MicroRNA-132 protects hippocampal neurons against oxygen-glucose deprivation–induced apoptosis

Zu-Zhen Sun, Zhan-Yun Lv, Wen-Jing Tian and Yan Yang

Abstract

Hypoxic-ischemic brain injury (HIBI) results in death or long-term neurologic impairment in both adults and children. In this study, we investigated the effects of microRNA-132 (miR-132) dysregulation on oxygen-glucose deprivation (OGD)-induced apoptosis in fetal rat hippocampal neurons, in order to reveal the therapeutic potential of miR-132 on HIBI. MiR-132 dysregulation was induced prior to OGD exposure by transfection of primary fetal rat hippocampal neurons with miR-132 mimic or miR-132 inhibitor. The effects of miR-132 overexpression and suppression on OGD-stimulated hippocampal neurons were evaluated by detection of cell viability, apoptotic cells rate, and the expression of apoptosis-related proteins. Besides, TargetScan database and dual luciferase activity assay were used to seek a target gene of miR-132. As a result, miR-132 was highly expressed in hippocampal neurons following 2 h of OGD exposure. MiR-132 overexpression significantly increased OGD-diminished cell viability and reduced OGD-induced apoptosis at 12, 24, and 48 h post-OGD. MiR-132 overexpression significantly down-regulated the expressions of Bax, cytochrome c, and caspase-9, but up-regulated BCl-2. Caspase-3 activity was also significantly decreased by miR-132 overexpression. Furthermore, FOXO3 was a direct target of miR-132, and it was negatively regulated by miR-132. To conclude, our results provide evidence that miR-132 protects hippocampal neurons against OGD injury by inhibiting apoptosis.

Keywords

apoptosis, FOXO3, hippocampal neuron cells, hypoxic-ischemic brain injury, microRNA-132, oxygen-glucose deprivation

Introduction

Hypoxic-ischemic brain injury (HIBI) is a leading cause of mortality and morbidity, which is caused by intrapartum or late antepartum brain hypoxia and ischemia.1 This potentially devastating event leads to death or long-term neurologic impairment in both adults and children, with fetal distress and hypoxia caused by situations such as the umbilical cord wrapped around the neck.2–3 Despite the recent advances in perinatal care, perinatal hypoxia–ischemia remains a tragic cause of neonatal death and/or severe neurologic disorders.4 In particular, ischemia and hypoxia in the perinatal period occurs in 1–6 of every 1000 live-births5,6 and approximately 40% of newborn infants with HIBI die during the neonatal period, while 30% suffer lifelong neurologic impairment.7,8 The consequences of
HIBI require complex and continuous management and new treatments are urgently needed to improve patient outcomes.

MicroRNAs (miRNAs) represent a class of highly conserved small, non-coding RNA molecules that are widely expressed in eukaryotic cells. These molecules regulate protein expression by binding to specific sequences in messenger RNA (mRNA) molecules, most commonly in the 3'-UTR.9,10 MiRNAs are critically involved in controlling a wide variety of biological processes including differentiation, proliferation, and apoptosis.11 It has been reported that approximately 70% of known miRNAs are expressed specifically or enriched in the brain12 and play important roles in the development and function of the nervous system. Using oxygen-glucose deprivation (OGD) to mimic an ex vivo model of HIBI, miR-210 has been found to exert neuroprotective effects by inhibiting PC12 cells apoptosis.13,14 Furthermore, a recent in vivo study provided the evidence that miR-210 inhibition might be a potential therapeutic approach to combat neonatal ischemic hypoxia.15 Reduced levels of miR-659-3p have been shown to correlate with increased levels of neuroprotective progranulin in some brain injury conditions caused by OGD and hypoxic stress.16

In the previous study, miR-132 has been identified as a conserved and neuron-enriched miRNA.17 Knockdown or mutation of miR-132 caused severe intracranial hemorrhage and disruption of brain vascular integrity in zebrafish larvae.18 Besides, it was also delineated that overexpression of miR-132 in primary cultures of hippocampal neurons or delivered directly into the CA1 of living rats by means of the lentiviral expression system prior to induction of ischemia afforded robust protection against ischemia-induced neuronal death.19 These studies indicate that inhibition of miR-132 expression plays an important role in neuronal cell death under ischemic conditions.

