

M₃ subtype of muscarinic acetylcholine receptor inhibits cardiac fibrosis via targeting microRNA-29b/beta-site app cleaving enzyme 1 axis

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Background: Previous studies have confirmed that choline exerts anti-fibrotic effect in the heart by activating the M_3 subtype of muscarinic acetylcholine receptor (M_3 receptor), but the mechanism remains to be clarified. MicroRNA-29b (miR-29b) plays an important role in the fibrotic process and can directly target collagen to resist myocardial fibrosis. This study investigated whether miR-29b is involved in the anti-fibrotic effect of activating M_3 receptor.

Methods: Proliferation of cardiac fibroblasts was induced by transforming growth factor (TGF)- β 1 *in vitro*. The expression of miR-29b in cardiac fibroblasts was detected by quantitative real-time reverse transcription polymerase chain reaction (qRT-PCR). Protein levels of collagens I, connective tissue growth factor (CTGF), α -smooth muscle actin (α -SMA) and beta-site app cleaving enzyme 1 (BACE1) were determined by Western blot analysis. Fibroblast-myofibroblast transition was identified by immunofluorescence staining. Proliferation and migration of cardiac fibroblasts as indicated by transwell and scratch assays.

Results: The expression of miR-29b decreased when treated with TGF-β1 (P=0.0389) and increased after choline stimulated (P=0.0001). Overexpression of miR-29b could reverse the high expression of collagen I (P<0.0001), α-SMA (P=0.0007), and CTGF (P=0.0038) induced by TGF-β1, whereas inhibition of miR-29b had a tendency to even further increase the expression of fibrosis markers. Meanwhile, inhibition of miR-29b could reverse the anti-fibrotic effect of choline, increasing the expression of collagen I (P=0.0040), α-SMA (P=0.0001), and CTGF (P=0.0185), and promoting the fibroblast proliferation and migration. Moreover, BACE1 protein level, increased after TGF-β1 treatment (P=0.0037) and reversed by overexpression of miR-29b (P=0.0493). Choline could reduce the increase of BACE1 induced by TGF-β1 (P=0.0264), and 4-diphenylacetoxy-N-methyl-piperidine methiodide (4-DAMP) increased the expression of BACE1 (P=0.0060). Furthermore, overexpression of BACE1 could reverse the protective effect of miR-29b in cardiac fibrosis, increasing the protein level of collagen I (P=0.0404).

Conclusions: The results suggested that M_3 receptor activation could exert cardioprotective effects in cardiac fibrosis by mediating miR-29b/BACE1 axis.

Keywords: Cardiac fibrosis; M₃ subtype of muscarinic acetylcholine receptor (M₃ receptor); microRNA-29b (miR-29b); beta-site app cleaving enzyme 1 (BACE1)

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Introduction

Background

Cardiovascular disease is one of the main causes of hospitalization and mortality, and is a large public health burden in the world (1). Cardiac fibrosis is a common pathophysiological process occurring along with a variety of diseases, including arrhythmia (2), heart failure (3) and sudden cardiac death (4). It is a pathological change from cardiac compensation to decompensation and can be caused by numerous factors. The fibroblast-myofibroblast transformation is a hallmark change in myocardial fibrosis, characterized by the secretion of extracellular matrix proteins including collagen deposition. In myofibroblasts, many endoplasmic reticulum form aggregates and express contractile proteins such as α -smooth muscle actin (α -SMA) and extracellular matrix proteins (5). Subsequently, myofibroblasts undergo migration and contraction, promoting the development of cardiac fibrosis. Several signaling pathways have been reported to be involved in the development of cardiac fibrosis, including transforming

Highlight box

Key findings

- Choline produces the anti-fibrotic effects through activating M₃ subtype of muscarinic acetylcholine receptor (M₃ receptor) to upregulate microRNA-29b (miR-29b) in cardiac.
- Beta-site app cleaving enzyme 1 (BACE1), a downstream target gene of miR-29b, is involved in the anti-fibrotic effects of activating of M, receptor.

What is known and what is new?

- Anti-fibrotic effects of miR-29b are known.
- The anti-fibrotic effects produced by activation of M₃ receptor is mediated by upregulation of miR-29b, and we identified BACE1 as a novel target for myocardial fibrosis that could be involved in this anti-fibrotic pathway.

What is the implication, and what should change now?

- This further illustrates the role of activating M₃ receptor in exerting anti-fibrotic effects and provides ideas for clinical treatment.
- Further explore the role of BACE1, the target identified in this study, in the progression of myocardial fibrosis and heart failure.

growth factor $-\beta$ (TGF- β) (6), mitogen-activated protein kinase (MAPK) (7), wingless/integrated (Wnt) (8), reninangiotensin-aldosterone system (RAAS) (9) and other signaling pathways.

