MicroRNA-187 inhibits pentylenetetrazol-induced neuronal apoptosis and alleviates development of epilepsy in epileptic rats by regulating SPRY1 expression

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Original Research Article

Abstract

**Purpose:** To explore the role of microRNA-187 on the pathological process of epilepsy.

**Methods:** The seizure score of epileptic rats was evaluated according to Racine’s scale. Real-time quantitative polymerase chain reaction (RT-qPCR) was performed to determine the expression levels of microRNA-187 (miR-187). Western blot technique was conducted to assess the expression levels of caspase 3, B-cell lymphoma-2 (BCL-2), and poly (ADP-ribose) polymerase (PARP) and activation of phosphatase and tensin homolog (PTEN)/PI3K/AKT cascade. Caspase 3 colorimetric assay kit was employed to evaluate the activity of caspase 3. Dual-luciferase reporter gene system was used to explore the regulating mechanisms of miR-187 and protein sprouty homolog 1 gene (SPRY1).

**Results:** The results showed that miR-187 was aberrantly downregulated in the hippocampus regions of pentylenetetrazol (PTZ)-treated rats compared to normal rats (p < 0.05). Furthermore, PTZ promoted caspase 3-dependent neuronal apoptosis by increasing the expression of pro-apoptosis protein PARP and decreasing the expression levels of BCL-2 in rats. On the other hand, overexpression of miR-187 downregulated SPRY1 as well as PTEN (p < 0.05), thereby activating the downstream PI3K/AKT signaling pathway. Notably, the effects of upregulated miR-187 on neuronal apoptosis and epilepsy development in PTZ-induced rats was reversed by the concomitant overexpression of SPRY1 (p < 0.05).

**Conclusion:** The results of this research show that overexpressed miR-187 alleviates the development of PTZ-induced neuronal apoptosis and epilepsy in epileptic rat models by regulating SPRY1 expression. These findings can hopefully be beneficial for the discovery of new therapeutic strategies for epilepsy treatment.

**Keywords:** Epilepsy, Neuronal apoptosis, miR-187, SPRY1
INTRODUCTION

Epilepsy seriously endangers the human health worldwide and degrades the life quality of human beings to a large extent [1]. Currently, there are several methods for epilepsy treatment and monitoring that help alleviate the symptoms of patients with epilepsy to some extent [2]. However, the therapeutic effects of these treatment methods are limited due to a lack of adequate knowledge regarding the underlying mechanisms of epilepsy development.

Neuronal apoptosis is closely associated with epilepsy, and targeting neuronal apoptosis has been shown to be a potential therapeutic method [3]. Nevertheless, the underlying mechanisms involved in regulating neuronal apoptosis in epilepsy are still not fully delineated.

Small non-coding RNA microRNA (miRNA) contains approximately 22 nucleotides on average [4], functions as a gene expression regulator, and participates in the pathogenesis of multiple diseases. Recently, miRNA was reported to be closely related to neurodegenerative disorders and the development of epilepsy [5]. For example, suppression of miR-141 has been shown to prevent against neuronal apoptosis in epilepsy [6]. Targeting of miR-21-5p protects rats against seizure damage by regulating the phosphatase and tensin homolog (PTEN)-mTOR signal pathway [7]. Also, miR-187 participated in the regulation of multiple diseases pathogenesis and progression [8].

In addition, by analyzing the miRNA expression profile of GSE49850, miR-187 was identified as playing a critical role in the pathogenesis of epilepsy [9]. Furthermore, the levels of miR-187 were closely related with epilepsy progression in both epileptic rat models and patients with temporal lobe epilepsy [10]. In spite of these reports and findings, the regulatory role of miR-187 in epilepsy development remains still unclear. Uncovering the underlying mechanisms might be helpful for the discovery of novel therapeutic agents for epilepsy treatment in clinic.

Protein sprouty homolog 1 gene (SPRY1) was crucial for regulating cell proliferation, migration, apoptosis and neurological functions [11]. In addition, sprouty homolog 1 is the downstream target of several miRNAs [12]. Hence this study aimed at investigating the role of miR-187-SPRY1 interaction in the development of PTZ-induced neuronal apoptosis and epilepsy.