In this study, OGD was established in primary hippocampal neurons to mimic an in vitro model of HIBI, since miR-132 has been reported to exert neuroprotective functions under OGD but not under hydrogen peroxide.20 The effects of miR-132 on OGD-induced apoptosis were assessed, and the underlying mechanism(s) in which miR-132 protected hippocampal neurons was detected. The findings of this study will add to the growing literature that targeting neuroprotective pathways controlled by miR-132 may represent a therapeutic strategy for the treatment of HIBI.

Materials and methods

Hippocampal neuron isolation and culture

Hippocampal neuron was isolated from specific pathogen-free grade fetal Sprague-Dawley (SD) rats (14 days, Vital River Laboratories, Beijing, China). All the animal experiments conducted in this study were approved by the Animal Ethics Committee of Affiliated Hospital of Jining Medical University and were performed according to the instructions of our institute.

For hippocampal neuron isolation, rats were anesthetized with chloral hydrate (2 mL/kg body weight) and then hippocampus was removed and cut into small chunks (~2 mm³). After washing with Hanks’ balanced salt solution (HBSS, Bogoo, Shanghai, China) twice, hippocampus was dissociated in 0.125% trypsin (Sigma-Aldrich, St. Louis, MO, USA) for 40 min at 37°C. Cells were collected by centrifugation at 400 g for 10 min, suspended in neurobasal (NB) medium (Gibco, Carlsbad, CA, USA) containing 2% B27 supplement (Gibco), 500 µM l-glutamine (Gibco), and penicillin-streptomycin (100 U/mL, Sigma-Aldrich). Cells (75,000 in 50 µL medium) were plated in four-well plates (13-mm diameter wells) coated with poly-d-lysine (0.05 mg/mL) and incubated at 37°C with 95% air and 5% CO₂.

OGD model

For OGD, hippocampal neurons were incubated in glucose-free Dulbecco’s modified Eagle’s medium (DMEM, Gibco) at 37°C in a hypoxic incubator with 0.2% O₂, 5% CO₂, and 94.8% N₂. Control neuron cells (non-OGD conditions) were cultured in glucose-free DMEM at 37°C with 95% air and 5% CO₂. Hippocampal neurons were subjected to OGD for 2h. Glucose-free medium was then replaced with NB medium and cultures were returned to a normoxic 95% air and 5% CO₂ incubator.

MiRNA transfection

The miR-132 mimic, RNA-132 inhibitor, and corresponding negative control were designed and synthesized by GenePharma Co., Ltd. (Shanghai,
Transfections were performed using Lipofectamine 2000 reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s instructions. The final concentration of miR-132 mimic, miR-132 inhibitor, and control was 50 nM, 200 nM, and 100 nM. At 48 h after transfection, the cells were collected for the use in the following experiments.

**Cell viability assay**

Cell viability was assessed using the Cell Counting Kit-8 (CCK-8, Dojindo Molecular Technologies, Gaithersburg, MD). The miR-transfected hippocampal neurons were seeded into 96-well plates (5 × 10^3 cells/well) for adherence. After 2 h of exposure in OGD conditions, and another 12–48 h incubation in the fresh NB medium, 10 µL CCK-8 solution was added, and the cultures were incubated for 1 h at 37°C. Cell viability was determined by measuring the absorbance of the culture supernatants at 450 nm using a Microplate Reader (Bio-Rad, Hercules, CA, USA).

