



# Basic fibroblast growth factor induces miR-134 upregulation in astrocyte for cell maturation



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## ABSTRACT

Evidence suggests that neuronal microRNAs (miRs) contribute to synaptic plasticity, although a role of glial miRs have been unknown. Growth factors including brain-derived neurotrophic factor (BDNF) regulate neuronal functions via upregulation of miRs, while possible influences on expression/function of glial miRs have not been fully understood. Here, we report that basic fibroblast growth factor (bFGF) increased miR-134 expression in astrocyte. The miR-134 was upregulated through stimulating extracellular signal-regulated kinase and phosphatidylinositol 3-kinase signaling, because inhibitors for each signaling blocked the miR-134 induction by bFGF. We also found upregulation of glial fibrillary acidic protein (astrocyte marker) and decreased extracellular concentration of glutamate after miR-134 overexpression and bFGF application, suggesting that astroglial cell maturation is enhanced by bFGF through induction of miR-134.

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## 1. Introduction

MicroRNAs (miRs), a highly conserved noncoding small RNAs, have been recognized to be a regulator for various physiological aspects including neural development via inhibiting the function of mRNA of target genes [1]. Both negative (e.g. miR-34a, -134, -137, and -219) and positive (e.g. miR-124, -132, and -188) functions of neural miRs in the neuronal maturation and synaptic plasticity have been demonstrated [2]. Recently, it has been suggested that the dysregulation of miR-132 is involved in the abnormality of neurodevelopment and neuro-morphology in schizophrenia [3]. In addition, interfering of miR-134 function in mice by antagomir leads to reduction of evoked seizures and neuroprotection [4], suggesting a possible therapeutic role of miRs in brain diseases.

Studies have demonstrated that the miR expression is regulated by growth factors in neurons [5–7]. For example, BDNF/TrkB (brain-derived neurotrophic factor and its high affinity receptor) system upregulates expression levels of miR-212/132 through activating MSK (mitogen- and stress-activated kinase) 1 and CREB

(cAMP-response element binding protein) pathways [5]. Glial cell line-derived neurotrophic factor (GDNF) also regulates miRs (e.g. miR-188-5P, miR-434-3P, and miR-340-3P) expression, and protects dopaminergic cell line against cell death induction caused by 6-hydroxydopamine [7]. Previously, we showed that basic fibroblast growth factor (bFGF) enhanced miR-132 expression not only in cortical neurons but also in astrocytes, while BDNF-dependent miR-132 induction was observed in only neurons and plays a role for synaptic protein expression [8]. On the other hand, miR-125b [9] and miR-155 [10] have been identified as astroglial miRs. However, function of glial miRs induced by growth factors has been poorly understood. Especially, though bFGF is a well-known regulator for cell differentiation and survival in neurons [11], glial miR induced by bFGF and its physiological role has not been clarified.

MiR-134, one of brain-specific miRs, localizes in dendrites, and regulates dendritic spine size via suppressing levels of Limk1 (Lim-domain-containing protein kinase 1) mRNA in hippocampal neurons [12]. MiR-134 has a role in neuronal development such as proliferation of neural precursor cell and embryonic neuronal maturation [13]. Increased miR-134 is important for the dendritic maturation induced by BDNF through regulating myocyte enhancing factor 2 (Mef2), a transcription factor [14]. These reports show functions of neuronal miR-134 and its interaction with growth factors including BDNF.

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In the present study, we found that significant miR-134 upregulation after bFGF stimulation in astrocytes, and that overexpression of miR-134 increased expression levels of astroglial marker proteins and glutamate transporter, suggesting a novel role of miR-134 in astroglial maturation and glial function.

