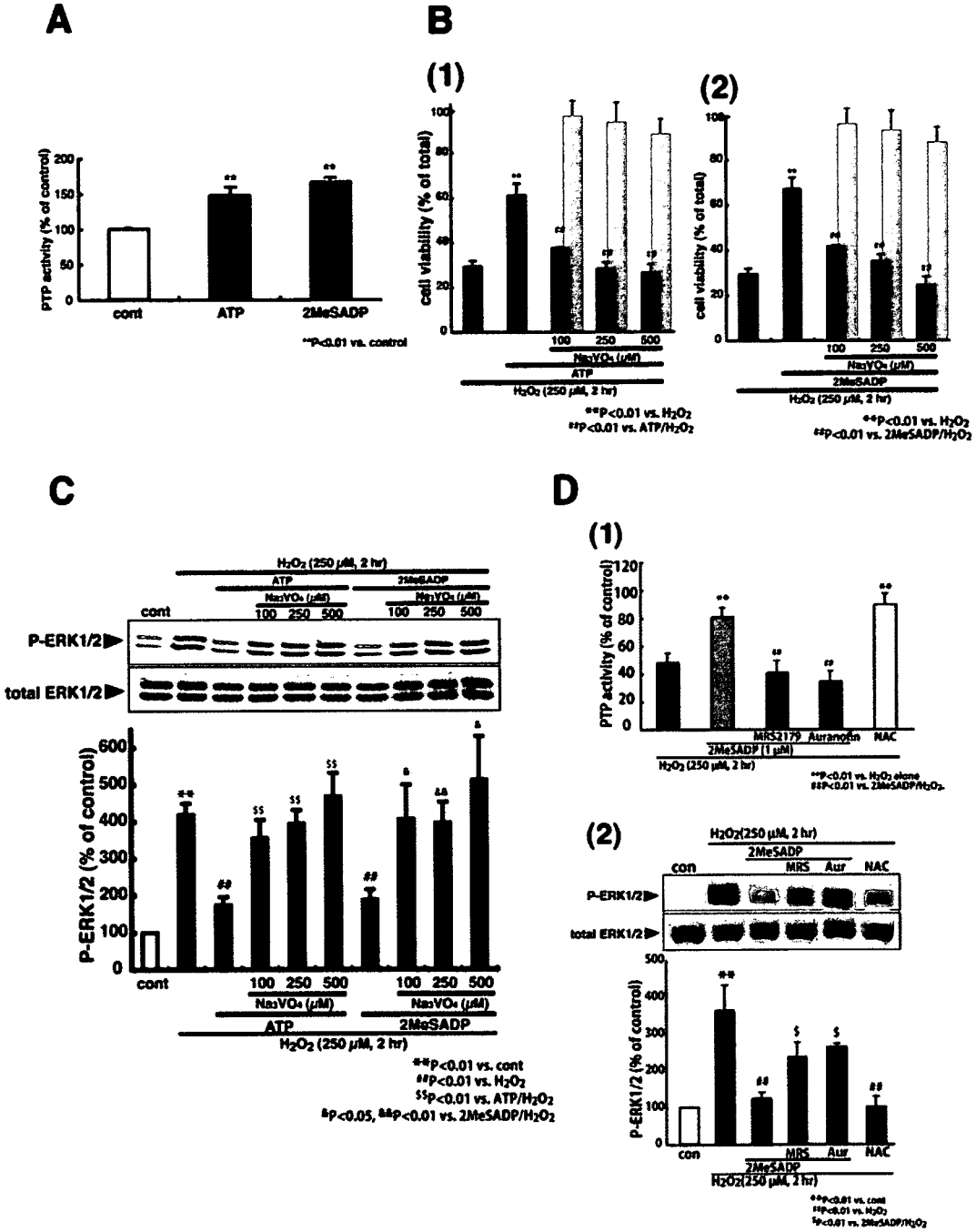

Fig. 7. The effect of ATP-induced PTP upregulation on H_2O_2-evoked cell death and ERK1/2 activation. **A**: ATP- and 2MeSADP-induced an increase of the PTP activity. ATP (100 μM, 24 h) and 2MeSADP (1 μM, 24 h) significantly increased the PTP activity (ATP: 148.3% ± 8.8%; 2MeSADP: 168.8% ± 4.8%, vs. control). Asterisks show significant difference from control (**$P < 0.01$ vs. control, Student's t-test). **B**: PTP participates in ATP- and 2MeSADP-induced cytoprotective effect. The PTP inhibitor Na_3VO_4 (100–500 μM, 1 h) reversed the ATP(100 μM, 24 h)- and 2MeSADP(1 μM, 24 h)-induced protective effect concentration-dependently (black column). Na_3VO_4 alone did not affect the cell viability of the astrocytes (light gray column). Asterisks show significant difference from the response evoked by H_2O_2 (**$P < 0.01$ vs. 250 μM H_2O_2, Student's t-test). Sharps show significant difference from the response by ATP/H_2O_2 or $2MeSADP/H_2O_2$ (##$P < 0.01$ vs. ATP/H_2O_2 or $2MeSADP/H_2O_2$, Student's t-test). **C**: PTP participates in ATP- and 2MeSADP-induced inhibition of H_2O_2-evoked ERK1/2 activation. Na_3VO_4 (100–500 μM, 1 h) reversed the ATP(100 μM, 24 h)- and 2MeSADP(1 μM, 24 h)-induced inhibition of H_2O_2-evoked ERK1/2 activation. The cells were treated with ATP and 2MeSADP 24 h before and during H_2O_2 treatment. Na_3VO_4 was added to the cells 1 h before the H_2O_2 (250 μM) treatment. Asterisks show significant difference from control (**$P < 0.01$ vs. control, Student's t-test). Sharps show significant difference from H_2O_2 (##$P < 0.01$ vs. H_2O_2, Student's t-test). Dollar marks show significant difference from ATP/H_2O_2 or $2MeSADP/H_2O_2$ ($$P < 0.05$, $$$P < 0.01$ vs. ATP/H_2O_2 or $2MeSADP/H_2O_2$, Student's t-test). **D**: The effect of MRS2179, auranofin, and NAC on PTP and ERK1/2 activity. (1) 2MeSADP (1 μM) restored the H_2O_2 (250 μM, 2 h)-decreased PTP activity and the effect was reversed by MRS2179 (10 μM) and auranofin (1 μM). NAC (10 mM) restored the PTP activity decreased by H_2O_2. Asterisks show significant difference from the response evoked by H_2O_2 (**$P < 0.01$ vs. 250 μM H_2O_2, Student's t-test). Sharps show significant difference