

miR-124 promotes the neuronal differentiation of mouse inner ear neural stem cells

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Abstract. MicroRNAs (miRNAs or miRs) act as key regulators in neuronal development, synaptic morphogenesis and plasticity. However, their role in the neuronal differentiation of inner ear neural stem cells (NSCs) remains unclear. In this study, 6 miRNAs were selected and their expression patterns during the neuronal differentiation of inner ear NSCs were examined by RT-qPCR. We demonstrated that the culture of spiral ganglion stem cells present in the inner ears of newborn mice gave rise to neurons *in vitro*. The expression patterns of miR-124, miR-132, miR-134, miR-20a, miR-17-5p and miR-30a-5p were examined during a 14-day neuronal differentiation period. We found that miR-124 promoted the neuronal differentiation of and neurite outgrowth in mouse inner ear NSCs, and that the changes in the expression of tropomyosin receptor kinase B (TrkB) and cell division control protein 42 homolog (Cdc42) during inner ear NSC differentiation were associated with miR-124 expression. Our findings indicate that miR-124 plays a role in the neuronal differentiation of inner ear NSCs. This finding may lead to the development of novel strategies for restoring hearing in neurodegenerative diseases.

Introduction

Sensorineural hearing loss is a severe hearing impairment that affects millions of individuals worldwide. Moreover, neuronal regeneration is uncommon in the mature inner ear (1,2), in contrast to neurogenesis in the adult brain, such as in the subventricular zone (SVZ) of the lateral ventricles and the dentate gyrus of the hippocampus (3). Inner ear neural stem cells (NSCs) may exist in the spiral ganglion. Since NSCs can proliferate and differentiate into neurons (4-6), they may be used to regenerate damaged neurons in the inner ear. However, the intrinsic 'self-repair' capacity of inner ear NSCs is inactive *in vivo* for unclear reasons (unsuitable microenvironment, lack of guidance, etc.). Therefore, successfully harnessing the differentiation potential of inner ear NSCs may restore hearing. While the molecular mechanisms underlying the neuronal differentiation of inner ear NSCs are not yet fully understood, microRNAs (miRNAs or miRs) may play an important role in this process.

miRNAs are small non-coding RNAs that affect mRNA stability or inhibit translation by binding complementary sequences in the 3'-translated regions (3'-UTRs) of target mRNAs (7-10). Thus, miRNAs regulate multiple biological functions, including cell proliferation, differentiation and apoptosis (11,12). miRNA expression studies in mammals using microarrays and reverse transcription-quantitative PCR (RT-qPCR) have demonstrated that miRNAs are expressed in the developing nervous system and in mature neurons (13-18). miRNAs have crucial functions in neuronal development and plasticity (19,20). The miR-17 family has been shown to play an integral role in the regulation of neuronal differentiation (21). miR-124 expression has been shown to be upregulated during neuronal differentiation (22,23). Several lines of evidence have indicated that miR-124 regulates neuronal differentiation by inhibiting small C-terminal domain phosphatase (SCP1), a component of the REST-silencing transcription factor (REST)/neuron-restrictive silencer factor (NRSF) transcriptional repression complex, and by targeting polypyrimidine tract

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binding protein 1 (PTBP1), a global repressor of brain-specific alternative splicing (24,25). Moreover, global miRNA expression profiling by microarray analysis, RT-qPCR, and/or *in situ* hybridization have revealed several miRNAs with distinct spatio-temporal expression patterns in the embryonic and post-natal mouse inner ear (25-28). While miRNAs regulate the development, morphogenesis and function of the inner ear (29-33), it is not yet known whether miRNAs also regulate the neuronal differentiation of inner ear NSCs.

In this study, 6 miRNAs (miR-124, miR-132, miR-134, miR-20a, miR-17-5p and miR-30a-5p) were selected based on their involvement in neuronal differentiation/neurogenesis as discussed in the 'Introduction' and in the 'Discussion'; their expression patterns during the neuronal differentiation of inner ear NSCs were examined by RT-qPCR. Our data demonstrate that miR-124 is important for the differentiation of inner ear NSCs into neurons. Our results revealed that miR-124 expression is upregulated during neuronal differentiation. Moreover, miR-124 increased the percentage of cells expressing neuron-specific class III β -tubulin in culture, and increased the neurite length in mouse inner ear NSCs. These changes were accompanied by changes in the expression of tropomyosin receptor kinase B (TrkB) and cell division control protein 42 homolog (Cdc42). TrkB is a receptor of brain-derived neurotrophic factor (BDNF), and participates in the regulation of neurogenesis, neurite outgrowth and in the survival of spiral ganglion neurons. The small GTP-ase, Cdc42, which regulates both microtubules and actin filaments in a number of cells, regulates the neurite extension of spiral ganglion neurons. Our data demonstrated that miR-124 promotes the neuronal differentiation of and neurite outgrowth in inner ear NSCs, and regulates the expression of TrkB and Cdc42 in inner ear NSCs.

Materials and methods

Animals. Post-natal day 1 (P1) C57BL/6 mice (n=135; Laboratory Animal Center of Sun Yat-sen University, Guangzhou, China) were used for the experiments. The animals were sacrificed according to policies set forth by the Animal Care and Use Committee of Sun Yat-sen University based on the National Institutes of Health guidelines for animal care. All animal experiments were also approved the Animal Care and Use Committee of Sun Yat-sen University.