EXPERIMENTAL

Establishment of animal models

Adult male rats weighing 260 to 300 g were maintained under standard experimental conditions of temperature of 25 °C, and humidity of 50–60 %. Animal feed and water were available ad libitum. The animal experiments were in compliance with the Guidelines for the Care and Use of Laboratory Animals” of the Institutional Animal Care and Use Committees (IACUCs) of the United States [24]. All animal studies were approved by the Animal Care and Use Committee of Guangxi Medical University (Approval Number: 2018-038-02). Pentylenetetrazol (PTZ) dissolved in artificial cerebrospinal fluid (ACSF) was administered according to the body weights of the rats (30 mg/kg) on alternate days for a maximum of 35 days. Then, each rat was placed in an isolated transparent box and observed individually for 30 min. Racine’s scale was used to evaluate the seizure score: 0 = no response; 1 = ear and facial cramping; 2 = myoclonic jerk without rearing; 3 = myoclonic jerk with rearing; 4 = change into the side position (e.g., clonic-tonic seizures); and 5 = change into the supine position, including generalized tonic-clonic convulsions.

Real-time quantitative polymerase chain reaction (RT-qPCR)

After treating rats with PTZ and successfully establishing the epileptic model, the hippocampus tissues of the rats were isolated and the TRLzol kit (Invitrogen Life Technologies, Carlsbad, California, USA) was employed to extract total RNA from the tissues. Reverse-transcription PCR analysis by iScript cDNA Synthesis Kit (Bio-rad, Hercules, CA, USA) and RT-qPCR analysis by HiScript II Q Select RT SuperMix (Vazyme Biotech Co. Ltd., Nanjing City, China) were purchased to evaluate gene expression. The primers of these genes are designed and listed in Table 1.

Western blot

The Western Blot technique was performed according to the previous studies. Briefly, Total proteins were acquired from the hippocampus tissues of the rats with or without PTZ treatment by using the RIPA lysis buffer purchased from Beyotime (Jiangsu, China). The bicinchoninic acid (BCA) kit (Pierce, Rockford, IL) was utilized to quantify the total proteins, which were then separated by using 10 % SDS/PAGE at 80 V for 2 h at 4 °C and transferred to polyvinylidene

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difluoride (PVDF) membranes (Millipore, Bedford, MA).

Table 1: Quantitative PCR primer sequences

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer sequence (strand)</th>
</tr>
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<tbody>
<tr>
<td>β-actin</td>
<td>Forward: 5'-CTCCATCTGGCCTCGGTG-T-3' Reverse: 5'-GCTGCTACCTCAACGTTCC-T-3'</td>
</tr>
<tr>
<td>U6</td>
<td>Forward: 5'-CTCGCTTTCGGACGACA-C-3' Reverse: 5'-AACGCTTCAAAATTCGCC-T-3'</td>
</tr>
<tr>
<td>miR-187</td>
<td>Forward: 5'-GGCUACAACACAGGCCGGCC-T-3' Reverse: 5'-GCCCGGGUCUCUGUGUAGGCC-T-3'</td>
</tr>
<tr>
<td>SPRY1</td>
<td>Forward: 5'-GCCTTCTTTGGATAGCCGTCAG-T-3' Reverse: 5'-TCATTGCTGCTCTTATGGCC-T-3'</td>
</tr>
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The PVDF membranes were probed with the antibodies including SPRY1 (1:1000, #ab75492, Abcam, Cambridge, UK), cleaved caspase 3 (1:1000, #ab32351, Abcam), GAPDH (1:5000, #ab181602, Abcam), PARP (1:500, #ab74290, Abcam), BCL-2 (1:1000, #ab32124, Abcam), PTEN (1:1000, #ab32199, Abcam), PI3K (1:1000, #ab32089, Abcam), AKT (1:1000, ab8805, Abcam) and p-AKT (1:1000, #ab1283, Abcam) overnight at 4 °C. After washing with TBST buffer (a mixture of Tris-buffered saline and Tween), the membranes were then incubated with the secondary antibody (1:5000, Santa Cruz Blotechnology, CA, USA) for 1 h at room temperature. The protein bands were then quantified by using a ECL Western Blot system purchased from Thermo Fisher Scientific (MA, USA).