from the response by $2MeSADP/H_2O_2$ (##$P < 0.01$ vs. $2MeSADP/H_2O_2$). (2) The effect of MRS2179, auranofin, and NAC on H_2O_2-evoked ERK1/2 activation. MRS2179 (10 μM) and auranofin (1 μM) reversed the ERK1/2 activity inhibited by 2MeSADP. NAC (10 mM) prevented the H_2O_2-evoked ERK1/2 activation. Asterisks show significant difference from control (**$P < 0.01$ vs. control, Student's t-test). Sharps show significant difference from H_2O_2 (##$P < 0.01$ vs. H_2O_2, Student's t-test). Dollar marks show significant difference from $2MeSADP/H_2O_2$ ($$P < 0.05$ vs. $2MeSADP/H_2O_2$, Student's Results were expressed as means t-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$).

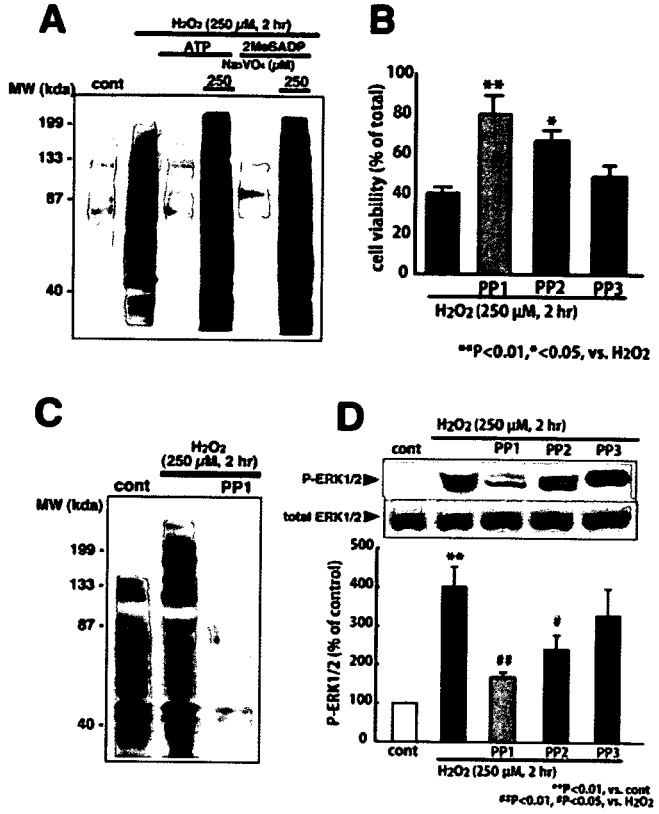

Fig. 8. The effects of the src family on the H_2O_2-evoked cell death and ERK1/2 activation. **A**: ATP and 2MeSADP prevented the H_2O_2-evoked protein tyrosine phosphorylation via PTP upregulation. H_2O_2 (250 μM) evoked the protein tyrosine phosphorylation. Na_3VO_4 (250 μM) markedly inhibited the ATP-(100 μM, 24 h) and 2MeSADP (1 μM, 24 h)-induced prevention of the H_2O_2-evoked protein tyrosine phosphorylation. Na_3VO_4 was applied to the cells 1 h before and during the H_2O_2 treatment. **B**: The effect of selective src family inhibitors on the H_2O_2-evoked cell death. PP1 and PP2 (250 nM) prevented the H_2O_2-evoked cell death but PP3 (250 nM) did not. Asterisks show significant difference from the response evoked by H_2O_2 ($^*P < 0.05$, $^{**}P < 0.01$ vs. 250 μM H_2O_2, Student's *t*-test) . **C**: The effect of PP1 on H_2O_2-evoked protein tyrosine phosphorylation. PP1 (250 nM) strongly inhibited the H_2O_2-evoked protein tyrosine phosphorylation. **D**: The effect of selective src family inhibitors on H_2O_2-evoked ERK1/2 activation. PP1 and PP2 (250 nM) inhibited H_2O_2-evoked ERK1/2 activation but PP3 (250 nM) did not. PP1, PP2, and PP3 were applied to the cells 1 h before and during the H_2O_2 treatment. Asterisks show significant difference from control ($^{**}P < 0.01$ vs. control, Student's *t*-test). Sharps show significant difference from H_2O_2 ($^{\#\#}P < 0.01$ vs. H_2O_2, Student's *t*-test). Results were expressed as means ± SEM of triplicate measurements ($n = 3$).