Inner ear NSC cultures, neuronal differentiation and immunostaining. The isolation and culture of NSCs from the spiral ganglia of newborn C57BL/6 mice were performed as previously described (4). Spiral ganglia, isolated from 5 mice, were digested in 200 μ l of 0.125% trypsin in phosphate-buffered saline (PBS; Gibco/Thermo Fisher Scientific, Waltham, MA, USA) at 37°C for 15 min. The cells were carefully triturated with plastic 200 μ l pipette tips, centrifuged and suspended in 2 ml Dulbecco's modified Eagle's medium (DMEM)/high glucose and F12 medium (mixed 1:1; Gibco/Thermo Fisher Scientific) supplemented with N2 and B27 supplements (Invitrogen/Thermo Fisher Scientific), epidermal growth factor (EGF; 20 ng/ml), basic fibroblast growth factor (bFGF; 20 ng/ml), insulin-like growth factor 1 (IGF-1; 20 ng/ml) (PeproTech, Rocky Hill, NJ, USA). The cell suspension was passed through a 70-mm cell strainer (BD Falcon, Franklin Lakes, NJ, USA) to

remove cell clumps. For neuronal differentiation, mouse inner ear NSCs at 5 days were harvested and cultured in laminin and poly-L-lysine-coated dishes in DMEM-F12 (1:1) medium supplemented with 20 ng/ml of neurotrophin-3 (NT-3), 20 ng/ml of BDNF (both from PeproTech), 10 ng/ml of leukemia inhibitory factor (LIF) (R&D Systems, Inc., Minneapolis, MN, USA), 2 mM of L-glutamine (Gibco/Thermo Fisher Scientific) and 3 mM of KCl (Sigma, St. Louis, MO, USA) for 7 days. Half of the differentiation medium was replaced every other day. The NSCs were allowed to differentiate for up to 14 days (day 14). Cell proliferative ability was assessed by BrdU incorporation assay as previously described (34). Undifferentiated and differentiated inner ear NSCs were fixed with 4% paraformaldehyde and immunostained with antibodies specific for BrdU (C2181; 1:10,000; Sigma), TuJ1 (class III β -tubulin; MO15013; 1:250; Neuromics, Edina, MN, USA), p27^{Kip1} (554069; 1:150; BD Biosciences, San Jose, CA, USA), ATP-binding cassette sub-family G member 2 (Abcg2; sc-25822; 1:150; Santa Cruz Biotechnology, Inc., Dallas, TX, USA), Sox2 (SAB2104660; 1:250; Sigma), Nestin (MO15012; 1:200; Neuromics), TrkB (ab18987; 1:200; Abcam, Cambridge, UK) and Cdc42 (sc-8401; 1:150 Santa Cruz Biotechnology, Inc.). Nuclei were counterstained with Fluoroshield Mounting Medium with 4',6-diamidino-2-phenylindole (DAPI; ab104139; Abcam).

RNA isolation and reverse transcription-quantitative PCR (RT-qPCR). The cells were collected in cold D-Hank's solution (Gibco/Thermo Fisher Scientific), and RNA was extracted using TRIzol reagent (Invitrogen/Thermo Fisher Scientific) and column-purified with an RNeasy kit [Qiagen (Suzhou) Translational Medicine Co., Ltd., Suzhou Industrial Park, China]. Reverse transcription was performed using M-MLV reverse transcriptase (Roche Diagnostics, Mannheim, Germany) according to the instructions provided by the manufacturer. The sequences of the primers were as follows: Nestin forward, 5'-GCCGAGCTGGAGCGCGA GTTAGAG-3' and reverse, 5'-GCAAGGGGGAAGAGAA GGATGTCTG-3'; Sox2 forward, 5'-ACCCGGGCCTCAA CGCTCACG-3' and reverse, 5'-TCCCCTTCTCCAGTT CGCAGTCC-3'; p27^{Kip1} forward, 5'-CTGGAGCGGAT GGACGCCAGAC-3' and reverse, 5'-CGTCTGCTCCAC AGTGCCAGC-3'; Abcg2 forward, 5'-GCTGTGGAGC TGTTCTAGTGG-3' and reverse, 5'-GCTAAAGTA CTGAAGCCAGGAC-3'; Nanog forward, 5'-CCCTTCCC TCGCCATCACA-3' and reverse, 5'-ACCGCTTGC ACTTCATCCTTTG-3'; TuJ1 forward, 5'-TAGACCCCA GCGGCAACTAT-3' and reverse, 5'-GTTCCAGG TTCCAAGTCCACC-3'; and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) forward, 5'-AACGGGAAG CCCATCACC-3' and reverse, 5'-CAGCCTTGGCAG CACCAG-3'. Primers were synthesized by Invitrogen. Transcripts of GAPDH were amplified and used as an internal reference. Quantitative PCR (qPCR) was performed using 2 μ l of SYBR-Green/ROX qPCR Master Mix (Suzhou GenePharma Co., Ltd., Suzhou, China). All procedures were subjected to a TaqMan miRNA assay (Applied Biosystem 7500) according to the manufacturer's instructions. U6 was employed as an endogenous control (for miRNAs) to normalize data using the $2^{-\Delta\Delta C_t}$ method. Probe and primers specific for

miR-124, miR-132, miR-134, miR-20a, miR-17-5p, miR-30a-5p, or for U6 RNA internal control were purchased from LC Sciences (Hangzhou, China). PCR and qPCR analyses of at least 3 independent cultures were performed.

Western blot analysis. For protein extraction, the cells were washed twice and resuspended in lysis buffer containing 50 mM of Tris-HCl, pH 7.4, 150 mM of NaCl, 1% NP-40, 10% glycerol, 1 mM of sodium orthovanadate, 1 mM of PMSF and a protease inhibitor cocktail (Shanghai Roche Pharmaceutical Co. Ltd., Shanghai, China). Subsequently, 20 μ g of cell lysates were separated by 8-15% of resolving sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), transferred onto polyvinylidene fluoride membranes, and immunoblotted with antibodies according to the manufacturer's instructions. The antibodies used in western blot analysis included β -actin (sc-47778; 1:3,000; Santa Cruz Biotechnology, Inc.), Tuj1 (MO15013; 1:1,000, Neuromics), TrkB (ab18987; 1:1,000; Abcam) and Cdc42 (sc-8401; 1:1,000; Santa Cruz Biotechnology, Inc.).