Muscarinic acetylcholine receptors (M receptors) belong to the heterotrimeric G protein-coupled receptors (GPCR) family and can be bound by endogenous agonist acetylcholine (10), as well as choline (11), to excite parasympathetic nerve endings and act differently depending on the downstream effector. M3 subtype of muscarinic acetylcholine receptor (M3 receptor) is one of the M receptor subtypes, which is involved in the development of a variety of cardiovascular diseases. In ischemic arrhythmias, choline protects against myocardial ischemia-induced arrhythmias in rats through activating the M₃ receptor, mainly by reducing intracellular calcium overload (12). Activation of M₃ receptor can also prevent Angiotensin II-induced cardiac hypertrophy by inhibiting reactive oxygen species (ROS)-mediated p38 MAPK activation and regulating Ca2+-mediated calcineurin signal transduction pathway (13). The above results indicate the important role of M3 receptor in cardiomyocytes. Our pilot study revealed that M₃ receptor was expressed in cardiac fibroblasts in almost the same abundance as cardiomyocytes under basal conditions, and activation of M3 receptor inhibited cardiac fibrosis and reduced collagen secretion in mice by modulating the classical pathways of fibrosis, TGF-β1/Smad2/3 and p38 MAPK signaling pathways (14).

In this study, we used Kunming mice as the research subjects. The mice we used were characterized by easy rearing, high reproduction rate, easy accessibility, small inter-individual differences, and high disease resistance and adaptability, which improved the efficiency of our experiments and facilitated the observation of the results of parallel experiments. Meanwhile, they're close to humans in terms of biological evolution, with similar tissue organs and cellular functions and good genetic homology. Cells in primary culture have just been separated from living tissues and are closer to the morphology, structure and functional activities of the original tissues in the organism, and are able to represent signal transduction *in vivo* very closely, thus allowing for better cellular disease models to be established through primary cells.

Rationale and knowledge gap

 M_3 receptor belongs to the class of GPCRs that mediate many physiological responses in eukaryotes, and it is not clear whether new molecules are involved in this antifibrotic process after the activation of M_3 receptor, nor what its upstream and downstream factors are. We explored new regulatory factors and targets of this process in the present study.

Micro-RNAs (miRNAs) belong to a highly conserved, endogenous non-coding RNA family with a short length of approximately 22 nucleotides, and these molecules posttranscriptionally regulate gene expression by binding to the 3'-untranslated region of targeted messenger RNAs by partial complementarity (15). MicroRNA-29 (miR-29) forms a miRNA seed family, which consists of several members including miR-29a, miR-29b and miR-29c (16). Mature miR-29 is highly conserved in human, mouse and rat. MiR-29b is involved in a variety of pathways that regulate cardiovascular disease, and it may be useful for treatment or diagnostic of diseases (17). While a part of roles of miR-29b remain poorly understood, recent studies have provided important insights into the relevance of miR-29b to fibrotic diseases, such as the miR-29b plays a protective role in renal fibrosis (18), liver fibrosis (19), and pulmonary fibrosis (20). Moreover, miR-29b plays an important role in myocardial fibrosis (21).

 M_3 receptor can stimulate the nuclear factor kappa-B (NF- κ B) signaling pathway to inhibit the expression of miR-376b-5p and produce a cardioprotective effect (22). M_3 receptor overexpression shortens cardiac action potential duration through downregulation of miR-1 and prevents arrhythmias in mice with myocardial ischemia-reperfusion (23). The connection between M_3 receptor and miRNAs prompted us to propose that M_3 receptor might regulate miRNA to alter cardiac function and vice versa, miRNAs might act as downstream factors mediating the beneficial effects of M_3 receptor.

Due to the important role of miR-29b in the fibrotic process, and the unexplored relationship between the M_3 receptor and miR-29b, we investigated whether miR-29b was involved in the anti-fibrotic effect of M_3 receptor. Meanwhile, miRNAs usually act on targets to play their functions, whether a new target of miR-29b in myocardial fibrosis can be found is another purpose of this paper. It also provides new insights for clinical studies of myocardial fibrosis.

Objective

The goal of the present study was to explore the potential connection between M₃ receptor and miRNAs in the setting of cardiac fibrosis with a focus on miR-29b that is known to be a regulator of cardiac fibrosis. We assumed that the expression of miR-29b may be raised after activating the M₃ receptor, and the downstream target gene was controlled to affect fibrosis. We present this article in accordance with the ARRIVE reporting checklist (available at https://cdt. amegroups.com/article/view/10.21037/cdt-23-309/rc).