## 2. Materials and methods

### 2.1. Antibodies and materials

Anti-phosphorylated or total extracellular signal-regulated kinase1/2 (ERK1/2), and anti-phosphorylated or total Akt antibodies were purchased from cell signaling technology Inc. (Danvers, MA, USA). U0126 and LY294002 were also from the cell signaling technology Inc. Both anti-S-100 and anti-aldehyde dehydrogenase 1 family, member L1 (ALDH1L1) antibodies were obtained from Abcam plc (Cambridge, UK). Anti-glial glutamate transporter GLT-1 (EAAT2) antibody was purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Anti-glial fibrillary acidic protein (GFAP) and anti- $\beta$ -actin antibodies were obtained from Chemicon International, Inc. (Temecula, CA, USA) and Sigma-Aldrich (St. Louis, MO, USA), respectively. Taqman<sup>®</sup> microRNA assays (miR-9: 000583, miR-124a: 001182, miR-132: 000457, miR-134: 001186, and miR-16: 000391), Taqman<sup>®</sup> microRNA reverse transcription kit, epidermal growth factor (EGF), Alexa Fluor<sup>®</sup> 488 goat anti-rabbit IgG, Hoechst 33342, Dulbecco's modified Eagle medium: Nutrient Mixture F-12 (DMEM/F12), and minimum essential medium (MEM) from Life technologies corporation (Carlsbad, CA, USA) were used in this study. Takeda Pharmaceutical Company Ltd., and Sumitomo Co. Ltd. kindly donated BDNF. bFGF was purchased from PeproTech (Rocky Hill, NJ, USA). Insulin-like growth factor-1 (IGF-1) and GDNF were purchased from R&D systems (Minneapolis, MN, USA) and Wako Pure Chemical Industries Ltd. (Osaka, Japan), respectively.

### 2.2. Cell cultures

Primary cortical neurons and astrocytes were prepared from cerebral cortex of Wistar rats at postnatal day 1–2 as described previously [8]. Neural cortical cells were cultured with DMEM/F12 containing heated-inactivated 5% horse serum and 5% fetal bovine serum (FBS), and 2  $\mu$ M cytarabine to prevent glial cell survival. At 4–5 days *in vitro* (DIV), the neuronal cultures were treated with 100 ng/ml BDNF or bFGF for 24 h. Astrocyte pure cultures were maintained with MEM-based growth medium supplemented with 100  $\mu$ g/l EGF, 20 mM glucose, 25 mM NaHCO<sub>3</sub>, 5% FBS, and 0.5 mM glutamine. When the confluency of the astrocytes was reached to 70–80%, culture medium was replaced with fresh one without EGF for 48 h before stimulation by growth factors. The pre-miR-134 (CAGGGUGUGUGACUGGUUGACCAGAGGGGCGUGCA CUUUGUUCACCCUGUGGGCCACCUAGUCACCAACCCUC) with 5'- and 3'-flanking region was inserted to BamH I/Hind III site of pBApo-CMV Neo vector (TAKARA BIO Inc., Shiga, Japan). Transfection of 3  $\mu$ g of the miR-134 expression vector into astrocytes was conducted with Lipofectamine 2000 (Life Technologies) according to manufacturer's protocol. After additional 48 h maintenance, transfected cells were used for experiments. We confirmed that the transfection efficiency was approximately 77.4  $\pm$  8.7% (the ratio: the number of GFP-positive cells to the number of total nuclear stained with Hoechst 33342) using GFP construct (pAc-GFP1-N1; Clontech laboratories, Inc., Mountain View, CA, USA).

### 2.3. The measurement of miRs expression

Extraction of total RNA including miRs from cultures was performed with mirVana<sup>™</sup> miRNA isolation kits (Ambion, Austin,

TX, USA) according to manufacturer's instruction. The concentration of total RNA was measured (as optical density, OD, at 260 nm) by using ND-1000 spectrophotometer (Nanodrop, Wilmington, DE, USA), and quality of RNA was determined with the ratio value of OD260/OD280 (>1.8). Then, 100 ng of total RNA were applied to reverse-transcription with Taqman<sup>®</sup> primer and Taqman<sup>®</sup> microRNA reverse transcription kit. The quantification of miR concentration in extracted samples was carried out by real-time PCR using specific Taqman<sup>®</sup> probe with 7900HT fast real-time PCR System (Life Technologies Corporation). The data was analyzed by SDS 2.2 real-time PCR data analysis software (Life Technologies Corporation). In this study, the each concentration value was normalized by levels of miR-16 in the same sample.