Electroporation. RNA oligoribonucleotides for miR-124 mimics (which mimic endogenous mature miR-124 molecules), non-specific miRNA (negative control) and miR-124 inhibitor [small interfering RNA (siRNA) targeting miR-124 (miR-124 siRNA)] were all purchased from Genepharma (Shanghai, China). For electroporation, mouse inner ear NSCs were dissociated into a single-cell suspension with 0.25% trypsin (Gibco/Thermo Fisher Scientific) at 37°C for 3 min. Subsequently, 1×10^6 cells were electroporated with 5 μ g of miR-124 mimics, non-specific miRNA or miR-124 siRNA using an Amaxa Mouse NSC Nucleofector kit (Lonza Group Ltd., Basel, Switzerland) following the manufacturer's instructions. RNA oligoribonucleotides specific for enhanced green fluorescent protein (eGFP) were used to monitor electroporation efficiency. Electroporated cells were plated on cover glasses coated with poly-L-lysine and laminin (R&D Systems, Inc.), and incubated in differentiation medium at 37°C. The transfected inner ear NSCs were then subjected to neuronal differentiation for up to 3 days.

Measurement of neuron percentage and neurite length. Transfected inner ear NSCs were fixed and immunostained at 3 days post-differentiation. The neuron percentage was determined as the number of cells stained positive for the neuron-specific marker, Tuj1, divided by the number of cells stained with Hoechst 33342 (the total number of differentiated inner ear NSCs) from 15 randomly selected fields at $\times 20$ magnification. All neurons within the image frame were counted. Cells located in clumps were excluded. Images were acquired using an Olympus microscope [Olympus IX71 Inverted Microscope and Olympus FV1000 confocal microscope; Olympus Inc. (China), Guangzhou, China] and quantitative analysis was carried out using ImageJ software (each image of $\times 20$ magnification generally captures at least 200 differentiated inner ear NSCs).

As previously described (35), we measured the entire length of the longest neurite extending from 95 differentiated neurons (identified by the presence of Tuj1), which was selected randomly (using a 40x objective lens). Neurites that

were not entirely in the frame were excluded. Neurite lengths were determined using ImageJ software. Processes that could not be distinguished from others were eliminated.

Statistical analysis. One-way ANOVA and Chi-square tests were used to analyze the data, and the differences between variables were considered to be statistically significance when the value was $P < 0.05$. All data are expressed as the means \pm SEM, unless otherwise stated.

Results

Culture and differentiation of inner ear NSCs. The cells were isolated from the mouse spiral ganglion and were cultured for 5 days. The cells formed floating solid neurospheres (Fig. 1A), which were positive for BrdU, Sox2 and other stem cell markers, such as Nestin, p27^{Kip1} and Abcg2 (Fig. 1B), confirming the self-renewal and proliferative capacity of these cells. To further characterize the stemness properties of the inner ear neurospheres, we confirmed the expression of markers of progenitor cells (Nestin, p27^{Kip1}, Abcg2 and Nanog) by RT-qPCR (Fig. 1C).

To initiate the differentiation of mouse inner ear NSCs into neurons, we plated the neurospheres onto a coated substratum in differentiation medium. Four hours after adherence, differentiated neurons that expressed neuron-specific class III β -tubulin (detected by Tuj1 antibody) began to sprout neurites from the attached neurospheres. Immunofluorescence staining demonstrated that the number of Tuj1-positive cells increased on day 3, and reached maximal levels on day 14 (Fig. 2A-D). To confirm the differentiation of NSCs, the expression of neuron-specific class III β -tubulin was examined by western blot analysis and by RT-qPCR (Fig. 2E-F). Proliferating spheres expressed low levels of neuron-specific class III β -tubulin, which is a marker of mature neurons. This phenomenon may be related to the differentiation of individual peripheral NSCs. During differentiation, the levels of neuron-specific class III β -tubulin increased on day 3 and reached the highest levels on day 14, which was consistent with the data obtained by immunofluorescence.

TrkB and Cdc42 are upregulated during the neuronal differentiation of inner ear NSCs. Although Cdc42 and TrkB have been shown to affect neuronal differentiation and polarity (36-40), the role of TrkB and Cdc42 in the neuronal differentiation of mouse NSCs remains unknown. In this study, we cultured inner ear NSCs in differentiation medium for 14 days; our results revealed that TrkB and Cdc42 expression were low in the undifferentiated inner ear NSCs, and their expression gradually increased during differentiation (Fig. 3C-D). Since TrkB and Cdc42 modulate neuronal differentiation and neurite extension, TrkB and Cdc42 may be involved in the differentiation of inner ear NSCs into neurons.

Expression of miRNAs during the differentiation of inner ear NSCs. To determine the function of miRNAs in the neuronal differentiation of inner ear NSCs, we compared the expression levels of 6 miRNAs between undifferentiated and differentiated inner ear NSCs. A total of 6 miRNAs were selected based on their established role in neuronal differentiation.