Methods

Neonatal mouse cardiac fibroblasts isolation and culture

Neonatal Kunming mice (1-3 days old) were purchased from the Animal Center of the Second Affiliated Hospital of Harbin Medical University (Harbin, China). The license number is SCXK(HeilongJiang)2019-001. The study was reviewed and approved by the Ethics Committee of the Second Affiliated Hospital of Harbin Medical University (Approval number: Ky2016-103. Approval time: 9 March 2016), in compliance with the "Laboratory animal-Guideline for ethical review of animal welfare" (GB/T 35892-2018) national guidelines for the care and use of animals. A protocol was prepared before the study without registration. We strive to use fewer animals required by experiments by improving experimental methods, and obtain more data with the minimum quantity of animals. Moreover, we painlessly execute animals and then take their hearts for in vitro studies, which is a relative substitute for live animals. Throughout the experiment, we treat experimental animals kindly, respect animal life, and thank animals for the effort for our human scientific research. No exclusion criteria was set during the experiment. The hearts of randomly selected neonatal mice were dissected and cut into 2-3 pieces and placed in Dulbecco's modified Eagle's medium (DMEM, HyClone, USA). The preparation was transferred into 50 mL centrifuge tubes, followed by the addition of prewarmed 0.25% trypsin for cleaning and digestion. The collected cell suspension was filtered, and cells were obtained by centrifugation at 1,000 r/min for 5 min. The cells were resuspended in DMEM containing 10% fetal bovine serum (FBS) and 5% penicillin/ streptomycin and then incubated in Petri dishes (60 mm) or other specifications of culture plates. The cells were

then cultured under a condition of 5% CO₂ and 95% air at 37 °C for 1.5 h to allow cardiac fibroblasts to adhere to the bottom of the Petri dishes and leave the isolated cardiac myocytes and other cells suspended. The cardiac fibroblasts were cultured under the same conditions for 48 h before being examined for morphology and density. The cells were incubated with 3 nM of 4-diphenylacetoxy-N-methyl-piperidine methiodide (4-DAMP) (Abcam, USA), 1 mM of choline (Sigma, USA) or 20 ng/mL of TGF-β1 (PeproTech, USA) as to be specified in the appropriate sections as described previously (14).

Transfection procedures

Prior to transfection, the cardiac fibroblasts were confirmed to be in good condition under a microscope and devoid of contamination, followed by incubation in fresh culture medium for 4-6 h. The cells were then transfected with 50 nM of miR-29b mimics or inhibitor using Opti-MEM (Invitrogen, USA) and X-treme GENE siRNA transfection reagent (Roche, Penzberg Germany) for 6 h according to the manufacturer's protocols. Next, the cells were kept in the medium containing 10% bovine serum. In separate experiments, cells were transfected with 1 µg/mL beta-site app cleaving enzyme 1 (BACE1) plasmid or empty plasmid (pcMV3-C-GFPSpark) through Lipofectamine 2000 (Invitrogen, USA) and X-treme for 6 h. The transfected cardiac fibroblasts were incubated for 24 h and then treated with drugs according to the previous experimental procedure (14). Briefly, the cardiac fibroblasts were pretreated with 3 nM 4-DAMP in the presence of 1 mM choline for 1 h and then incubated with 20 ng/mL TGF-β1 for 48 h. The sequence of miR-29b mimics was 5'-AACACUGAUUUCAAAUGGUGCUA-3' and the sequence of miR-29b inhibitor was 5'-UAGCACCAUUUGAAAUCAGUGUU-3'. The negative control sequence was 5'-UCUACUCUUUCUAGGAGGUUGUGA-3' and all these constructs were purchased from General Biosystems, Inc. (Anhui, China). BACE1 plasmid and empty plasmid were constructed by Sino Biological Inc. (Beijing, China).

Immunofluorescence staining

Cardiac fibroblasts were inoculated onto sterile coverslips in 24-well culture plates and treated accordingly as described in the transfection procedure. After treatment, coverslips were washed three times with cold phosphate buffered saline (PBS) and fixed with 4% paraformaldehyde for 30 min. The coverslips were washed three times again with cold PBS and then permeabilized with 4 mL PBS including bovine serum albumin 0.04 g and Triton X-100 10 µL for 20 min. Goat serum (Solarbio, Beijing, China) was mixed with PBS in an equal volume to block, and the cardiac fibroblasts were incubated at 37 °C for 1 h. Subsequently, the cells were incubated with collagen I antibody (Proteintech, Wuhan, China, 1:250) or α-SMA antibody (Abcam, USA, 1:250) at 4 °C overnight, followed by incubation with fluorescein isothiocyanate (FITC)-conjugated goat antirabbit IgG secondary antibody (ZSGB-Bio, China, 1:500) in the dark at room temperature for 1 h. DAPI (Beyotime, Haimen, China) was added to stain the nuclei for 5 min after washing with PBS, and all steps were protected from light. Images were taken with a laser scanning confocal microscope (Olympus, Fluoview1000, Tokyo, Japan).