2MeSADP-induced prevention disappeared when the astrocytes were treated with Na_3VO_4 to inhibit the PTP activity, suggesting that ATP and 2MeSADP would inhibit tyrosine phosphorylation through a pathway(s) mediated by PTP. As it has been reported that H_2O_2 especially activates the src tyrosine kinase family of PTK (Lee and Esselman, 2002; Nishida et al., 2000), we pharmacologically studied whether the src family participates in the H_2O_2-evoked cell death and ERK1/2 activation. When the selective src family inhibitors PP1 (250 nM) and PP2 (250 nM) (Hanke et al., 1996) were added to the cells 1 h before H_2O_2 treatment, the H_2O_2-evoked cell death in astrocytes was abolished (Fig. 8B). The inactive analogue PP3 (250 nM) did not inhibit the H_2O_2-evoked cell death. PP1 (250 nM) also inhibited the H_2O_2-evoked protein tyrosine phosphorylation (Fig. 8C). In addition, the H_2O_2-evoked ERK1/2 activation was inhibited by PP1 and PP2 (250 nM) but not by PP3 (Fig. 8D). PP1, PP2, and PP3 were added to the cells 1 h before H_2O_2 stimulation.

DISCUSSION

In the present study, we demonstrated that ERK1/2 and src family are important molecules that promote the H_2O_2-evoked astrocytic cell death, and that ATP upregulates PTP expression and its activity, thereby preventing the H_2O_2-evoked src family and following ERK1/2 activation, resulting in the protection of astrocytes against H_2O_2-evoked cell death.

We clearly showed that the H_2O_2-evoked activation of ERK1/2 and accumulation of P-ERK in nuclei were critical events that promote cell death in astrocytes. ATP itself, however, which exhibited a protective effect against H_2O_2 in our study, is also known to activate ERK1/2 in astrocytes (Neary et al., 1999, 2003; Panenka et al., 2001). In fact, ATP and 2MeSADP activated ERK1/2 in astrocytes. However, the ERK1/2 activation by ATP, in contrast to that by H_2O_2, was transient (5–15 min after stimulation) and did not affect the astrocyte cell viability. Furthermore, 2MeSADP-activated ERK1/2 seems to function rather as an essential signal that prevents cell death and induces the upregulation of oxidoreductases and PTP gene expression. Thus, such a discrepancy appears to result from spatio- and temporal-behavioral differences of P-ERK1/2. It is known that, after faint brain ischemia, neuronal cells acquire tolerance to a subsequent more serious ischemic injury (Chen and Simon, 1997; Dawson and Dawson, 2000; Schaller and Graf, 2002). The similarity between such preconditioning against ischemia and the preconditioning of ERK1/2 against H_2O_2-evoked cell death is very interesting.

In the present study, the extent of the ERK1/2 activity appeared to be important in the H_2O_2-evoked cell death because of the correspondence between the concentration-dependency of the H_2O_2-evoked ERK1/2 activity and that of the H_2O_2-evoked cell death (Shinozaki et al., 2005). Although recent studies also have reported that H_2O_2-activated ERK1/2 evokes cell death in glioma and osteoblastic cells (Choi et al., 2005; Levinthal and Defranco, 2005), the spatio- and temporal behavior of P-ERK1/2 remained unclear. P-ERK1/2 is known to translocate into the nucleus (Rosenberger et al., 2001) and accumulate there (Brand et al., 2001; Stanciu and DeFranco, 2002), thereby inducing neuronal death. In the immunocytochemical analysis, stimulation by 250 μM but not by 50 μM H_2O_2 for 2 h induced P-ERK1/2 accumulation in the nucleus. Most of the total ERK1/2 existed in the cytoplasm and was not affected by H_2O_2 (50 or 250 μM). Accordingly, it is conceivable that only P-ERK1/2, activated by high concentrations (i.e. 250 μM) of H_2O_2, accumulates in the nucleus and induces cell

death. In a spatiotemporal analysis of P-ERK1/2, although there was a time-dependent activation of ERK1/2 by 50 and 250 μM H_2O_2 at the whole cell level, the fraction of N = E was increased transiently (~15 min after stimulation) by 50 μM H_2O_2 but time-dependently by 250 μM. These results suggest that the long-term accumulation of P-ERK1/2 into the nucleus participates in the H_2O_2-evoked cell death.