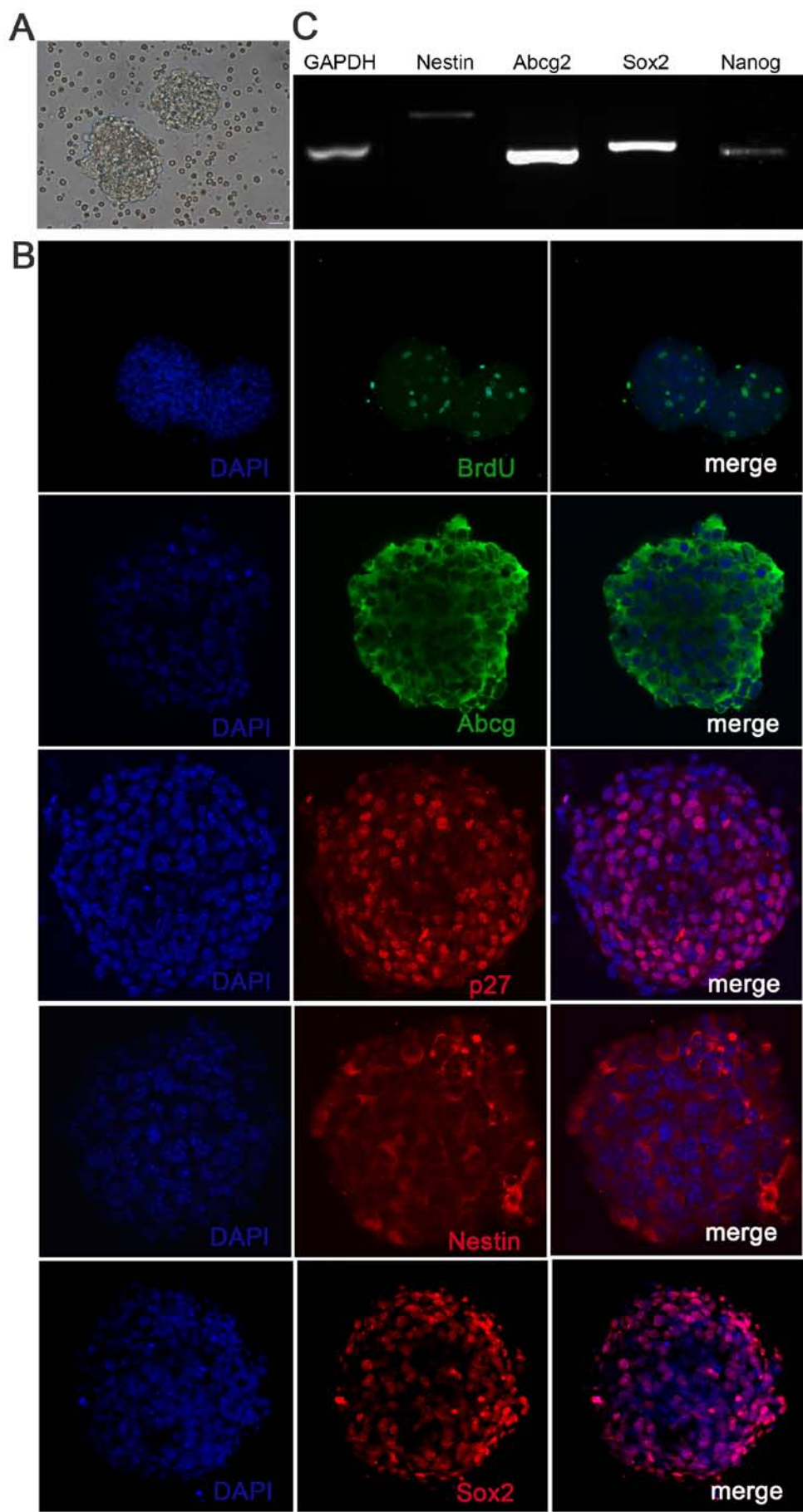


Figure 1. Inner ear neural stem cells (NSCs) isolated from newborn mice proliferate and form floating neurospheres. Neurospheres were immunostained after 5 days under proliferative conditions. (A) Typical morphology of a spiral ganglion-derived neurosphere after 5 days of culture in suspension. (B) Neurospheres were positive for BrdU, Sox2, Nestin, Abcg2 and p27^{Kip1}. (C) Analysis of stemness-related genes in inner ear NSCs by RT-qPCR.