Western blot

Cells were rinsed with PBS and lysed using Radio Immunoprecipitation Assay Lysis buffer (RIPA) (Beyotime, Jiangsu, China) supplemented with protease inhibitor mixture. Protein concentration was determined using the Bicinchoninic Acid (BCA) Protein Assay Kit (Beyotime, Shanghai, China) according to the manufacturer's instructions. Configured the system with RIPA and sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) 5× Loading Buffer (Beyotime, Jiangsu, China). The protein sample (70 µg) was fractionated by SDS-PAGE (10% polyacrylamide gel) and transferred to a nitrocellulose membrane. The nitrocellulose membranes were blocked in 5% nonfat milk for 2 h and then incubated with primary antibodies at 4 °C in the refrigerator overnight. Primary antibodies included collagen I (1:1,000), α-SMA (1:1,000), CTGF (1:500), BACE1 (1:1,000), and glyceraldehyde-3phosphate dehydrogenase (GAPDH, 1:1,000). Antibodies against collagen I and CTGF were purchased from Proteintech (Wuhan, China). Antibody against α-SMA was purchased from Abcam (USA) and antibody against BACE1 was purchased from ABclonal (Hubei, China). Antibody against GAPDH was purchased from ZSGB-Bio (China). The membranes were washed three times with PBS-Tween (0.5%), secondary antibodies were used to incubate the membranes in the dark at room temperature for 1 h, and then scanned to capture blot band images with Imaging System (LI-COR Biosciences, Lincoln, NE, USA).

Quantitative real-time reverse transcription polymerase chain reaction (qRT-PCR)

Total RNA from cardiac fibroblasts was extracted using TRIzol™ Reagent (Thermo, USA) according to the manufacturer's protocol. RNA concentration was determined by Nanodrop 2000 and was reverse transcribed into complementary DNA. Quantification of miR-29b was done by LC480 Real-time PCR system (Roche, USA) using SYBR Green PCR Master Mix (Roche, USA) PCR cycling protocol: 95 °C for 10 min, 40 cycles at 95 °C for 15 s, 60 °C for 30 s and 72 °C for 30 s. U6 was used as an internal control and the amount of miR-29b was calculated according to $2^{-\Delta\Delta Ct}$. The sequences of primers used in this study were: miR-29b, forward: 5'-GCGGTAGCACCATTTGAAATCAGT-3' and reverse: 5'-ATCCAGTGCAGGGTCCGAGG-3'; U6 forward: 5'-GCTTCGGCACATATACTAAAAT-3' and reverse: 5'-CGCTTCACGAATTTGCGTGTCAT-3'.

Scratch test

Cardiac fibroblasts were plated in a 6-well plate, incubated in DMEM containing 10% FBS and 5% penicillin/ streptomycin. The cells were cultured under a condition of 5% CO_2 and 95% air at 37 °C for 48 h before the culture medium was changed to DMEM for 4–6 h. The cells were wounded by scratching using a 10 µL tip, and non-adherent cells were washed with culture medium. Transfection reagent, choline and TGF- β 1 treatment were carried out as described in transfection procedures and photographs were taken under a microscope immediately. Photographs were taken at 24 and 48 h after TGF- β 1 treatment. The percentages of wound closure area of cell scratches over time were determined using ImageJ software.

Transwell migration assay

Transwell chambers with 8- μ m pore size membranes (Corning Life Sciences, Lowell, MA, USA) were used to detect the migration of cardiac fibroblasts. The transfected cells were harvested and reconstituted in serum-free buffer at 5×10^5 cells/mL, and 100 μ L of the cell suspension was taken and added to the chambers. Meanwhile, 600 μ L of DMEM containing 10% FBS, TGF- β 1, and choline was added to the lower chamber. After incubation, the chamber was cleaned with PBS and fixed with 4% paraformaldehyde for 30 min, followed by staining with crystal violet staining

solution for 20 min. The chambers were examined under a microscope.