**Apoptosis assay**

Cell apoptosis analysis was performed by flow cytometry using the FITC-conjugated Annexin V apoptosis detection kit (BD Pharmingen, San Jose, CA, USA). The miR-transfected hippocampal neurons were seeded into six-well plates (5 × 10^5 cells/well) for adherence. After 2 h of exposure in OGD conditions, and 12–48 h of incubation in the fresh NB medium, 10 µL FITC and 10 µL propidium iodide (PI) were added to the cultures. The samples were then incubated in the dark at 4°C for 30 min in the dark. Flow cytometric analysis was conducted using a FACScalibur (Beckman Coulter, Fullerton, CA, USA), and the data were analyzed using FlowJo software (Tree Star, San Carlos, CA, USA).

**RNA extraction and quantitative real-time polymerase chain reaction analysis**

Total RNA was extracted from hippocampal neurons using TRIzol reagent (Life Technologies Corporation, Carlsbad, CA, USA) according to the manufacturer’s instructions. Complementary DNA (cDNA) was synthesized using a PrimeScript RT reagent Kit (Takara, Dalian, China), and quantitative real-time polymerase chain reaction (qRT-PCR) was performed using SYBR Premix Ex Taq II Reagent Kit (Takara) according to the manufacturer’s instructions. MiR-132 expression was normalized to the endogenous snRNA U6. Relative miR-132 expression was calculated using the 2^(-ΔΔCt) method, where ΔCt = (Ct target gene – Ct U6), and Ct represents the cycle threshold.

**Western blot analysis**

Hippocampal neurons were collected and lysed for 30 min in lysis buffer (Beyotime Biotechnology, Shanghai, China) containing 1 mM phenylmethane sulfonyl fluoride (Beyotime Biotechnology). The cells were then centrifuged at 400g for 10 min at 4°C. Protein concentrations in the cell lysates were determined by the BCA protein assay (Beyotime Biotechnology). Equal amounts of cell lysates (20 µg) were separated by 10%–12% sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) before transfer to polyvinylidene difluoride (PVDF) membranes. After blocking for 30 min in phosphate-buffered saline (PBS) containing 1% bovine serum albumin with 0.05% Tween 20, the membranes were incubated at 4°C overnight with primary antibodies for the detection of BCL-2, Bax, cytochrome c, caspase-9, and FOXO3 (1:1000 dilution, Abcam, Cambridge, MA, USA). Glyceraldehyde 3-phosphate dehydrogenase (GAPDH; 1:1000 dilution, Immuno Way Biotechnology, USA) was detected as an internal control. After three washes with Tris-buffered saline with Tween 20 (TBST), the membranes were incubated for 1 h at room temperature with alkaline phosphatase-anti-rabbit IgG (1:1000 dilution; ZSGB-BIO, Beijing, China) or alkaline phosphatase-rabbit anti-goat IgG (1:1000 dilution; ZSGB-BIO). Western Blue® stabilized substrate alkaline phosphatase (Promega, Madison, WI, USA) was used to visualize the immunoreactive signals. Band intensities of the air-dried membranes were analyzed using an image analyzer (Bio-Rad Laboratories, Hercules, CA, USA) and Image J v1.50 software.

**Caspase-3 activity assay kit**

Hippocampal neurons were collected by centrifugation at 600g for 5 min at 4°C. The supernatant was discarded and cells were washed once with PBS. Cell lysates were prepared as described in western blot analysis, and caspase-3 activity was
measured using the colorimetric caspase-3 activity assay kit (Beyotime Biotechnology) according to the manufacturer’s instructions. After the addition of the caspase-3 substrate Ac-DEVD-pNA (2 mM), the reaction mixture was mixed and incubated at 37°C for 60–120 min. The absorbance was measured immediately at 405 nm using a spectrophotometer.