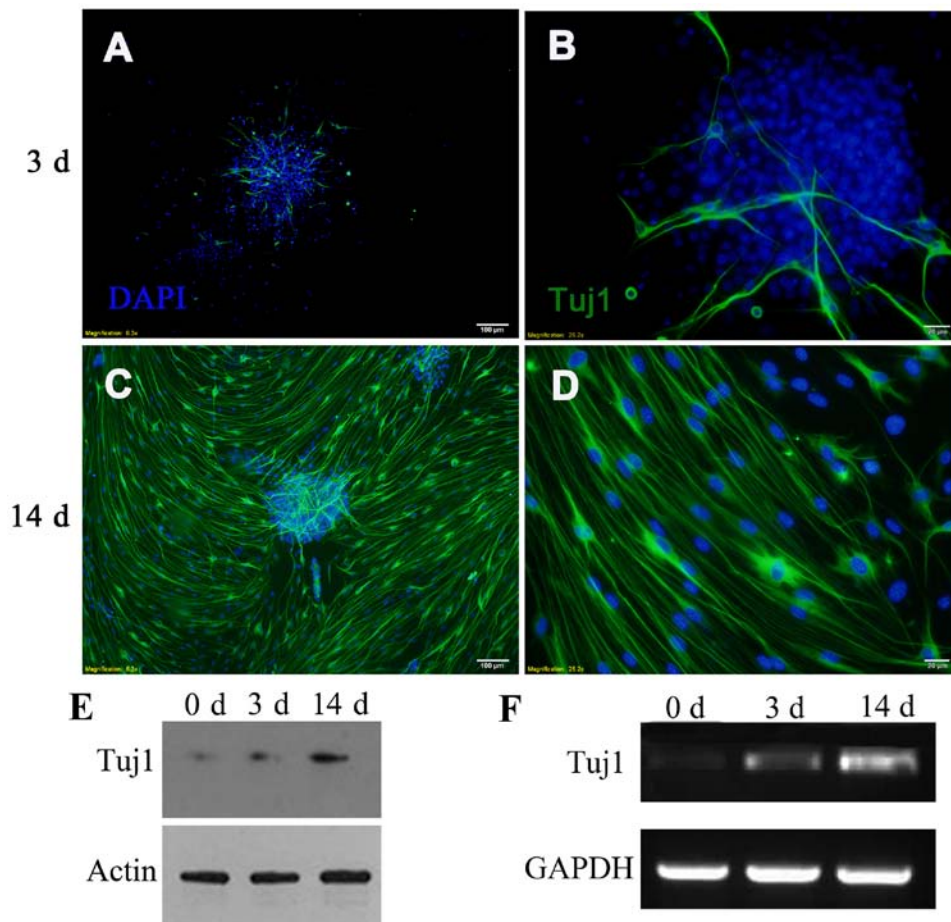


Figure 2. Tuj1 expression increases during neuronal stem cell (NSC) differentiation. (A) Two representative images of inner ear NSC differentiation (days 3 and 14). Mature neurons were visualized with Tuj1 (green), and the nuclei were visualized with 4',6-diamidino-2-phenylindole (DAPI) (blue). (A and C) Lower magnification (x10) scale bars, 100 μm; (B and D) higher magnification (x40) scale bars, 20 μm. Neuron-specific class III β-tubulin expressions (E) and mRNA that encode neuron-specific class III β-tubulin (F) in undifferentiated cells (0 days) and during differentiation (3 and 14 days). d, days.

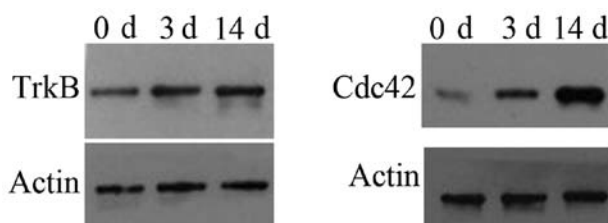


Figure 3. The expression of cell division control protein 42 homolog (Cdc42) and tropomyosin receptor kinase B (TrkB) during the differentiation of inner ear neural stem cells (NSCs). Western blot analysis of Cdc42 and TrkB in undifferentiated (0 day) and differentiated NSCs (3 and 14 days). d, days.

We set the expression level in undifferentiated inner ear NSCs as the baseline, and compared it to the expression in differentiated inner ear NSCs on days 3 and 14 using the $2^{-\Delta\Delta Ct}$ method. The levels of miR-17-5p decreased by day 3 ($p < 0.05$) and increased by day 14 ($p < 0.05$); however, the overall expression level of miR-17-5p did not show any significant change over the 14-day period (Fig. 4). Similarly, no significant change in miR-20a expression levels were observed during neuronal differentiation ($p > 0.05$). By contrast, the miR-30a-5p expression levels increased and the miR-134 levels significantly decreased during the differentiation of NSCs on day 14,

suggesting that miR-30a-5p and miR-134 play a role in neuronal differentiation. The expression levels of 2 miRNAs, miR-124 and miR-132 were altered on days 3 and 14 of differentiation. The levels of miR-124 increased 6-fold, while the levels of miR-132 decreased 4-fold during the entire 14-day period. In this study, we focused on the role of miR-124 in the differentiation of inner ear NSCs.

miR-124 promotes the neuronal differentiation of and neurite outgrowth in inner ear NSCs. Inner ear NSCs were transfected with RNA oligonucleotides for miR-124 mimics, non-specific miRNA and miR-124 inhibitor. The transfection efficiency was 50-60%, as determined by eGFP (Fig. 5A and B). In addition, we measured the miR-124 expression levels in differentiated inner ear NSCs by RT-qPCR. Three days following electroporation with miR-124 inhibitor, the miR-124 levels were decreased by 2.13-fold. By contrast, the electroporation of miR-124 mimics led to a 3.02-fold increase in miR-124 expression (Fig. 5C).

Next, we determined the effect of miR-124 on the percentage of neuron and neurite outgrowth. We found that the overexpression of miR-124 in differentiated inner ear NSCs increased neuronal differentiation and neurite outgrowth, whereas the knockdown of miR-124 had an opposite effect (Fig. 6A-C). The results revealed that at 3 days of differentia-