### 2.4. Western blotting

Total lysate from astroglial cultures were collected with lysis buffer which composed of 1% SDS, 10 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub>, 10 mM Tris-HCl, 10 mM NaF, 5 mM EDTA, 2 mM NaVO<sub>4</sub> and 1 mM PMSF. The protein concentration in each sample was measured with Pierce<sup>®</sup> BCA protein assay (Thermo scientific, IL, USA), before equivalent amounts of total proteins were applied to each electrophoresis. Separated proteins on acryl-amide gel were blotted to PVDF membrane (Millipore, MA, USA), then, incubation with primary antibody after blocking with 5% skim milk diluted in TBS {0.1 M Tris-HCl (pH 7.4) in saline} was performed. The mouse- (Jackson ImmunoResearch Europe Ltd., Suffolk, UK) or rabbit- (Rockland Immunochemicals, Inc., Gilbertsville, PA, USA) secondary antibody was used after several times of washing with TBS. The immunoreactivity visualized by Immunostar<sup>®</sup> reagents (Wako Pure Chemical Industries Ltd., Osaka, Japan) and ECL films (GE healthcare UK Ltd., Buckinghamshire, UK) was obtained. The optical density of each protein expression was determined by CS Analyzer Version 3.00.1011 (ATTO Co., Tokyo, Japan).

### 2.5. Cell survival

MTT {3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide, yellow tetrazolium salt} assay was performed as previously reported [15]. Briefly, cultured medium was exchanged to fresh DMEM/F12 containing 0.5 mg/ml MTT. After 1.5 h incubation with the MTT solution, cell viability which can be recognized as deep blue formazan product from the yellow tetrazolium salts was measured at OD570 with iMark micro plate reader (Bio-Rad Laboratories, Hercules, CA, USA).

### 2.6. Immunohistochemistry

Cultured astrocytes were fixed with 4% paraformaldehyde in phosphate buffered saline (PBS) for 20 min. After several time washing with PBS, incubation with anti-GFAP antibody (1:1000) with 0.2% Triton-X (sigma) and 10% FBS in PBS was performed over-night followed by washing with fresh PBS. After incubation with secondary antibody, immunofluorescence image was captured with a fluorescent microscope (Observer Z1, Carl Zeiss, Oberkochen, Germany). The value of fluorescence intensity from 32 to 37 cells selected randomly in 4 dishes for each experimental condition was calculated.

### 2.7. The measurement of glutamate release

The amount of extracellular glutamate in cultured astroglial cells was measured according to the previous study [15]. Briefly, treatment with bFGF (24 h) or overexpression of miR-134 (48 h after transfection) was performed before washing 4 times with the modified Hepes-buffered Krebs Ringer solution (KRH; containing

130 mM NaCl, 10 mM glucose, 5 mM KCl, 1.8 mM  $\text{CaCl}_2$ , 1.2 mM  $\text{NaH}_2\text{PO}_4$ , 1% BSA, and 25 mM Hepes, pH 7.4). Then, the cultured astroglial cells were incubated with fresh KRH buffer for 20 min. Afterward, collected KRH samples were applied to the high-performance liquid chromatography (HPLC; Shimadzu Co., Kyoto, Japan).

### 2.8. Statistical analysis

Data were presented as mean  $\pm$  standard deviation (SD). Statistical analysis between two groups was performed with student's *t*-test. The data among multiple groups were analyzed by one- or two-way analysis of variance (ANOVA) with Bonferroni's multiple comparison *post hoc* test (SPSS Japan, Tokyo, Japan). Probability values less than 0.05 was defined as statistically significant.

## 3. Results

### 3.1. bFGF increased miR-134 expression in astrocytes

As we previously reported [8], miR-132, (but not miR-9, or -124a), was increased by BDNF and bFGF application in cultured cortical neurons (Fig. 1A). bFGF induced the expression of neural miR-134 although the increment was lower than miR-132 increase (Fig. 1A). Because it has been reported that astrocytes also express bFGF receptor [16], we tested whether bFGF increases brain-specific miRs in astrocyte. Though the expression levels of