Dual luciferase activity assay

The fragments from FOXO3 containing the predicted miR-132 binding site was amplified by PCR and cloned into a pmirGLO Dual-luciferase miRNA Target Expression Vector (Promega) to form the reporter vector FOXO3-wild-type (FOXO3-wt). To mutate the putative binding site of miR-132 in the FOXO3, the sequences of putative binding site were replaced and were named as FOXO3-mutated-type (FOXO3-mt). Cells were co-transfected with the reporter constructs and miR-132 mimic or miR-132 inhibitor, or their control using Lipofectamine 2000 (Invitrogen). Reporter assays were done using the dual-luciferase assay system (Promega) following the manufacturer’s information.

Statistical analysis

All statistical analyses were performed using GraphPad Prism software version 6.0 (GraphPad Software Inc., San Diego, CA, USA) and the data are presented as means ± standard derivations (SD) of three independent experiments. Differences among groups were analyzed using repeated-measure one-way analysis of variance (ANOVA) comparison test. A value of $P<0.05$ was considered to indicate statistical significance.

Results

MiR-132 was highly expressed in hippocampal neurons following OGD exposure

To determine if miR-132 expression is involved in hypoxic-ischemic induced neuronal death, the expression of miR-132 in OGD-stimulated hippocampal neurons was monitored. qRT-PCR analysis results showed that (Figure 1(a)) the expression of miR-132 in hippocampal neurons was increased to 1.97 ± 0.14 folds following OGD exposure ($P<0.001$), indicating miR-132 was significantly up-regulated by OGD stimulus. Next, we transfected miR-132 mimic, miR-132 inhibitor, or their control into hippocampal neurons and detected the transfection efficiency by qRT-PCR. As shown in Figure 1(b), when compared with the control group, cells transfected with the miR-132 mimic exhibited significant overexpression ($P<0.001$), while transfection with the miR-132 inhibitor resulted in a significant decrease in expression ($P<0.01$).
Effects of miR-132 expression on OGD-diminished cell viability

The effects of miR-132 dysregulation on hippocampal neurons viability following OGD were analyzed by CCK-8 assay. We found that miR-132 overexpression resulted in a significant viability increase on the cells cultured for 12–48 h ($P < 0.05$ or $P < 0.01$) when compared to the control group. In contrast, miR-132 suppression resulted in a significant decrease in cell viability at the same time points ($P < 0.05$ or $P < 0.01$, Figure 2).

Effects of miR-132 expression on apoptosis-associated proteins

To further investigate the underlying mechanism(s) via which miR-132 expression impacted OGD-stimulated hippocampal neurons, we focused on the expression changes in several apoptosis-related proteins. We found that miR-132 overexpression resulted in a significant decrease in the expression of Bax, cytochrome c, and caspase-9 on the cells cultured for 12–48 h ($P < 0.05$ or $P < 0.01$, Figure 4(a)–(d)). It seemed that miR-132 overexpression had no impact on the expression of BCL-2 on the cells cultured for 12 and 24 h, while significant up-regulation of BCL-2 was observed in miR-132 overexpressing cells cultured for 48 h ($P < 0.05$). In contrast, miR-132 suppression resulted in a significant decrease in BCL-2 expression and significant increases in Bax, cytochrome c, and caspase-9 expression on cells cultured for 12–48 h ($P < 0.05$, $P < 0.01$, or $P < 0.001$).

Effects of miR-132 expression on apoptosis-associated proteins

The effects of miR-132 expression on OGD-induced apoptosis in hippocampal neurons were analyzed by flow cytometry after FITC-Annexin V and PI double-staining. As shown in Figure 3(a) and (b), miR-132 overexpression significantly decreased the proportion of apoptotic neurons compared with that of cells transfected with the control ($P < 0.05$ at 12 and 24 h; $P < 0.01$ at 48 h), while miR-132 suppression resulted in a significant increase in the proportion of apoptotic cells ($P < 0.05$ at 12 h; $P < 0.01$ at 24 h; $P < 0.001$ at 48 h).

Effects of miR-132 expression on caspase-3 activity

To further elucidate the role of miR-132 expression on hippocampal neurons apoptosis following OGD, the activity of caspase-3 was detected using a colorimetric assay. As shown in Figure 5, the activity of caspase-3 was significantly decreased in miR-132 overexpressing cells, while was increased in miR-132-suppressing cells at 12, 24, and 48 h post-OGD ($P < 0.05$, $P < 0.01$, or $P < 0.001$).