In physiological conditions, it is suggested that the P-ERK1/2 translocated into the nucleus is dephosphorylated by MAPK phosphatase (MAPKP), especially by the ERK1/2 selective phosphatase MAPK phosphatase-3 (MKP-3) (Dowd et al., 1998; Groom et al., 1996) and is exported from the nucleus depending on MKP-3 (Karlsson et al., 2004). The interaction between MKP-3 and ERK1/2 requires arginine residues of MKP-3 (Nichols et al., 2000). Because many amino acids, including arginines, in protein are oxidized by H_2O_2 (Amici et al., 1989; Moskovitz et al., 2002; Stadtman and Berlett, 1997; Taborsky, 1973), the interaction of MKP-3 and ERK1/2 and the nuclear export of ERK1/2 may be affected by H_2O_2. Furthermore, the ERK1/2 inactivating enzyme MAPKP is inactivated by H_2O_2 (Foley et al., 2004; Levinthal and Defranco, 2005). In conditions with oxidative stress, it is conceivable that the dephosphorylation and nuclear export of activated-ERK1/2 are attenuated because of the decreased MAPKP function and the association between MAPKP and ERK1/2, which thereby induces the prolonged activation and nuclear accumulation of ERK1/2.

With regard to the upstream molecule that activates ERK1/2 in response to H_2O_2, we found that the src family is important. Furthermore, we demonstrated that activation of $P2Y_1$ receptors inhibits the activation of src family and subsequent signaling cascades by upregulating the PTP expression and activity. We previously reported that ATP upregulates the thiol-containing protein TrxR (Shinozaki et al., 2005). The enzymatic activity of PTP requires reduction of the cystein residue in its active center (Cho et al., 2004; Persson et al., 2004). Accordingly, the redox state of the cystein residue in PTP is considered to crucially affect its phosphatase activity. The decreased activity of PTP in an oxidative state is recovered by adding thiol-containing protein such as glutathione (Salmeen et al., 2003) and thioredoxin (Lee and Esselman, 2002). In fact, the $P2Y_1$ receptor activation-induced restoration of the PTP activity was reversed by auranofin, indicating that TrxR restores the PTP activity. Additionally, the antioxidant NAC restored the PTP activity decreased by H_2O_2 (250 μM). The protective effect of PTP induced by ATP requires either an increase in the amount/activity of PTP or a reduction, in which the upregulated TxrR would have a critical role (Shinozaki et al., 2005). Furthermore, such ATP-induced oxidoreductases may preserve the MKP-3 activity, thereby enhancing the dephosphorylation and inactivation of ERK1/2 and the nuclear export of ERK1/2.

PTP and PTK regulate protein tyrosine phosphorylation in close coordination with each other. ATP prevented the H_2O_2-induced protein tyrosine phosphorylation by increasing the PTP activity. The inhibition of the H_2O_2-evoked cell death by PP1 and PP2 and the tyrosine phosphorylation by PP1 indicates that the src tyrosine kinase family is related to the H_2O_2-evoked cell death. Additionally, PP1 and PP2 inhibited the H_2O_2-evoked ERK1/2 activation, indicating that ERK1/2 is activated following src family activation. As the src family is activated by cystein oxidation independently of tyrosine phosphorylation (Akhand et al., 1999; Pu et al., 1996), under oxidative stress such as by H_2O_2 treatment or in condition of ischemia/reperfusion-injury, the src family could be activated independent of tyrosine phosphorylation. In contrast to src activation, PTP and MAPKP are inactivated by oxidation of the cystein residue under oxidative conditions (Cho et al., 2004; Foley et al., 2004; Meng et al., 2004). Briefly, the "irresponsive to control" signal transduction could be caused under oxidative conditions.

In conclusion, we clearly demonstrated that the src family activation followed by strong ERK1/2 activation and prolonged P-ERK1/2 accumulation into the nucleus participated in the H_2O_2-evoked cell death of astrocytes. ATP/$P2Y_1$ receptor activation inhibits H_2O_2-evoked ERK1/2 activation by preventing the H_2O_2-evoked src activation via upregulation of PTP expression/activity. Our present findings suggest that the gliotransmitter ATP protects astrocytes against oxidative stress by counteracting the intracellular signaling pathway that evokes cell death.

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Microglial α7 Nicotinic Acetylcholine Receptors Drive a Phospholipase C/IP_3 Pathway and Modulate the Cell Activation Toward a Neuroprotective Role

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Microglia perform both neuroprotective and neurotoxic functions in the brain, with this depending on their state of activation and their release of mediators. Upon $P2X_7$ receptor stimulation, for example, microglia release small amounts of TNF, which protect neurons, whereas LPS causes massive TNF release leading to neuroinflammation. Here we report that, in rat primary cultured microglia, nicotine enhances $P2X_7$ receptor-mediated TNF release, whilst suppressing LPS-induced TNF release but without affecting TNF mRNA expression via activation of α7 nicotinic acetylcholine receptors (α7 nAChRs). In microglia, nicotine elicited a transient increase in intracellular Ca^{2+} levels, which was abolished by specific blockers of α7 nAChRs. However, this response was independent of extracellular Ca^{2+} and blocked by U73122, an inhibitor of phospholipase C (PLC), and xestospongin C, a blocker of the IP_3 receptor. Repeated experiments showed that currents were not detected in nicotine-stimulated microglia. Moreover, nicotine modulation of LPS-induced TNF release was also blocked by xestospongin C. Upon LPS stimulation, inhibition of TNF release by nicotine was associated with the suppression of JNK and p38 MAP kinase activation, which regulate the post-transcriptional steps of TNF synthesis. In contrast, nicotine did not alter any MAP kinase activation, but enhanced Ca^{2+} response in $P2X_7$ receptor-activated microglia. In conclusion, microglial α7 nAChRs might drive a signaling process involving the activation of PLC and Ca^{2+} release from intracellular Ca^{2+} stores, rather than function as conventional ion channels. This novel α7 nAChR signal may be involved in the nicotine modification of microglia activation towards a neuroprotective role by suppressing the inflammatory state and strengthening the protective function.