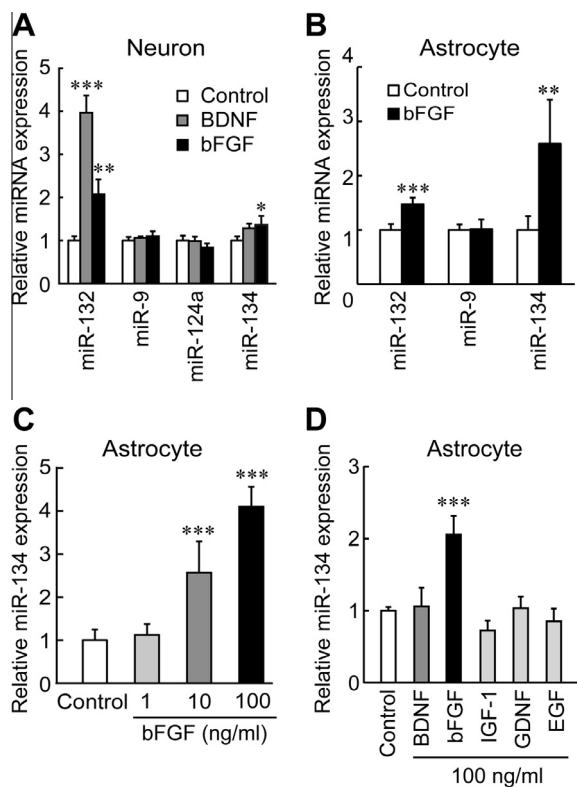
miR-132 were significantly increased by bFGF in astrocytes, an induction of miR-134 was much higher compared with that of miR-132 (Fig. 1B). To confirm the positive effect of bFGF on glial miR-134 expression, a dose-dependency of the expression was examined. Glial miR-134 expression was dose-dependently enhanced by bFGF (Fig. 1C). Previously, we confirmed significant activation of ERK, one of the intracellular signaling molecules, after bFGF, IGF-1, or EGF addition to astrocytes, suggesting an existence of specific receptor for each factor [8]. Interestingly, bFGF induced marked upregulation of miR-134 in astroglial cultures, while the other growth factors (including BDNF, IGF-1, GDNF, and EGF) did not increase miR-134 levels (Fig. 1D).

### 3.2. The involvement of ERK and PI3K signaling in the bFGF-induced miR-134

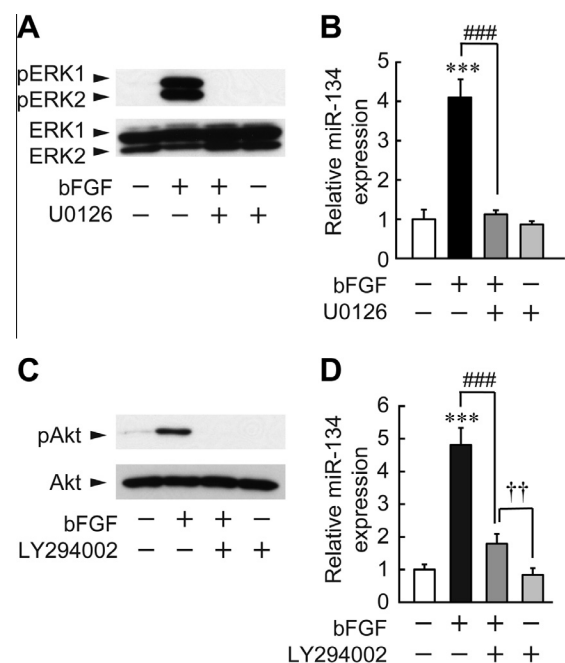
bFGF activates ERK and phosphatidylinositol 3-kinase (PI3K) signaling in astrocytes [17]. To clarify the contribution of intracellular signaling to the bFGF-induced miR-134, astrocytes were exposed to bFGF in the presence of MEK (MAP kinase kinase, upstream of ERK) inhibitor, U0126. U0126 completely inhibited both ERK activation and miR-134 upregulation induced by bFGF (Fig. 2A and B). LY294002, PI3K inhibitor, also inhibited Akt activation and significantly diminished the miR-134 induction by bFGF (Fig. 2C and D). These results suggest that ERK and Akt signaling are involved in the miR-134 induction by bFGF.

### 3.3. Overexpression of miR-134 did not affect cell survival in astrocytes

To investigate physiological role of glial miR-134, we performed miR-134 overexpression. Transfection of the miR-134 construct



**Fig. 1.** bFGF induced upregulation of miR-134 in astrocytes. (A) The expression of miRs (miR-132, -9, -124a, and -134) in cortical neurons after BDNF or bFGF stimulation. Cortical neurons at DIV 4–5 were exposed to 100 ng/ml BDNF or bFGF for 24 h ( $n = 4$ ), respectively. (B) Changed expression of miRs in astroglia after bFGF incubation for 24 h at 100 ng/ml ( $n = 5$ ). (C) Dose-dependency of astroglial miR-134 upregulation. bFGF (1–100 ng/ml) was added ( $n = 6$ ). (D) BDNF, bFGF, IGF-1, GDNF, or EGF were applied to astroglial cultures to evaluate their effect on miR-134 induction ( $n = 6$ ). The expressions of miRs were determined with real-time PCR. Error bars represent SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