Effects of miR-132 expression on FOXO3 expression

It is well known that repression of mRNAs by miRNAs is an important mechanism for regulation of expression during cell fate specification, apoptosis, and metabolism. Therefore, we sought the mRNA targets of miR-132 in order to further understand the mechanism via which miR-132 protected hippocampal neurons from OGD exposure. Using TargetScan database, we found that none of Bax, caspase-9, and cytochrome c was a target of miR-132, while FOXO3 was predicted as a potential target of miR-132 (Figure 6(a)). To confirm this prediction, dual luciferase activity assay was performed, and we found that the luciferase activity was significantly reduced by co-transfection with...
FOXO3-wt and miR-132 mimic (P<0.01, Figure 6(b)), suggesting FOXO3 was a direct target of miR-132. Then, qRT-PCR and western blot analyses were performed to assess the regulatory effects of miR-132 on FOXO3. As shown in Figure 6(c) and (d), both the mRNA and protein levels of FOXO3 were down-regulated by miR-132 overexpression, while was up-regulated by miR-132 suppression (P<0.01 or P<0.001), indicating that FOXO3 was negatively regulated by miR-132.

**Discussion**

MiRNAs are essential for normal brain development and for establishing the functional connectivity of the brain. MiR-132 has been widely reported as a neuron-enriched miRNA which plays important roles in the development and activity-dependent plasticity of neural systems. In this study, we showed that miR-132 was highly expressed in hippocampal neurons following OGD exposure. Overexpression of miR-132 protected hippocampal neurons from OGD-diminished viability and OGD-induced apoptosis by regulation of BCL-2, Bax, cytochrome c, caspase-9, and caspase-3. Furthermore, FOXO3 was a direct target of miR-132 and was negatively regulated by miR-132. To our knowledge, this is the first demonstration that miR-132 protects hippocampal neurons against OGD injury by inhibiting apoptosis.

Altered miRNA expression has been linked with a variety of brain diseases and injury, including HIBI. Previous studies have showed that miR-132 was abnormally expressed in ischemic brain damage, while the description of how hypoxic-ischemic alters miR-132 expression profiles is
controversial. Lusardi et al.\textsuperscript{25} demonstrated that miR-132 was reduced in a preconditioning model of focal stroke. In contrast to Lusardi et al., Keasey et al. demonstrated that miR-132 was increased following OGD.\textsuperscript{20} Yao et al.\textsuperscript{26} also reported hypoxia-induced up-regulation of miR-132 in Schwann cells. In this study, we employed qRT-PCR to explore the expression of miR-132 in

**Figure 4.** Overexpression of miR-132 altered the expression of apoptosis-related proteins in hippocampal neurons following OGD. (a–d) Primary hippocampal neurons were transfected with miR-132 mimic or miR-132 inhibitor before exposure to OGD for 2 h. Expression of apoptosis-related proteins in hippocampal neurons was analyzed by Western blotting at 12, 24, and 48 h post-OGD. Data represent the mean ± SD of three independent experiments. \(^{*}P<0.05, \quad ^{**}P<0.01, \quad ^{***}P<0.001\) compared to control group.
OGD-stimulated hippocampal neurons, and our findings were consistent with the studies of Keasey et al. and Yao et al. that miR-132 was up-regulated following OGD. This finding hit that increased expression of miR-132 may be involved in OGD-induced damage in hippocampal neurons.