Statistical analysis

The different stages of the experiment (allocation period, experiment implementation) were handled by different person. Anyone who randomized the experiment by labeled numbers only, without knowing what kind of treatment each group received. Results the evaluation and data analysis results were entered according to the corresponding numbers of each group by another person. GraphPad Prism 9.0 was used for data analysis. The experimental data are presented as the mean ± SEM. One-way analysis of variance (ANOVA) followed by a *post hoc* Tukey test for multiple group comparisons. Statistical significance was considered when a value of P was <0.05.

Results

Activation of M₃ receptor by choline upregulates miR-29b and downregulates fibrotic markers in cardiac fibroblasts

Previous study has demonstrated that choline possesses anti-fibrotic properties through inhibiting the TGF-β1/ Smad and p38 MAPK pathways upon activating M₃ receptor (14). However, how M3 receptor activation leads to fibrosis inhibition remains elusive. To this end, we detected the expression of miR-29b in cardiac fibroblasts with different treatment conditions. The results showed the level of miR-29b was downregulated in cardiac fibroblasts induced by TGF-β1 (P=0.0389), whereas upregulated after further addition of choline (P=0.0001), and 4-DAMP, a competitive antagonist of M₃ receptor reversed the effect of choline (P=0.0013) (Figure 1A). Meanwhile, we detected that the administration of choline and 4-DAMP under physiological conditions did not affect the expression of miR-29b (Figure S1), which may be related to other mechanisms. We also detected the expression of M_3 receptor when transfected with miR-29b mimics and miR-29b inhibitor. Transfection efficiency of miR-29b mimics (Figure S2A) and miR-29b inhibitor (Figure S2B) were shown in Figure S2. There was no effect on the protein level of M_3 receptor when transfected with miR-29 mimics or miR-29b inhibitor (Figure S3), so we suggested that miR-29b may play a role in myocardial fibrosis as a downstream of the M3 receptor. These data suggested that miR-29b might be involved in the anti-fibrotic effect afforded by choline via activating the

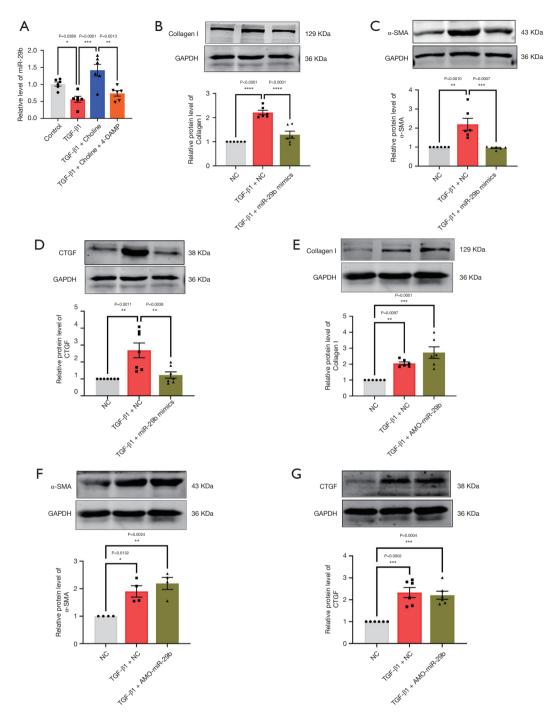


Figure 1 The anti-fibrotic effects of miR-29b and it participates in the role of activating M, receptor in cardiac fibroblasts. (A) qRT-PCR was performed to detect the relative expression of miR-29b in TGF-β1-induced cardiac fibroblasts when choline and 4-DAMP stimulation (n=6). (B-G) Relative protein level of collagen I (B, n=6; E, n=6), α-SMA (C, n=6; F, n=4), and CTGF (D, n=7; G, n=6) in cardiac fibroblasts detected by western blot. ****, P<0.001; ***, P<0.001; ***, P<0.01; *, P<0.05. GAPDH was included as a control. The data are expressed as the mean ± SEM. miR-29b, microRNA-29b; TGF-β1, transforming growth factor beta 1; 4-DAMP, 4-diphenylacetoxy-N-methylpiperidine methiodide; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; NC, negative control; α-SMA, α-smooth muscle actin; CTGF, connective tissue growth factor; AMO-miR-29b, miR-29b inhibitor; M3 receptor, M3 subtype of muscarinic acetylcholine receptor; qRTPCR, quantitative real-time reverse transcription polymerase chain reaction; SEM, standard error of mean.