Key words: microglia; nicotinic acethylcholine receptor; ATP; lipopolysaccharide; tumor necrosis factor

Nicotinic acetylcholine receptors (nAChRs) contribute to brain functions such as the modulation of synaptic transmission, memory formation, and neuroprotection (MacxDermot et al., 1999; Cordero-Erausquin et al., 2000; Laudenbach et al., 2002; Daijas-Bailador and Wonnacott, 2004). Among nAChRs, α7 and α4β2 are the most abundant subunits in the brain (Buisson and Bertrand, 2002). The α7 nAChRs are ion channels forming homo-pentamers with high Ca^{2+} permeability, which exhibit a low affinity for acetylcholine (ACh) and nicotine and are desensitized rapidly. In contrast, α4β2 nAChRs are considered to exhibit a highly affinity for ACh and nicotine and are desensitized slowly (Cordero-Erausquin et al., 2000). Although these nAChRs have been regarded to be expressed mainly in neuronal cells, recent evidence suggests that the α7 nAChRs also function in nonexcitable cells, such as endothelial cells, keratinocytes, and T cells (Sharma and Vijayaraghavan, 2002).

Microglia are the primary immune cells in the central nervous system (CNS). Under pathological conditions such as brain damage and stroke, they are rapidly activated, phagocytose dead cells, and secrete various cytokines, including tumor necrosis factor (TNF), interleukin-1β

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(IL-1β), and IL-6 (Aliosi, 2001; Shigemoto-Mogami et al., 2002). TNF, a proinflammatory and cytotrophic cytokine up-regulated in the brain in response to various insults or injury (Barone et al., 1997; Shohami et al., 1999), is released mainly by microglia and astrocytes around the injured area. Lipopolysaccharide (LPS), one of the main components of gram (−) bacterial outer membranes, induces the release of large amounts of TNF from macrophages and microglia, leading to inflammation and possibly neuronal destruction (Kim et al., 2000). Microglia also release moderate amounts of TNF in response to extracellular ATP via activation of the ionotrophic $P2X_7$ receptor (Hide et al., 2000; Suzuki et al., 2004). This manner of TNF release protects neurons against glutamate cytotoxicity (Suzuki et al., 2004).

Recently, it was revealed that α7 nAChRs are expressed in mouse brain microglia and are involved in the suppression of neuroinflammation (Shytle et al., 2004). In this study, we found that, in rat primary cultured microglia, the activation of α7 nAChRs by nicotine enhanced protective TNF release induced by ATP stimulation, in addition to having a suppressive effect on LPS-induced TNF release via α7 nAChRs. Furthermore, the present results demonstrate that these microglial α7 nAChRs have different properties from conventional neuronal α7 nAChRs, not functioning as ion channels but coupling to phospholipase C (PLC) activation and Ca^{2+} mobilization from IP_3-sensitive Ca^{2+} stores. We also investigated the signaling mechanism by which α7 nAChR activation could modulate TNF release in response to LPS (TLR4 activation) and ATP or BzATP ($P2X_7$ receptor activation).

MATERIALS AND METHODS

Reagents

Reagents were obtained from the following sources: all reagents for cell culture, (−)nicotine hydrogen tartrate, LPS, ATP, 2′- and 3′-O-(benzoyl-benzoyl) adenosine 5′-triphosphate (BzATP), α-bungarotoxin, methyllycaconitine, and xestospongin C were from Sigma Chemical Co. (St. Louis, MO); U73122 from Calbiochem (San Diego, CA); BAPTA-AM from WAKO Pure Chemical Industries (Osaka, Japan); rat TNF ELISA kit from Biosource International (Camarillo, CA); polyclonal antibody against α7 nAChRs from Santa Cruz Biotechnology (Santa Cruz, CA); antibody kits for p42/p44 (ERK), JNK, and p38 MAP kinase from Cell Signaling Technology (Beverly, MA). All other reagents were purchased from commercial sources and were of the highest available purity.

Cell Culture

Microglia were obtained from the primary cell cultures of neonatal rat brains as previously described (Nakajima et al., 1992). After 7–16 days in culture, microglia were prepared as a floating cell suspension. Aliquots ($1.5–2.0 \times 10^5$ cells) were transferred to the wells of a 24-well plate and allowed to adhere at 37°C for 45 min. Unattached cells were removed by rinsing with serum-free Dulbecco's modified Eagle's medium (DMEM).

TNF Assay

Microglia were incubated with 0.4 ml of serum-free DMEM with or without drugs for 3 hr. TNF was assayed in 50-μl samples by using a rat TNF ELISA kit according to the manufacturer's instructions.