MiR-132 is enriched in brain tissue and plays important roles in neuronal differentiation and development. The neuroprotective functions of miR-132 have been well established. At a cellular level, miR-132 regulates neurite growth and arborization, and synaptic structure and function. At a system level, miR-132 is essential for experience-dependent structural and functional plasticity of the visual cortex. A previous study has demonstrated that miR-132 could reduce OGD-induced neuronal death, as evidenced by the increase in cell viability. In consistence with the previous study, we found that miR-132 increased OGD-diminished viability in hippocampal neurons. Moreover, we for the first time revealed that miR-132 reduced OGD-induced apoptosis, which further confirmed the neuroprotective role of miR-132 in OGD-induced injury.

To further elucidate the underlying mechanism(s) via which miR-132 protected hippocampal neurons from OGD-induced injury, we conducted western blot analysis for detecting the expression
of several apoptosis-related proteins. We found that miR-132 overexpression prior to OGD exposure resulted in significant down-regulation of Bax, cytochrome c, and caspase-9 and a significant up-regulation of BCL-2. These data further confirmed the neuroprotective functions of miR-132 on OGD-induced injury and provided the evidence that miR-132 plays an anti-apoptotic role in regulating the expression of pro- and anti-apoptotic proteins. However, miR-132 silencing has been reported to decrease recurrent epileptic seizures by reducing neuron cell apoptosis, with reduced expression of both BCL-2 and Bax.30 This discrepancy may be due to differences in the models investigated and highlights the complexity of the control of cell processes exerted by miRNAs.

Cytochrome c release has been implicated in mitochondrial control of apoptosis and activation of caspase-3.31 Caspase-3 and caspase-9 are critically involved in the process of programmed cell death in the central nervous system.32,33 Thus, we further detected the caspase-3 activity to reveal whether miR-132 protects OGD-induced apoptosis via regulation of caspase-3. As a result, we found that miR-132 overexpression significantly reduced caspase-3 activity. Thus, our findings indicate that miR-132 protects hippocampal neurons against OGD injury by inhibiting apoptosis via a mechanism that involves regulation of caspase activity and the expression of BCL-2 and Bax. Although further studies are required to confirm the results of this study in vivo, it has recently been reported that miR-132 protects against neuronal cell death in a mouse model of intracerebral hemorrhage.34

FOXO transcription factors are pivotal regulators in maintaining cellular homeostasis.35 Among the mammalian FOXO family members, FOXO3 is highly homologous in its protein sequences and has been reported as a pro-apoptotic factor in tumor cells.36,37 FOXO3 is also associated with oxidative stress.38 In a previous study, it has been demonstrated that miR-132 plays an important role in hematopoietic stem cells cell-cycling, differentiation, and apoptosis via buffering FOXO3 protein expression.39 In this study, we showed that FOXO3 was a direct target of miR-132, and it was negatively regulated by miR-132. These findings suggested that miR-132 plays a role in buffering perturbations in the expression of FOXO3 in response to OGD, and miR-132 protected OGD-induced apoptosis possibly via inhibition of FOXO3.

In summary, this study demonstrates that miR-132 protects hippocampal neurons against OGD-induced apoptosis via a mechanism that possibly involves the regulation of FOXO3, and caspases, as well as Bax and BCL-2. These findings provide an improved understanding of the potential of miR-132 as a therapeutic target for fetal HIBI.

Declaration of conflicting interests
The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Funding
The author(s) received no financial support for the research, authorship, and/or publication of this article.