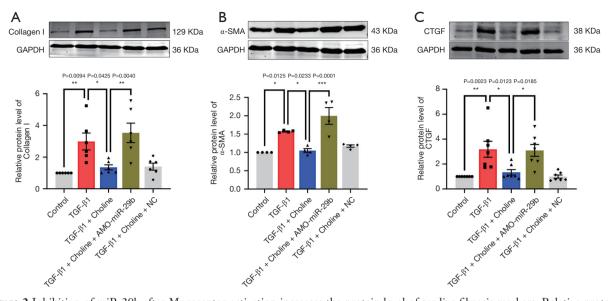


Figure 2 Inhibition of miR-29b after M_3 receptor activation increases the protein level of cardiac fibrosis markers. Relative protein level of collagen I (A, n=6), α-SMA (B, n=4), and CTGF (C, n=7) in cardiac fibroblasts detected by western blot. ***, P<0.001; **, P<0.01; *, P<0.05. GAPDH was included as a control. The data are expressed as the mean \pm SEM. GAPDH, glyceraldehyde-3-phosphate dehydrogenase; TGF-β1, transforming growth factor beta 1; AMO-miR-29b, miR-29b inhibitor; NC, negative control; miR-29b, microRNA-29b; α-SMA, α-smooth muscle actin; CTGF, connective tissue growth factor; M_3 receptor, M_3 subtype of muscarinic acetylcholine receptor; SEM, standard error of mean.

M₃ receptor when myocardial fibrosis occurs.

We then overexpressed or inhibited the expression of miR-29b in cardiac fibroblasts to detected the expression of cardiac fibrosis markers such as collagen I, α -SMA and CTGF. The primary endpoint of this study was defined as the protein level of collagen I, α -SMA and CTGF. Compared with the negative control group (NC), the TGF- β 1 group showed high expression of collagen I (P<0.0001), α -SMA (P=0.0010), and CTGF (P=0.0011), whereas miR-29b overexpression reversed such an abnormal expression upregulation of collagen I (P<0.0001), α -SMA (P=0.0007), and CTGF (P=0.0038) (*Figure 1B-1D*), and at the same time, the inhibition of miR-29b expression did not have such an effect on the expression of collagen I, α -SMA, and CTGF (*Figure 1E-1G*).

MiR-29b mediates the anti-fibrotic action of M_3 receptor activation in cardiac fibroblasts

The above results suggested that miR-29b might act as an anti-fibrotic miRNA downstream of the M_3 receptor. To clarify this thought, we performed the following experiments. The primary endpoint of this study was defined as the protein level of collagen I, α -SMA and

CTGF. First, we demonstrated that choline reversed the overproduction of collagen I protein in the presence of TGF- β 1 (P=0.0425) (*Figure 2A*). Notably, the anti-fibrotic effect of choline was counteracted by miR-29b inhibitor in cardiac fibroblasts (P=0.004), but not by the NC construct (P>0.99) (*Figure 2A*). Similar results were observed with α -SMA (P=0.0001) and CTGF proteins (P=0.0185) (*Figure 2B*,2C). This further confirmed the importance of miR-29b in the anti-fibrotic effect of M_3 receptor.

Second, we used immunofluorescence to detect the changes of expression and subcellular distribution of collagen I and α -SMA in cardiac fibroblasts under different treatment conditions. The results uncovered that the expression of collagen I and α -SMA of cardiac fibroblasts which stimulated by TGF- β 1 were increased, as indicated by the enhanced fluorescence intensity (P<0.0001). α -SMA served as an indicator of fibroblast-myofibroblast transition, demonstrating that TGF- β 1 can promote the conversion of fibroblasts to myofibroblasts. Choline abrogated TGF- β 1-induced upregulation of collagen I and α -SMA (P<0.0001), and these beneficial effects of choline were mitigated after transfection with miR-29b inhibitor (P<0.0001) (*Figure 3A-3D*). We also verified whether overexpression of miR-29b could affect the level of collagen I and α -SMA