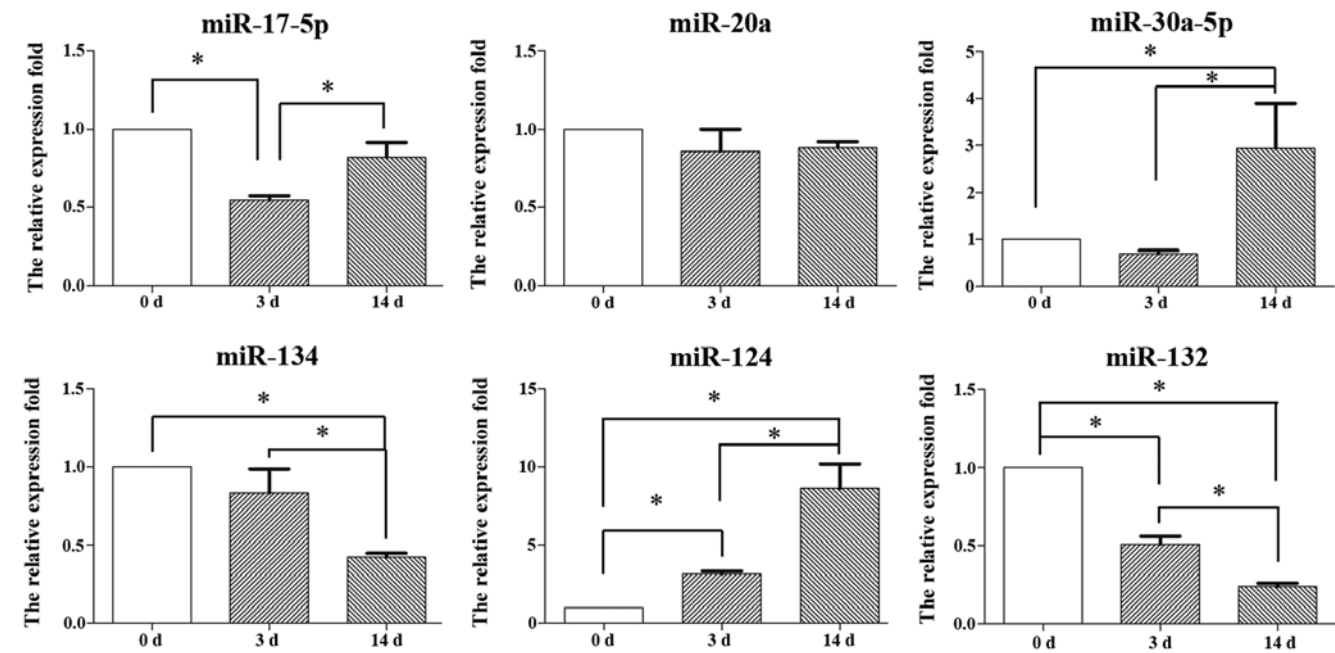


Figure 4. Results of RT-qPCR of the miRNA expression in mouse inner ear during the differentiation of neural stem cells (NSCs). Expression of miRNAs in undifferentiated (0 days) or differentiated (3 or 14 days) inner ear NSCs. The RT-qPCR data were normalized to U6 expression levels. The fold change was expressed as the ratio of miRNA expression in differentiated and undifferentiated cells (day 0) using the $2^{-\Delta\Delta Ct}$ method. * $p < 0.05$, statistically significant difference. d, days.

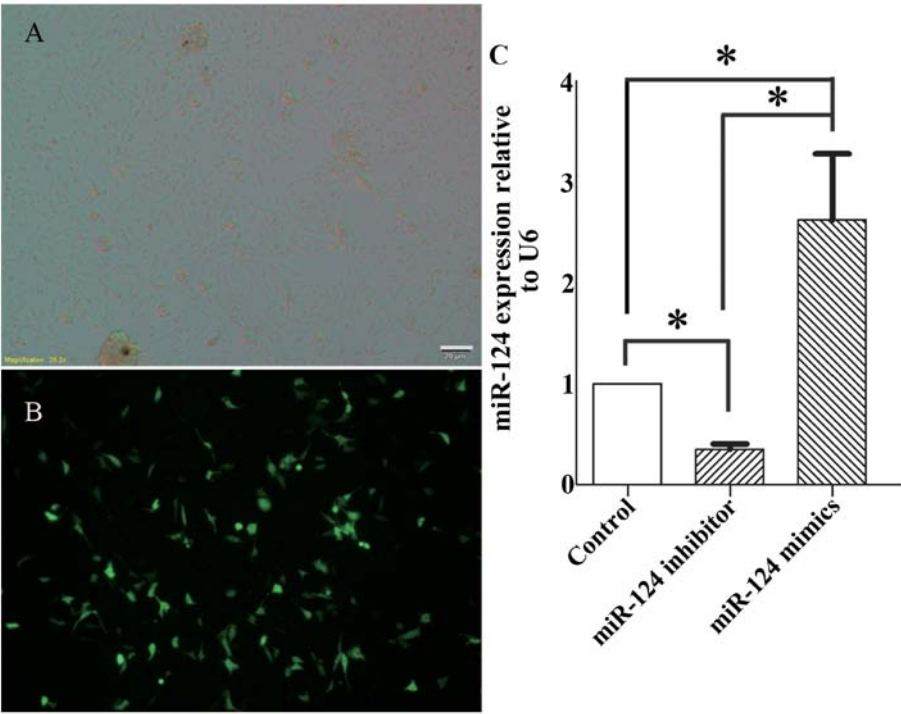


Figure 5. Transfection efficiency of inner ear neural stem cell (NSC) cultures after differentiation for 3 days. (A and B) Images of inner ear NSCs transfected with RNA oligoribonucleotides for enhanced green fluorescent protein. Scale bars, 100 μm . (C) miR-124 expression levels in transfected cells were determined by RT-qPCR and were normalized to U6 levels. * $p < 0.05$, statistically significant difference.

tion, $3.53 \pm 0.46\%$ of the cells transfected with miR-124 mimics were Tuj1-positive cells, while the percentage of Tuj1-positive cells in the controls was $1.99 \pm 0.06\%$ ($p < 0.05$; Fig. 6D). Moreover, the average neurite length of the Tuj1-positive cells transfected with the miR-124 mimics was $113.22 \pm 7.31 \mu\text{m}$, which was significantly higher than that of the control cells

($98.83 \pm 4.18 \mu\text{m}$; $p < 0.05$) (Fig. 6E). By contrast, $1.05 \pm 0.25\%$ of the cells transfected with miR-124 inhibitor were Tuj1-positive, while the percentage of Tuj1-positive cells in the controls was $1.99 \pm 0.06\%$ ($p < 0.05$; Fig. 6D). Additionally, the average neurite length of the Tuj1-positive cells transfected with miR-124 inhibitor was reduced to $83.12 \pm 3.40 \mu\text{m}$, which was