References
1. Xu LX, Lv Y, Li YH, et al. (2017) Melatonin alleviates brain and peripheral tissue edema in a neonatal rat model of hypoxic-ischemic brain damage: The involvement of edema related proteins. BMC Pediatrics 17: 90.
2. Chen A, Xiong LJ, Tong Y, et al. (2013) The neuroprotective roles of BDNF in hypoxic ischemic brain injury. Biomedical Reports 1: 167–176.
3. Fitch RH, Alexander ML and Threlkeld SW (2013) Early neural disruption and auditory processing outcomes in rodent models: Implications for developmental language disability. Frontiers in Systems Neuroscience 7: 58.
4. Lawn JE, Cousens S and Zupan J (2005) 4 million neonatal deaths: When? Where? Why? Lancet 365: 891–900.
5. Hristova M, Rocha-Ferreira E, Fontana X, et al. (2016) Inhibition of signal transducer and activator of transcription 3 (STAT3) reduces neonatal hypoxic-ischaemic brain damage. Journal of Neurochemistry 136: 981–994.
6. Lai Z, Zhang L, Su J, et al. (2016) Sevoflurane post-conditioning improves long-term learning and memory of neonatal hypoxia-ischemia brain damage rats via the PI3K/Akt-mPTP pathway. Brain Research 1630: 25–37.
7. Charon V, Proisy M, Ferre JC, et al. (2015) Comparison of early and late MRI in neonatal hypoxic-ischemic encephalopathy using three assessment methods. Pediatric Radiology 45: 1988–2000.
8. Li D, Li X, Wu J, et al. (2015) Involvement of the JNK/FOXO3a/Bim pathway in neuronal apoptosis after hypoxic-ischemic brain damage in neonatal rats. PLoS ONE 10:e0132998.
9. Lal A, Navarro F, Maher CA, et al. (2009) miR-24 Inhibits cell proliferation by targeting E2F2, MYC, and other cell-cycle genes via binding to “seedless”
3'UTR microRNA recognition elements. *Molecular Cell* 35: 610–625.

10. Zhao Y, Samal E and Srivastava D (2005) Serum response factor regulates a muscle-specific microRNA that targets Hand2 during cardiogenesis. *Nature* 436: 214–220.

11. Flynt AS and Lai EC (2008) Biological principles of microRNA-mediated regulation: Shared themes amid diversity. *Nature Reviews Genetics* 9: 831–842.

12. Nowak JS and Michlewski G (2013) miRNAs in development and pathogenesis of the nervous system. *Biochemical Society Transactions* 41: 815–820.

13. Qiu J, Zhou XY, Zhou XG, et al. (2013) Neuroprotective effects of microRNA-210 against oxygen-glucose deprivation through inhibition of apoptosis in PC12 cells. *Molecular Medicine Reports* 7: 1955–1959.

14. Qiu J, Zhou XY, Zhou XG, et al. (2015) MicroRNA-210 knockdown contributes to apoptosis caused by oxygen glucose deprivation in PC12 cells. *Molecular Medicine Reports* 11: 719–723.

15. Ma Q, Dasgupta C, Li Y, et al. (2016) Inhibition of microRNA-210 provides neuroprotection in hypoxic-ischemic brain injury in neonatal rats. *Neurobiology of Disease* 89: 202–212.

16. Piscopo P, Grasso M, Fontana F, et al. (2016) Reduced miR-659–3p levels correlate with progranulin increase in hypoxic conditions: Implications for frontotemporal dementia. *Frontiers in Molecular Neuroscience* 9: 31.

17. Magill ST, Cambronne XA, Luikart BW, et al. (2010) microRNA-132 regulates dendritic growth and arborization of newborn neurons in the adult hippocampus. *Proceedings of the National Academy of Sciences of the United States of America* 107: 20382–20387.

18. Xu B, Zhang Y, Du XF, et al. (2017) Neurons secrete miR-132-containing exosomes to regulate brain vascular integrity. *Cell Research*. Epub ahead of print 21 April 2017. DOI: 10.1038/cr.2017.62.

19. Hwang JY, Kaneko N, Noh KM, et al. (2014) The gene silencing transcription factor REST represses miR-132 expression in hippocampal neurons destined to die. *Journal of Molecular Biology* 426: 3454–3466.

20. Keasey MP, Scott HL, Bantoumas I, et al. (2016) MiR-132 is upregulated by ischemic preconditioning of cultured hippocampal neurons and protects them from subsequent OGD toxicity. *Journal of Molecular Neuroscience* 59: 404–410.

21. Bartel DP (2004) MicroRNAs: Genomics, biogenesis, mechanism, and function. *Cell* 116: 281–297.

22. Yang Y, Sandhu HK, Zhi F, et al. (2015) Effects of hypoxia and ischemia on microRNAs in the brain. *Current Medicinal Chemistry* 22: 1292–1301.