Isolation of Total RNA, RT-PCR, and Real-Time Quantitative RT-PCR

Total RNA was isolated from microglia with Trizol reagent (Life Technologies, Inc.) according to the manufacturer's protocol. The amounts of tRNA was determined by using absorption of light at 260 and 280 nm.

The expression of mRNA of α7 nAChR subunit was detected by nested RT-PCR in rat microglia. RT-PCR was performed with an Access RT-PCR system (Promega, Madison, WI). Two sets of primers used to identify α7 were directed toward portions of exons 9–10: the first sets, forward, 5′-TCATGCTGCTTGTGGCTGAG-3′ and reverse sets, 5′-CCAATTCTCACCCTCCAGATTCTC-3′, and the second sets, forward, 5′-GCAACATCTGATTCTGTGCCCTTG-3′ and reverse, 5′-TCTGCGCATTTCCTACTTGA-3′. Thermocycling conditions were as follows: 95°C for 30 min; 45 cycles of 95°C for 15 sec, 60°C for 60 sec; PCR was performed in a tube with a total volume of 50 μl of reverse transcriptase mixture containing 1 mM $MgSO_4$, 0.2 mM dNTP, 0.1 U/μl Tfl DNA polymerase, 1 U/μl AMV RT.

The expression levels of mRNA for TNF were measured by real-time quantitative RT-PCR (ABI Prism model 7700 sequence detection system; PE Applied Biosystems, Foster City, CA) as described previously (Suzuki et al., 2004). RT-PCR was carried out by using TaqMan one-step RT-PCR Maser Mix reagents kit according to the manufacturer's protocol (Applied Biosystems). The sequences of the forward and reverse primers were 5′-ACAAGGCTGCCCCGACTAC-3′ and 5′-TCCTGGTATGAAATGGCAAACC-3′ respectively. The TaqMan fluorogenic probe was 5′-6FAM-TGCTCCTCACCCACACCGTCAGC-TAMIRA-3′. During PCR amplification, 5′ nucleotidase activity of AmpliTaq Gold DNA polymerase cleaves the TaqMan probe separating the 5′ reporter dye from the 3′ quencher dye, resulting in increased fluorescence of the reporter. The threshold cycle, C_T, which correlates inversely with the target mRNA levels, was measured as the cycle number at which reporter fluorescent emission increases above a threshold level. The TNF mRNA levels were corrected for the C_T values of glyceraldehydes-3-phosphate dehydrogenase (GAPDH) RNA by using VIC probe according to the manufacturer's protocol.

Measurement of Intracellular Ca^{2+} Concentration in Microglia

Microglia were plated on glass coverslips with a silicon rubber wall (Flexiperm; Heraeus Biotechnology, Hanau, Germany) at a density of 3×10^5 cells/well and maintained in 10% CO_2. After rinsing, the cells were loaded with 2.5 μM fura-2 acetoxymethyl ester for 20 min at room temperature. Fura-2-loaded cells were placed on a fluorescence-image microscope and stimulated by nicotine. The fura-2 fluorescence was measured with excitation at 340 and 380 nm and at

an emission wavelength of 510 nm. The video image output was digitized by an Argus 50 color image processor (Hamamatsu Photonics, Shizuoka, Japan) as described elsewhere (Hide et al., 1997).

Western Blot Analysis

Western blots were performed for the analysis of α7 nAChRs expression and for the analysis of ERK (p44/42), JNK, and p38 activation using each MAP kinase antibody. In brief, the cells were washed with phosphate-buffered saline, lysed by adding sodium dodecyl sulfate (SDS) sample buffer, and sonicated. After heating to 95°C for 5 min, the protein samples were separated by SDS-polyacrylamide gel electrophoresis (PAGE) and blotted onto polyvinylimide difluoride (PVDF) membranes. The membranes were blocked with blocking buffer containing 5% skim milk for 3 hr at room temperature and incubated with primary antibody with gentle agitation overnight at 4°C. After washing, the membranes were incubated for 1 hr at room temperature with horseradish peroxidase-conjugated secondary antibody and horseradish peroxidase-conjugated anti-biotin antibody to detect biotinylated protein markers. The membranes were then washed and incubated with Lumi GLO, and the proteins were detected by exposure to X-ray films.

Whole-Cell Patch Clamp

We performed whole-cell patch clamp for microglia according to the methods previously described (Matsubayashi et al., 2004). Microglia were transferred into a recording chamber, and the chamber was perfused continuously with external bath solution at 2 ml/min. The ionic composition of the external solution was as follows (in mM): NaCl, 165; KCl, 5; glucose, 10; $CaCl_2$, 2; HEPES, 5; and the pH of the solution was adjusted to 7.3 with NaOH. Experiments were carried out at room temperature. In voltage-clamp modes, drug-induced currents were recorded from microglia according to the standard whole-cell patch clamp technique by using an Axopatch 200A patch clamp system (Axon Instruments, Foster City, CA). Patch pipettes were pulled from borosilicate capillary glass (World Precision Instrument-Japan, o.d. 1.5 mm) and had resistances between 3 and 7 MΩ when filled with an internal solution with the following ionic composition (in mM): CsCl, 80; CsF, 80; $MgCl_2$, 2; HEPES, 10; Cs-EGTA, 10 (pH adjusted to 7.3 with CsOH). The series resistance of the patches was 10–25 MΩ and was not compensated. The signals of currents were filtered at 2 kHz and directly sampled by a microcomputer using the program pCLAMP (Axon Instruments). The drugs were delivered to the microglia from a pore 400 μm in diameter at the apex of a U-shaped tube ("U-tube") positioned about 50 μm directly above the target microglia. With this drug application system, the drug solution was ejected from the pore when the solenoid valve was closed electronically, and then the solution surrounding target microglia was quickly exchanged to drug solution at the desired concentrations. The whole-cell currents were recorded at −60 mV of holding potential.