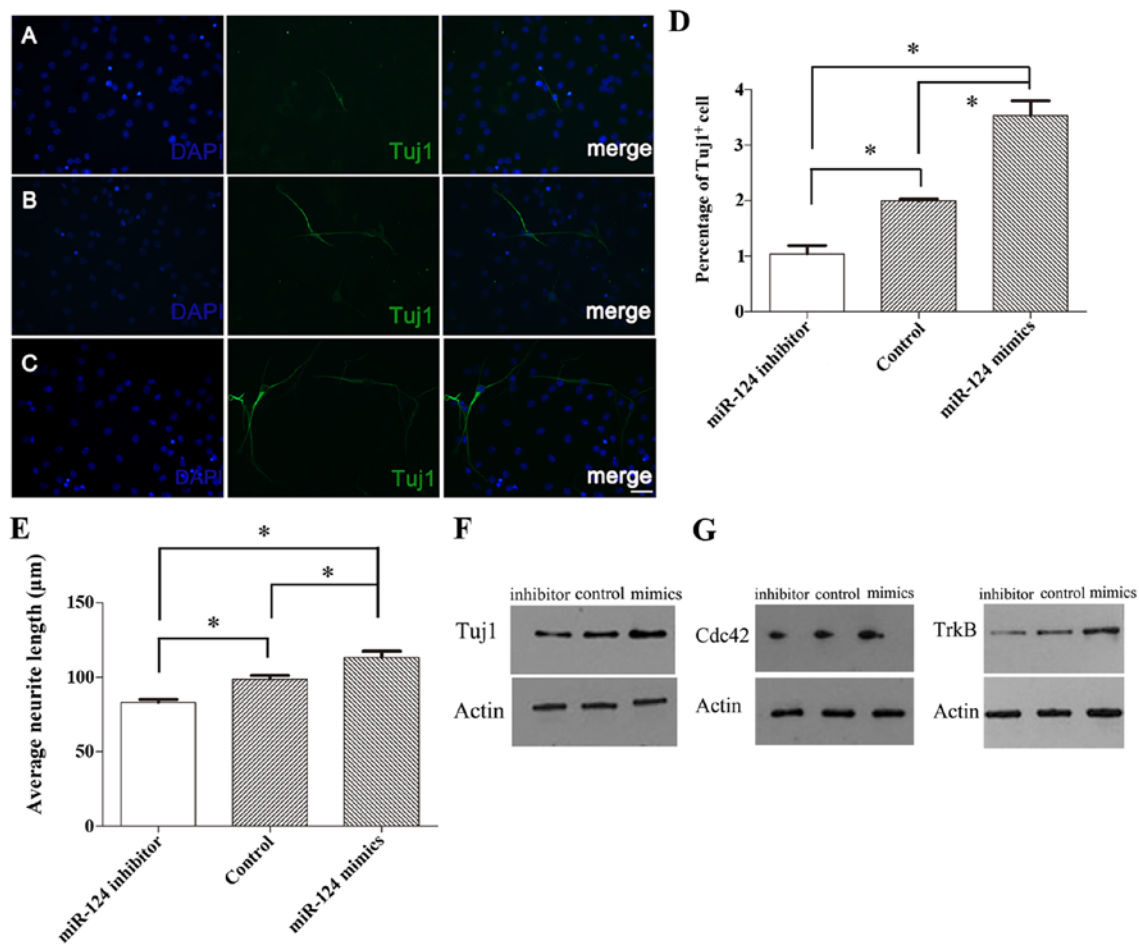


Figure 6. miR-124 promotes the neuronal differentiation of and neurite outgrowth in inner ear neural stem cells (NSCs). Inner ear NSCs transfected with (A) miR-124 inhibitor, (B) control, or (C) miR-124 mimics were differentiated into neurons for 3 days. Cells were immunostained with TuJ1 antibody (red) and nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI) (blue). (D) Quantification of TuJ1-positive cells in differentiated inner ear NSCs 3 days after transfection. (E) Analysis of average neurite length examined 3 days after differentiation. (F) Western blot analysis of neuron-specific class III β -tubulin in neurons derived from inner ear NSCs 3 days after differentiation. (G) Levels of tropomyosin receptor kinase B (TrkB) and cell division control protein 42 homolog (Cdc42) in inner ear NSCs transfected with miR-124 inhibitor or miR-124 mimics. Actin was used to confirm equal loading. * $p < 0.05$, statistically significant difference. d, days.

significantly lower than that of the controls ($98.83 \pm 4.18 \mu\text{m}$; $p < 0.05$) (Fig. 6E).

We confirmed that inner ear NSCs transfected with miR-124 mimics had higher levels of neuron-specific class III β -tubulin compared to the control cells, while inner ear NSCs transfected with miR-124 inhibitor had a reduced expression of neuron-specific class III β -tubulin (Fig. 6F). These findings are in agreement with the immunofluorescence observations and confirmed the role of miR-124 in neuronal differentiation.

miR-124 affects the expression of TrkB and Cdc42 in the neuronal differentiation of mouse inner ear NSCs. TrkB and Cdc42 play a role in neuronal differentiation and neurite elongation. We demonstrated that Cdc42 and TrkB were upregulated during neuronal differentiation. Our findings revealed that miR-124 promoted the differentiation of inner ear NSCs, which led us to investigate whether miR-124 modulates TrkB and Cdc42 expression in inner ear NSCs. We evaluated the TrkB and Cdc42 expression levels in inner NSCs transfected with miR-124 by western blot analysis (Fig. 6G). Compared to the control cells, miR-124 overexpression increased Cdc42 and TrkB expression during neuronal differentiation. Conversely,

miR-124 knockdown caused a marked decrease in the TrkB and Cdc42 protein levels, suggesting that Cdc42 and TrkB act as downstream effectors of miR-124 during inner ear NSC differentiation.