**Fig. 2.** The involvement of MAPK/ERK and PI3K signaling cascades in the bFGF-induced miR-134. (A) Activated ERK1/2 (phosphorylated ERK1/2, pERK1/2) and total ERK1/2 by bFGF were examined. Astrocytes were exposed to bFGF (100 ng/ml, for 20 min) in the presence or absence of U0126 (10  $\mu\text{M}$ ). (B) Astroglial miR-134 expression after treatment with 100 ng/ml bFGF in the presence or absence of U0126 ( $n = 4$ –6). (C) Activated Akt (phosphorylated Akt) and total Akt after bFGF application. Astrocytes were exposed to 100 ng/ml bFGF for 20 min with or without LY294002 (10  $\mu\text{M}$ ). (D) Astroglial miR-134 levels after bFGF stimulation with or without LY294002 ( $n = 4$ ). Error bars represent SD. \*\*\* $P < 0.001$  vs control, \*\* $P < 0.01$ , ### $P < 0.001$ .

drastically increased miR-134 expression (Fig. 3A). Because bFGF plays a role in cell proliferation and differentiation in astrocytes [18,19], we checked cell viability after exogenous miR-134 induction, and found that this increment of miR-134 had no effect on cell viability (Fig. 3B). The basal levels of pERK and pAkt were not changed by the increase of miR-134 (Fig. 3C).

### 3.4. miR-134 overexpression enhances glial maturation

Next, when we measured an immunoreactivity of glia fibrillary acidic protein (GFAP, astrocytic cytoskeleton protein), increased intensity of immunofluorescence by the miR-134 overexpression was observed (Fig. 4A). In addition to increased GFAP, upregulation of other glial specific proteins including ALDH1L1 and S-100 expression after the miR-134 overexpression in astroglial cultures was observed (Fig. 4B). As expected, bFGF also increased the expression levels of astroglial markers, GFAP and ALDH1L1 (Fig. 4C).

To further assess physiological role of glial miR-134, we also examined the glial glutamate transporter, GLT-1, in the miR-134 overexpressed astroglial cultures, because the regulation of transmitter concentration through specific transporters is a main function of glial cells to maintain neurotransmission [20]. Importantly, expression levels of GLT-1 were increased by the overexpression of miR-134 and treatment of bFGF (Fig. 4D and E). Furthermore, both miR-134 overexpression and bFGF treatment decreased the extracellular glutamate concentration (Fig. 4F).

## 4. Discussion

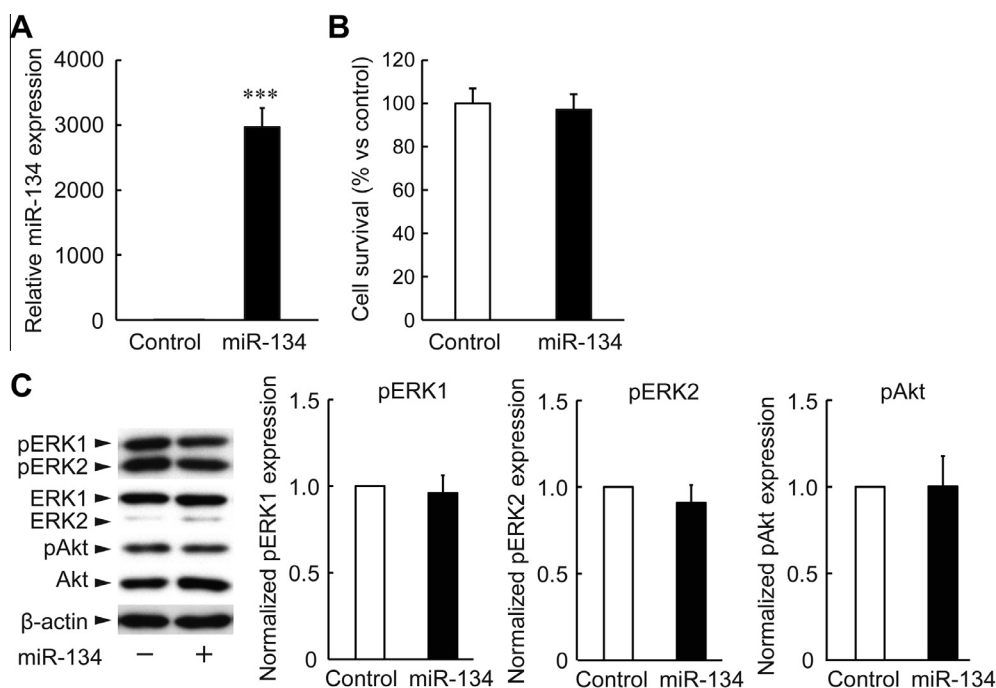
Here, we found that bFGF increased miR-134 expression in astrocytes. The upregulation of miR-134 was inhibited by co-application with an inhibitor for ERK- or Akt-signaling, respectively. Overexpression of miR-134 caused an increase in the expression

of glial specific proteins and glutamate transporter, and decreased extracellular glutamate concentration.