Statistical Analysis

Statistical analysis of data was performed via one-way ANOVA and Dunnett's test.

RESULTS

Nicotine Inhibits LPS-Induced TNF Release but Enhances ATP- or BzATP-Induced TNF Release From Microglia

LPS endotoxin derived from gram-negative bacteria induces a large amount of TNF release via toll-like receptor 4 (TLR4) in microglia and macrophages. Such unregulated TNF release should cause inflammation and neuronal destruction. On the other hand, extracellular ATP stimulates the release of TNF from rat microglia via activating $P2X_7$ receptors (Hide et al., 2000), and this TNF release seems to be well regulated and to be neuroprotective (Suzuki et al., 2004). To investigate whether nicotine could modulate such microglial cell function, we examined the effects of nicotine on TNF release from microglia activated by LPS (1 ng/ml), ATP (1 mM), or the $P2X_7$ receptor agonist BzATP (100 μM). Nicotine significantly inhibited LPS-induced TNF release in a concentration-dependent manner (Fig. 1A), confirming a previous report (Shytle et al., 2004). Unlike LPS stimulation, however, nicotine significantly enhanced the release of TNF in ATP- or BzATP-stimulated microglia (Fig. 1B,C). These results indicate that nicotine might control microglial cell function toward neuroprotective by suppressing LPS-stimulated massive TNF release and enhancing ATP-induced protective TNF release.

Nicotine Modulates TNF Release Via Activation of α7 nAChRs

To clarify which subtype of nAChRs is responsible for modulation of TNF release in microglia, we examined the effects of methyllycaconitine (MLA), an α7 nAChR blocker, on nicotine-induced modulation of TNF release. MLA (10 nM) reversed these bidirectional actions of nicotine, namely, the inhibition of LPS-induced TNF release and the increase of ATP-stimulated TNF release (Fig. 2A,B). MLA did not affect LPS- or ATP-induced TNF release by itself. These results suggest that nicotine modulates TNF release at least through activation of α7 nAChRs. The expression of α7 nAChRs was recently shown in mouse microglia (Shytle et al., 2004). In this study, we also confirmed the expression of α7 nAChR protein by Western blotting and their mRNA expression by nested RT-PCR in primary rat brain microglia (Fig. 2C,D).

Microglial α7 nAChRs Do Not Function as Ligand-Gated Ion Channels but Induce Ca^{2+} Release From Intracellular Ca^{2+} Stores Through PLC and IP_3 Pathway

It is reported that neuronal α7 nAChRs are ligand-gated ion channels with high permeablity to Ca^{2+}, and the activation of these receptors by nicotine induces Ca^{2+} influx (Albuquerque et al., 1995). In rat microglia, nico-

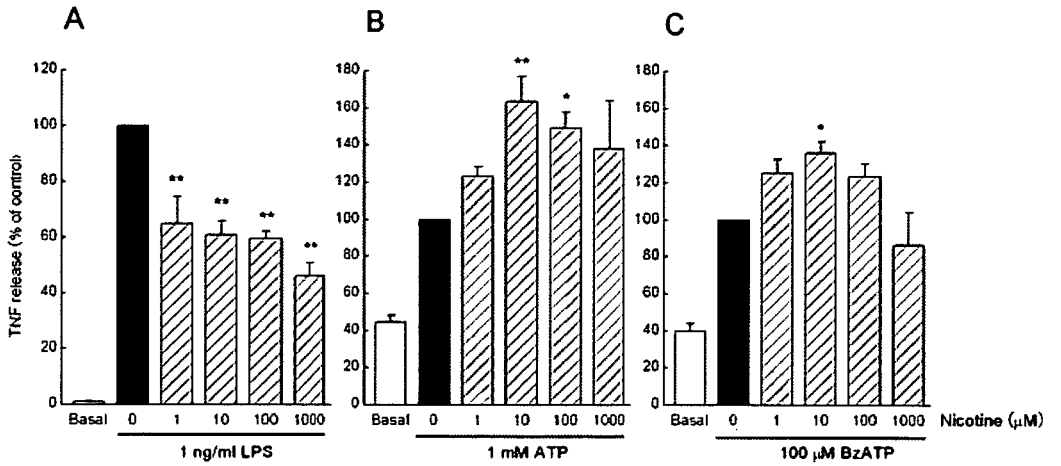

Fig. 1. Effects of nicotine on LPS-, ATP-, or BzATP-stimulated TNF release in microglia. The cells were treated with nicotine for 10 min and stimulated with 1 ng/ml LPS (**A**), 1 mM ATP (**B**), or 100 μM BzATP (**C**) for 3 hr. The released TNF was measured by ELISA. Values are expressed as mean ± SEM of percentage release compared with LPS or ATP/BzATP alone from three independent experiments. Values of 100% for the release of TNF in LPS-stimulated, ATP-stimulated, and BzATP-stimulated microglia were 11.7 ± 3.7 ng/10^6 cells, 0.42 ± 0.03 ng/10^6 cells, and 0.49 ± 0.09 ng/10^6 cells, respectively. $^*P < 0.05$, $^{**}P < 0.01$ vs. control (ANOVA with Dunnett's posttest).