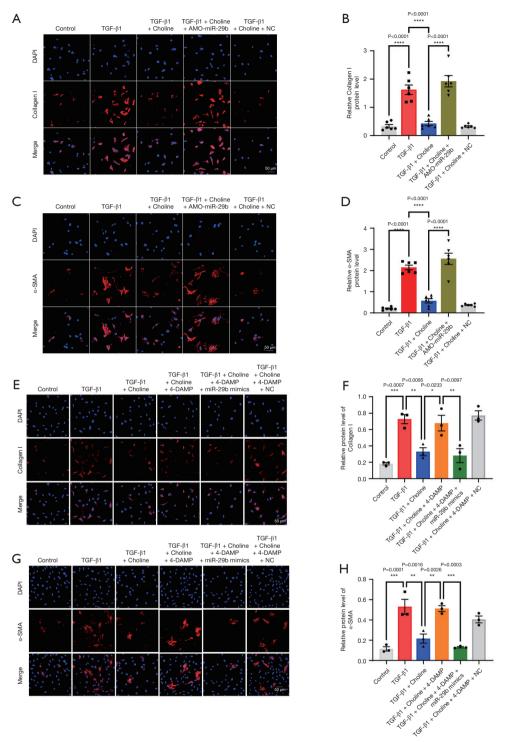


Figure 3 Inhibition of miR-29b reverses the anti-fibrotic effects of activating M_3 receptor and promotes the expression of collagen I and α-SMA. (A,E) Relative collagen I protein level in cardiac fibroblasts detected by immunofluorescence. Scale bar =50 μm. (B,F) Statistical results of immunofluorescence of collagen I (B, n=6; F, n=3). (C,G) Relative α-SMA protein level in cardiac fibroblasts detected by immunofluorescence. Scale bar =50 μm. (D,H) Statistical results of immunofluorescence of α-SMA (D, n=6; H, n=3). ****, P<0.001; ***, P<0.001; ***, P<0.001; **, P<0.05. The data are expressed as the mean ± SEM. TGF-β1, transforming growth factor beta 1; AMO-miR-29b, miR-29b inhibitor; miR-29b, microRNA-29b; NC, negative control; α-SMA, α-smooth muscle actin; 4-DAMP, 4-diphenylacetoxy-N-methylpiperidine methiodide; M_3 receptor, M_3 subtype of muscarinic acetylcholine receptor; SEM, standard error of mean.

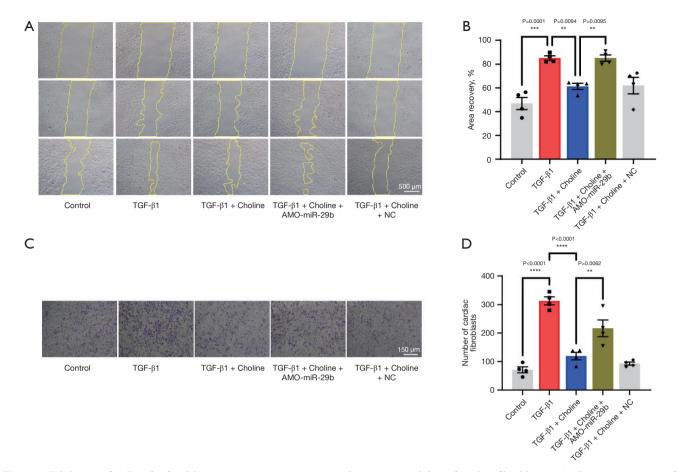


Figure 4 Inhibition of miR-29b after M_3 receptor activation promotes the migration ability of cardiac fibroblasts. (A) The migration rate of cardiac fibroblasts detected by scratch test. Scale bar =500 µm. (B) Statistical results about the migration rate of cardiac fibroblasts, n=4. (C) The migration ability of cardiac fibroblasts was detected by transwell (crystal violet staining). Scale bar =150 µm. (D) Statistical results about migration ability of cardiac fibroblasts, n=4. ****, P<0.0001; ***, P<0.001; ***, P<0.01. The data are expressed as the mean \pm SEM. TGF- β 1, transforming growth factor beta 1; AMO-miR-29b, miR-29b inhibitor; miR-29b, microRNA-29b; NC, negative control; M_3 receptor, M_3 subtype of muscarinic acetylcholine receptor; SEM, standard error of mean.

when M_3 receptor was inhibited by 4-DAMP under model conditions. It could be seen that in agreement with previous results, 4-DAMP reversed the effects of choline and increased the expression of collagen I (P=0.0233) and α -SMA (P=0.0026). These were counteracted by miR-29b mimics, which decreased the level of collagen I (P=0.0097) and α -SMA (P=0.0003) (*Figure 3E-3H*).

Third, activation of fibroblasts promotes cell proliferation and collagen secretion, so we examined the effects of miR-29b and M₃ receptor on the proliferative and migratory capacity of cardiac fibroblasts by two different methods. In the first set of experiments, we used the scratch test to measure the migratory ability of cardiac fibroblasts.

We found that TGF- β 1 promoted the migration of cardiac fibroblasts (P=0.0001), which was inhibited by choline (P=0.0094), indicating the anti-fibrotic effect of the M_3 receptor. Transfection of miR-29b inhibitor eliminated the protective effect of choline (P=0.0095) and aggravated the migration of cardiac fibroblasts (*Figure 4A*,4*B*). In the second set of experiments, we used the transwell assay to measure the migratory and invasive ability of cardiac fibroblasts. The results showed that the migration/invasion suppressing effects of choline were abolished following a knockdown of miR-29b by its antisense inhibitor (P=0.0062) (*Figure 4C*,4*D*). These consistent results suggested that miR-29b inhibitor promoted cardiac fibrosis even while the

M₃ receptor had been activated.