Discussion

NSCs are neural progenitor cells that have the ability to self-renew, proliferate and differentiate into lineages, such as neurons, oligodendroglia and astroglia (41). Thus, NSCs, which have regenerative potential may serve as an important source for replacing damaged neurons in hearing-impaired individuals. It is therefore important to understand the differentiation pathway of inner ear NSCs into neurons. While miRNAs are generally important in neuronal development and differentiation (19,20), the profiling of miRNAs in the inner ear has been a major research goal (27,28,42,43). Conditional Dicer knockout mice have major defects in the inner ear, including hair cell malformations, abnormal innervations, disrupted neurosensory development and hearing loss (29,33). A role of miR-96 in progressive hearing loss in mice and humans has been reported. Furthermore, miR-181a influences

proliferation within the basilar papilla (44), while miR-200b can cause severe malformations of the auditory and vestibular labyrinth by affecting the *Zeb1* pathway (45). miR-183 may affect zebrafish inner ear organization (31), as well as mouse inner ear function (46,47). However, the role of miRNAs in the neuronal differentiation of mouse inner ear NSCs warrants further investigation.

In this study, we examined the expression of 6 miRNAs in the differentiation of mouse inner ear NSCs into neurons. In this study, the levels of miR-30a-5p, an established inhibitor of BDNF expression (48), increased 14 days after neuronal differentiation, suggesting that miR-30a-5p plays a role in the late stages of neuronal differentiation or neurite plasticity. In contrast to previously published results (48-53), miR-134 and miR-132 were downregulated during the neuronal differentiation of inner ear NSCs. Although miR-20a has been shown to be downregulated in the neuronal lineage differentiation of unrestricted somatic stem cells (USSCs) from cord blood and NSCs (54,55), no changes in miR-20a were observed during differentiation of NSCs in our study. In addition, in our study, miR-17-5p expression increased during the first 3 days of differentiation and then increased on day 14; however, in a previous study, its expression decreased continued to decrease in differentiated SH-SY5Y cells (21). The differential expression of specific miRNAs may be due to specific neuronal types and time points being investigated.

In this study, we focused on the role of miR-124, which has been shown to enhance the development of mature neurons (56,57) and to play a role in the neuronal differentiation of several cell lines *in vivo* and *in vitro* (16,22,23,25,58). Moreover, miR-124 acts as a key mediator in regulating the differentiation of human embryonic and mesenchymal stem cells (59,60). We demonstrated that miR-124 was detectable at low levels in the undifferentiated inner ear NSCs; however, its expression gradually increased and peaked on day 14 of differentiation. The overexpression of miR-124 increased the percentages of neurons and neurite length, whereas the knock-down of miR-124 had an opposite effect, demonstrating that miR-124 is important for the *in vitro* neuronal differentiation of inner ear NSCs.

Cdc42 is a member of the Rho GTPases and regulates both microtubules and actin filaments in a variety of cells (61-63), and plays a role in the establishment of neuronal polarity (36,37,39,64). In addition, TrkB, the cognate receptor of BDNF, participates in multiple aspects of neuronal functions (65), modulating neurite outgrowth, structural plasticity and survival (36,38-40). Cdc42 functions downstream of BDNF stimulation (36,40,66,67). Furthermore, BDNF activates Cdc42 via the Cdk5-mediated phosphorylation of TrkB, which in turn influences dendritic growth (36). TrkB and Cdc42 play a role in neural development and plasticity. The upregulation of TrkB and Cdc42 expression during inner ear NSC differentiation suggests that TrkB acts upstream of Cdc42 in the regulation of the neuronal differentiation of and neurite outgrowth in mouse inner ear NSCs.

Recently, several studies have examined the function of Cdc42 in dendrite and axon development, as well as its regulation by miR-124 (58,68). It has been shown that miR-124 upregulates the expression of Cdc42 (24). Moreover, an increased TrkB level of has been shown to be accompanied by

the upregulation of miR-124 in exercise-dependent neuronal differentiation in the hippocampus of adult male rats (69).

In this study, we demonstrated that miR-124 promoted neuronal differentiation and neurite outgrowth, and we investigated the effects of miR-124 on TrkB and Cdc42 expression during the neuronal differentiation of and neurite outgrowth in inner ear NSCs. We demonstrated that TrkB and Cdc42 expression coincided with miR-124 during inner ear NSC differentiation. These results suggest that TrkB and Cdc42 may play an important role in miR-124-regulated neuronal differentiation. In support of this hypothesis, we found that miR-124 regulates their expression, raising the possibility that the effects of miR-124 on inner ear NSC differentiation and neurite outgrowth are mediated by TrkB and Cdc42.

Although TrkB promotes neuronal survival and plasticity (36,38-40), Cdc42 may have opposite effects on neurite outgrowth. Several studies have revealed that Cdc42 enhances dendritic growth or arborization (36,37,39), while other studies have demonstrated negative effects (58,68,70). These studies are consistent with our hypothesis, wherein Cdc42 promotes neurite outgrowth. In the inner ear, TrkB is enriched in cochlear neurons and plays a role in neuronal survival and structural plasticity (38,71,72). In contrast to our findings and the results reported by Brors *et al* (73), Rac/Cdc42 inhibition has been shown to enhance neurite formation and decrease neurite length in cochlear spiral ganglion explants in neonatal rat pups (74).

In conclusion, our results suggest that miR-124 promotes the neuronal differentiation of and neurite outgrowth in inner ear NSCs by regulating TrkB and Cdc42. Future studies are required to determine the mechanisms through which miR-124 controls TrkB and Cdc42 expression to regulate neuronal differentiation in the mouse inner ear.

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