Dual-luciferase reporter gene system

The fragments of human SPRY1 gene containing the binding sites were cloned into the pGL3-control vector by Ambion (Austin, TX) to generate SPRY1-WT vectors. The binding sites in in SPRY were then mutated by using a QuickChange site-directed Mutagenesis Kit purchased from Agilent Technologies (CA, USA) to generate SPRY1-MUT vectors.

Next, 293T cells were cultured under standard conditions and the above vectors were transfected into the cells. The Dual-Luciferase Reporter Gene System (Promega, Madison, WI, USA) was utilized to evaluate the luciferase activities of Firefly and Renilla after 48 h of transfection, which represented the regulating relationship between miR-187 and SPRY1.

Determination of caspase 3 activity

Tissues from the hippocampus of PTZ-treated rats were collected, and caspase 3 activity was evaluated by using a caspase 3 colorimetric assay kit purchased from Abcam (MA, USA). In brief, the tissues were collected, the lysis buffer solution was used to resuspend the above tissues. After that, the tissues were placed into the lysis buffer and a BCA kit (Pierce, IL, USA) was used to quantify the total proteins, and the supernatants were collected. Finally, the reaction buffer was then used to treat the supernatants and the activity of Caspase 3 was determined at the absorbance of 405 nm.

Statistical analysis

The data were collected and presented as mean ± standard deviation (SD) and analysed by SPSS 18.0 software. The Student’s t test for the comparison between two groups, and the one-way analysis of variance (ANOVA) for the comparisons among multiple groups. P < 0.05 means statistical significance.

RESULTS

Aberrant expression of miR-187 in the hippocampus tissues of PTZ-treated rats

To investigate the association between miR-187 and the development of epilepsy in rats, PTZ was used to establish epileptic rat models. The results showed that the epileptic rat models were successfully established based on the seizure score (Figure 1 A). In addition, the RT-qPCR results showed that miR-187 significantly downregulated in the hippocampus tissues of PTZ-treated rats compared to the control group (Figure 1 B).

Figure 1: The expression of miR-137 in the hippocampus tissues of PTZ-treated rats. (A) Seizure score of epileptic rats evaluated according to the Racine’s scale. (B) miR-187 level was evaluated by RT-qPCR in the hippocampus of epileptic rats (n = 6); **p < 0.05, *** p < 0.01
Effect of overexpressed miR-187 on neuronal apoptosis in PTZ-treated rat models

Based on the above findings, this study next explored the functions of miR-187 in PTZ-induced epileptic rats. Surprisingly, the results show that the development of PTZ-induced epilepsy in rats was attenuated by overexpression of miR-187 (Figure 2 A), which indicates that miR-187 might be a potential therapeutic agent for epilepsy. In addition, PTZ treatment downregulated miR-187 level and improved the activity of caspase 3 in neuronal cells in rat hippocampus, which were reversed by the concomitant overexpression of miR-187 (Figure 2 B and C). The expression levels of BCL-2 were downregulated, whereas the expressions of caspase 3 and PARP were upregulated by PTZ treatment in rat neuronal cells. The expressional changes of the above proteins were also reversed by the concomitant overexpression of miR-187 (Figure 2 D).

SPRY1 and its downstream PTEN/PI3K/AKT signal pathway are potential targets of miR-187

TargetScan software results showed that the 3’ UTR region of SPRY1 has miR-187 binding sites (Figure 3 A). In addition, the dual-luciferase reporter gene system results show that SPRY1 was targeted by miR-187 (Figure 3 A). Next, miR-187 mimics and inhibitor were successfully transfected into the hippocampus of normal rats (Figure 3 B). Further results show that SPRY1 was significantly decreased by overexpressed miR-187 and increased by downregulated miR-187 (Figure 3 C). The PTEN/PI3K/AKT cascade was proved to be the downstream target of SPRY1, which was also shown to be regulated by miR-187 (Figure 3 C). Besides, overexpressed miR-187 decreased the expression levels of PTEN and activated PI3K/AKT signal pathway, whereas knock-down of miR-187 had the opposite effects, which increased PTEN expression and inactivated PI3K/AKT signal pathway (Figure 3 C).