Evidence suggests important roles of miR-134 in neurons. Expression of neural miR-134 is linked to cell maturation and localized in dendrites to regulate hippocampal spine size via inhibiting Limk1 mRNA expression [12]. The translational repressor Pum-2 (Pumilio2) is another target of miR-134, which is involved in the dendritic outgrowth mediated by Mef-2 in an activity-dependent manner [14]. In this study, we found a glial role of miR-134 in relation to bFGF function in astrocyte. It is well known that bFGF regulates astroglial development including cell differentiation and proliferation [18,21]. Remarkably, FGF-2/FGF-5 double mutant mice exhibited significantly reduced astroglial marker proteins such as GFAP and S-100 in the prefrontal cortex [21]. Astroglial proliferation stimulated by bFGF was also reported [18]. In our pure astroglial cultures, overexpression of miR-134 and treatment with bFGF clearly increased not only the expression of glial specific proteins (GFAP, ALDH1L1 and S-100) but also the expression of GLT-1 without any effect on cell viability, suggesting that bFGF-increased miR-134 contributes to maturation of astrocytes.

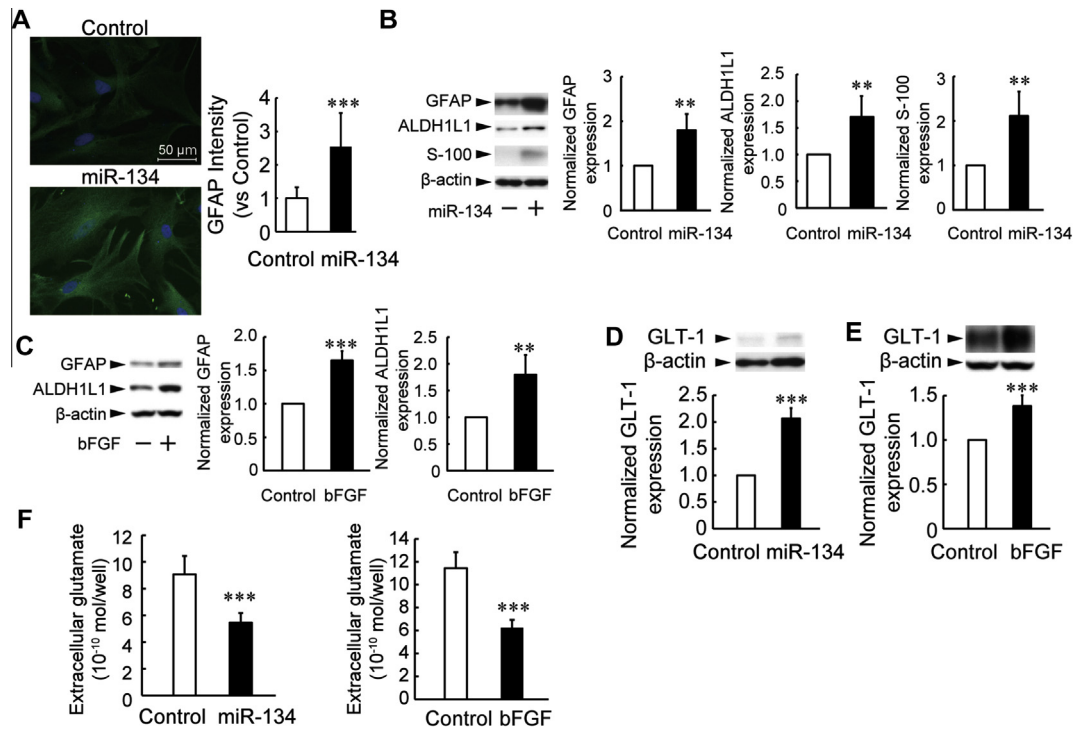
In both neuronal and astroglial cultures, we recently found an induction of miR-132 by bFGF through activating ERK pathway [8]. The ERK signal cascade is thought to have pivotal roles in the synaptic plasticity [22,23]. Importantly, ERK signaling stimulated by bFGF is involved in the process extension of astrocytes [17]. In the present study, it was revealed that miR-134 worked downstream of ERK activation in astrocytes and that the function of miR-134 is possibly cell maturation, since miR-134 overexpression resulted in increased levels of glial proteins and GLT-1.