23. Garberg HT, Huun MU, Baumbusch LO, et al. (2017) Temporal profile of circulating microRNAs after global hypoxia-ischemia in newborn piglets. *Neonatology* 111: 133–139.

24. Stammet P. Blood biomarkers of hypoxic-ischemic brain injury after cardiac arrest. *Seminars in Neurology* 37: 75–80.

25. Lusardi TA, Farr CD, Faulkner CL, et al. (2010) Ischemic preconditioning regulates expression of microRNAs and a predicted target, MeCP2, in mouse cortex. *Journal of Cerebral Blood Flow and Metabolism* 30: 744–756.

26. Yao C, Shi X, Zhang Z, et al. (2016) Hypoxia-induced upregulation of miR-132 promotes Schwann cell migration after sciatric nerve injury by targeting PRKAG3. *Molecular Neurobiology* 53: 5129–5139.

27. Schratt GM, Tuebing F, Nigh EA, et al. (2006) A brain-specific microRNA regulates dendritic spine development. *Nature* 439: 283–289.

28. Wayman GA, Davare M, Ando H, et al. (2008) An activity-regulated microRNA controls dendritic plasticity by down-regulating p250GAP. *Proceedings of the National Academy of Sciences of the United States of America* 105: 9093–9098.

29. Remenyi J, Hunter CJ, Cole C, et al. (2010) Regulation of the miR-212/132 locus by MSK1 and CREB in response to neurotrophins. *Biochemical Journal* 428: 281–291.

30. Huang Y, Guo J, Wang Q, et al. (2014) MicroRNA-132 silencing decreases the spontaneous recurrent seizures. *International Journal of Clinical and Experimental Medicine* 7: 1639–1649.

31. Porter AG and Janicke RU (1999) Emerging roles of caspase-3 in apoptosis. *Cell Death and Differentiation* 6: 99–104.

32. Kuida K, Haydar TF, Kuan CY, et al. (1998) Reduced apoptosis and cytochrome c-mediated caspase activation in mice lacking caspase 9. *Cell* 94: 325–337.

33. Kuida K, Zheng TS, Na S, et al. (1996) Decreased apoptosis in the brain and premature lethality in CPP32-deficient mice. *Nature* 384: 368–372.

34. Zhang Y, Han B, He Y, et al. (2016) MicroRNA-132 attenuates neurobehavioral and neuropathological changes associated with intracerebral hemorrhage in mice. *Neurochemistry International*. Epub ahead of print 9 December 2016. DOI: 10.1016/j.neuint.2016.11.011.

35. Eijkelenboom A and Burgering BM (2013) FOXOs: Signalling integrators for homeostasis maintenance. *Nature Reviews Molecular Cell Biology* 14: 83–97.

36. Sunters A, Fernández de Mattos S, Stahl M, et al. (2003) FoxO3a transcriptional regulation of Bim controls apoptosis in paclitaxel-treated breast cancer cell lines. *Journal of Biological Chemistry* 278: 49795.
37. Ho WC, Pikor L, Gao Y, et al. (2012) Calpain 2 regulates Akt-FoxO-p27(Kip1) protein signaling pathway in mammary carcinoma. *Journal of Biological Chemistry* 287: 15458–15465.

38. Sun Z, Yan B, Yu WY, et al. (2016) Vitexin attenuates acute doxorubicin cardiotoxicity in rats via the suppression of oxidative stress, inflammation and apoptosis and the activation of FOXO3a. *Experimental & Therapeutic Medicine* 12: 1879–1884.

39. Mehta A, Zhao JL, Sinha N, et al. (2015) The MicroRNA-132 and MicroRNA-212 cluster regulates hematopoietic stem cell maintenance and survival with age by buffering FOXO3 expression. *Immunity* 42: 1021–1032.