SPRY1 participated in the regulation of cell apoptosis and has also been proved to be regulated by miR-187 in the experiments. Hence, this study next explored whether miR-187 regulates neuronal apoptosis and epilepsy development in rats by regulating SPRY1. First, this study successfully overexpressed SPRY1 in the neuronal cells of the rats (Figure 4 A and B). The results show that the inhibiting effects of miR-187 on epilepsy development in PTZ-induced rat models were abrogated by synergistically overexpressing SPRY1 (Figure 4 C). In addition, the inhibiting effects of overexpressed miR-187 on PTZ-induced neuronal apoptosis were also reversed by SPRY1 overexpression (Figure 4 D and E).
Epilepsy is a chronic neurological disorder that greatly threatens human health [28], but the underlying mechanisms of the development of epilepsy are still unclear. Recent studies have showed that hippocampal neuronal apoptosis is critical for epilepsy [29]. Moreover, researchers reported that upregulation of Trem2 inhibits hippocampal neuronal apoptosis-mediated epilepsy via the PI3K/AKT signal pathway [29] and neuronal apoptosis-induced epilepsy in mice can be attenuated by sodium valproate [30]. These studies indicate that neuronal apoptosis is critical for epilepsy development and can be targeted to alleviate epilepsy.

MicroRNAs are small non-coding RNAs that contain an average of approximately 22 nucleotides [4] and function as gene regulators to modulate multiple genes expression. miRNAs have been reported to participate in the regulation of neuronal apoptosis [31] and epilepsy. MicroRNA-187 is a microRNA that has been reported to be closely related to epilepsy [9,10]. In addition, miR-187 has also been proved to regulate neuronal functions, including apoptosis, as well as neuronal development [33]. Hence, it is reasonable to speculate that miR-187 might be pivotal for neuronal apoptosis and epilepsy development. However, the detailed mechanisms are not fully delineated. The results showed that miR-187 was lowly expressed in the hippocampus regions of PTZ-treated rats. Furthermore, overexpressed miR-187 decreased PTZ-induced neuronal apoptosis and the development of epilepsy in rat models. These results are in accordance with previous studies [9,10] which also reported the protective role of miR-187 in epilepsy.

Sprotty homolog 1 is located on chromosome 4 of human beings and has been reported to participate in the regulation of various cell functions [34]. In addition, SPRY1 has been reported to regulate neuronal apoptosis [22,35] and predicted as the potential downstream target of miR-187 by using the TargetScan software, which indicated that miR-187 might alleviate PTZ-induced epilepsy in rats by targeting SPRY1. However, there are still no literature reports of a role of SPRY1 in the development of epilepsy.

Additional results show that overexpressed miR-187 downregulated the expression levels of SPRY1 and PTEN. Similarly, knock-down of miR-187 upregulated SPRY1 and PTEN expression levels, which indicates that SPRY1 and PTEN could be downregulated by overexpressed miR-187. Furthermore, PTEN inactivated PI3K/AKT cascade [36] and was crucial for the regulation of epilepsy development [37]. This results found that overexpressed miR-187 promoted the activation of PI3K/AKT signal pathway, which is in accordance with a previous study [38].

It is well known that miRNAs are pivotal for the regulation of cell functions [39]. This study found that miR-187 and SPRY1 are critical for neuronal apoptosis and epilepsy progression. Mechanistically, this study shows that upregulation of SPRY1 reverse the effects of overexpressed miR-187 on neuronal apoptosis and the development of epilepsy in PTZ-induced rat models. Taken together, miR-187 participates in the development of epilepsy by targeting SPRY1.

CONCLUSION

Overexpressed miR-187 inhibits neuronal apoptosis and the development of epilepsy in the hippocampus of PTZ-induced rat models by downregulating SPRY1 expression levels. These results can hopefully be helpful for the discovery of new therapeutic agents for epilepsy treatment in clinic.
DECLARATIONS

Acknowledgement

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Conflict of interest

No conflict of interest is associated with this work.

Contribution of authors

We declare that this work was done by the researchers listed in this article. All liabilities related with the content of this article will be borne by the authors. Limei Diao, Haichun Yu and Xianqiang Qiu designed all the experiments and revised the paper. Huan Li, Shunui Wang and Qian Yu performed the experiments, Huanxia Li, Ling Lu and Xianqiu Liao wrote the manuscript.

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REFERENCES