The increase of hippocampal mitogen-activated protein kinase (MAPK) phosphatase-1 (MKP-1), a negative regulator for ERK activity, caused depressive-like behaviors in rodents with reduced ERK activation [24]. The inactivation of ERK in the dentate gyrus was also observed in depressive-like mice following chronic treatment with corticosterone, and an antidepressant (amitriptyline) restored abnormalities in both ERK signal and behaviors [25]. In concert,



**Fig. 3.** The effect of miR-134 overexpression on astroglial cell survival. (A) The levels of miR-134 after control- or miR-134-coding vectors were transfected into astrocytes ( $n = 5$ ). (B) Cell survival after control- or miR-134-coding vectors transfection. Error bars represent SD. ( $n = 12$ ). Cell viability was measured with MTT assay. (C) The activated levels of ERK and Akt signaling in control- or miR-134-coding vector transfected astrocytes ( $n = 5$ ). The values are mean  $\pm$  SD. \*\*\* $P < 0.001$  vs control.





**Fig. 4.** Overexpression of miR-134 enhanced expression of glial markers and glutamate transporter, and decreased extracellular glutamate concentration. (A) Immunostaining with anti-GFAP antibody in control- or in miR-134 expression vector-transfected astrocytes. The immunofluorescence intensity was measured using control- ( $n = 32$ ) or miR-134-overexpressed cells ( $n = 37$ ) from 4 dishes for each condition. (B) Glial marker proteins (GFAP, ALDH1L1, and S-100) expression in control or miR-134 overexpression in astrocytes ( $n = 5$ ). (C) The expression levels of GFAP and ALDH1L1 at 24 h after bFGF (100 ng/ml) application in astrocytes ( $n = 5$ ). (D) miR-134 overexpression and (E) bFGF application increased GLT-1 expression in astrocyte ( $n = 5$ ). (F) Reduced concentration of extracellular glutamate in astroglial cultures after miR-134 overexpression and bFGF application ( $n = 6-12$ ). The values are mean  $\pm$  SD. \*\* $P < 0.01$ , \*\*\* $P < 0.001$  vs control.

activity and expression of ERK were attenuated in the tissues from prefrontal cortex and hippocampus in depressed suicide subjects [26]. Moreover, B-Raf, a Raf kinase family of serine/threonine kinase which regulates ERK signaling, was diminished in the prefrontal cortical and hippocampal tissues from suicide subjects [27]. Additionally, decreased plasma levels of miR-134 in patients with bipolar disorder have also been reported [28]. It is possible that miR-134, which is regulated by ERK signaling, is a useful biomarker for mental disorders, though more detailed relationship between ERK activity and miR-134 expression should be clarified. On the other hand, it is well known that the PI3K/Akt signal cascade works for survival promoting in a variety of cell populations [29,30]. As induction of glial miR-134 by bFGF was also inhibited in the presence of a PI3K inhibitor, cell protection effect might be achieved by miR-134 induction when the astrocyte cultures are exposed to severer conditions including oxidative insult.

It has been demonstrated that elevated hippocampal mRNA expression of FGF receptor 1 was observed in patients with major depressive disorder who exhibited reduced bFGF mRNA expression [31]. Post-mortem studies suggest that a dysfunction of FGF system in the prefrontal cortex is associated with depression [31,32]. In kainic acid-induced seizure model mice, hippocampal injection of locked nucleic acid (LNA) 3' cholesterol-conjugated oligonucleotides (antagomirs) targeting miR-134 prevents status epilepticus and cell death induction [4]. Indeed, neuronal miRs have a critical role in physiological cellular states including neuronal development and function [1,2]. On the other hand, although the astroglial miRs such as miR-125b and -155 have been identified [9,10], physiological function of astroglial miRs is poorly understood. Therefore, it is intriguing to explore whether altered levels of bFGF and miR-134 in brain tissues from depression patients is neuronal or glial response.

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