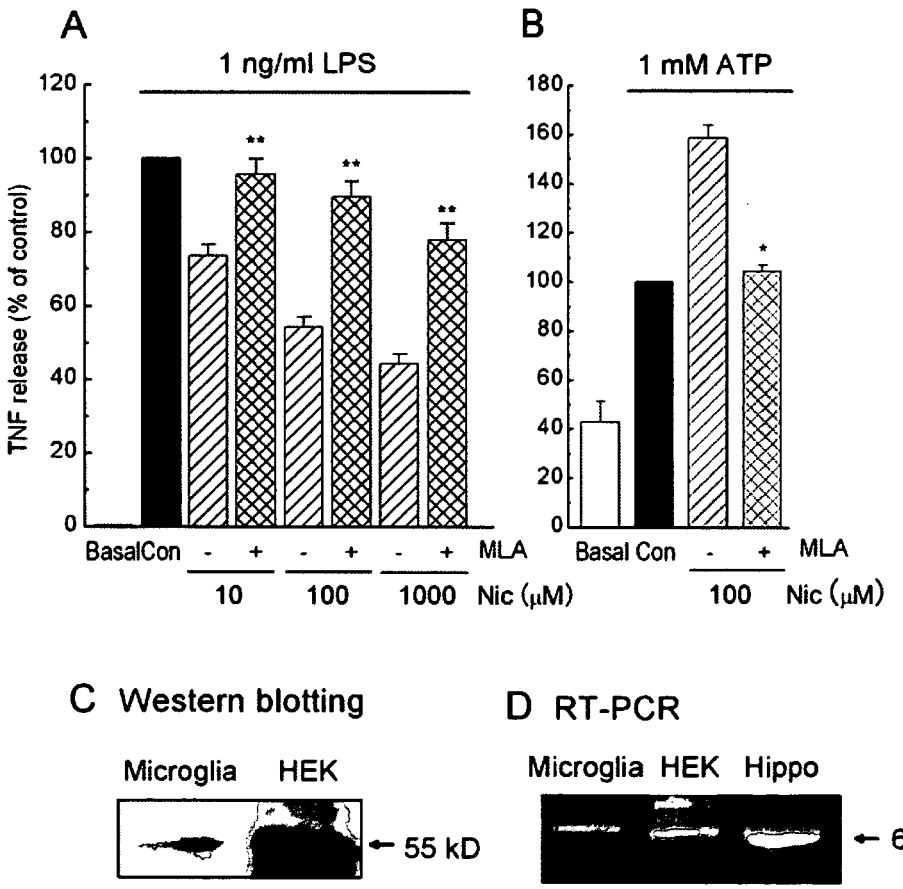

Fig. 2. The nicotine modulations of TNF release are mediated via α7 nAChRs expressed in microglia. **A,B:** Effects of methyllycaconitine (MLA), a specific antagonist of α7 nAChRs, on LPS- or ATP-induced TNF release from nicotine-treated microglia. The cells were treated with 10 nM MLA for 10 min before addition of nicotine, then stimulated with 1 ng/ml LPS (A) or 1 mM ATP (B) for 3 hr. The released TNF were measured by ELISA. Values are expressed as mean ± SEM of percentage of release compared with LPS alone from three independent experiments. Values of 100% for the release of TNF in LPS- and ATP-stimulated microglia were 20.5 ± 1.9 ng/10^6 cells (A) and 466.5 ± 15.6 pg/10^6 cells (B), respectively. $^*P < 0.05$, $^{**}P < 0.01$ compared with each nicotine-treated control (ANOVA with Dunnett's posttest). **C,D:** Protein and mRNA expression of α7 nAChRs in primary rat brain microglia. C: Western blotting. Cell lysates were analyzed with specific antibody against α7 nAChR. HEK 293 cells overexpressing α7 nAChRs were used as a positive control. D: Nested RT-PCR with primers specific for the α7 subunit as described in Materials and Methods.

tine induced a rapid and transient increase in the concentration of cytosolic Ca^{2+} (Fig. 3A). This Ca^{2+} response was strongly blocked by 10 nM MLA, an α7 nAChR blocker (Fig. 3B), showing that microglial α7 nAChRs are functional. Another selective α7 nAChR antagonist, α-bungarotoxin (10 nM), also blocked this response (Fig. 3C). Unexpectedly, however, the nicotine-elicited transient increase in intracellular Ca^{2+} concentration was not affected even in the absence of extracellular Ca^{2+} (Fig. 3D). Furthermore, nicotine-induced Ca^{2+} response