BACE1 participates in the anti-fibrotic effect by activating M3 receptor

It has been reported that miR-29b can fight against fibrotic diseases by directly targeting collagen I and collagen III (24), and in the present study, we screened for new targets of miR-29b against fibrosis. Four online prediction websites for miRNA targets were used, including miRDB (http:// www.mirdb.org/), miRTarBase (https://mirtarbase.cuhk. edu.cn/), TargetScan (http://www.targetscan.org/), and PicTar (https://pictar.mdc-berlin.de/), and Venn diagrams were drawn. A total of 19 target genes were identified common to the four databases mentioned above. Through literature review, we screened the target genes based on their expression, abundance in the heart and a good conservation among species. This procedure led us to focus on BACE1 as a new target gene of miR-29b for studying its anti-fibrotic properties (Figure 5A). Sequence alignment revealed multiple binding sites at the 3' non-coding region of BACE1 gene for miR-29b seed site (Figure 5B).

We next experimentally established the targeting relationship between miR-29b and BACE1. Western blot analysis showed that the protein level of BACE1 was increased in the presence of TGF-β1 (P=0.0037) but was reduced after transfection of miR-29b mimics (P=0.0493) (Figure 5C). Meanwhile, inhibition of miR-29b did not affect the effects of TGF-β1 (Figure 5D). We further detected the expression of BACE1 in choline or 4-DAMP treated cardiac fibroblasts. We found that TGF-β1-induced increase in the protein level of BACE1 was mitigated in cells treated with choline (P=0.0264), on the other hand, 4-DAMP, a competitive antagonist of M3 receptor, reversed the effect of choline (P=0.0060) (Figure 5E). We then overexpressed BACE1 by transfecting the cells with the plasmid carrying BACE1 for overexpression. The transfection efficiency of the plasmid was shown in Figure 5F. And we found that overexpression of BACE1 reduced the level of miR-29b (P=0.0010) (Figure 5G). Moreover, overexpression of BACE1 eliminated the anti-fibrotic effect elicited by miR-29b mimics (P=0.0404) and restored the expression of collagen I (Figure 5H). In all cases, the empty plasmid as a negative control for BACE1 overexpression did not produce any significant changes to the cardiac fibroblasts.

Discussion

Our previous results showed that choline suppresses cardiac fibroblast proliferation and collagen secretion via activating M_3 receptor, and this effect can be ascribed to the inhibition of the TGF- β 1/Smad and p38 MAPK pathways. However, its specific molecular mechanism has not been elucidated, and the present work was designed to shed light on it.

Key findings

The findings of the present study revealed a novel mechanism by which choline produces the anti-fibrotic action through activating the M_3 receptor to upregulate miR-29b, thereby repressing the expression of the downstream target gene BACE1: choline $\rightarrow M_3$ receptor activation $\uparrow \rightarrow \text{miR-29b}$ upregulation $\uparrow \rightarrow \text{BACE1}$ downregulation $\downarrow \rightarrow \text{collagen I} \downarrow \rightarrow \text{fibrosis inhibition} \downarrow$.

Strengths and limitations

In this study, we proposed for the first time that the activation of M_3 receptor can upregulate the expression of miR-29b to exert anti-fibrotic effect, and we identified a target, BACE1, that has never been reported in myocardial fibrosis, which supplemented the mechanism of the anti-fibrotic effects of M_3 receptor and further provided new evidence for the function of M_3 receptor in myocardial fibrosis.

However, there are some limitations in the present study. For example, how the M_3 receptor upregulates miR-29b expression is unclear. It is reported that TGF- β -Smad3-miR-29 isoforms may mainly contribute to the progression of cardiac fibrosis due to Cyclosporine A exposure (25). Previous studies confirmed that the activation of M_3 receptor by choline can inhibit the TGF- β /Smad signaling pathway, so we speculated that the fibrotic signaling pathway TGF- β /Smad may act as a bridge between M_3 receptor and miR-29b. This still needs to be confirmed by further experiments.

Furthermore, BACE1 was involved in the anti-fibrotic effects of increasing the expression of miR-29b caused by the activation of M₃ receptor, and overexpression of BACE1 eliminated the role of miR-29b mimics in cardiac fibrosis. However, it is still unclear whether BACE1 can directly affect the level of collagen I, or whether we can directly act on BACE1 to exert anti-fibrotic effects. This problem is

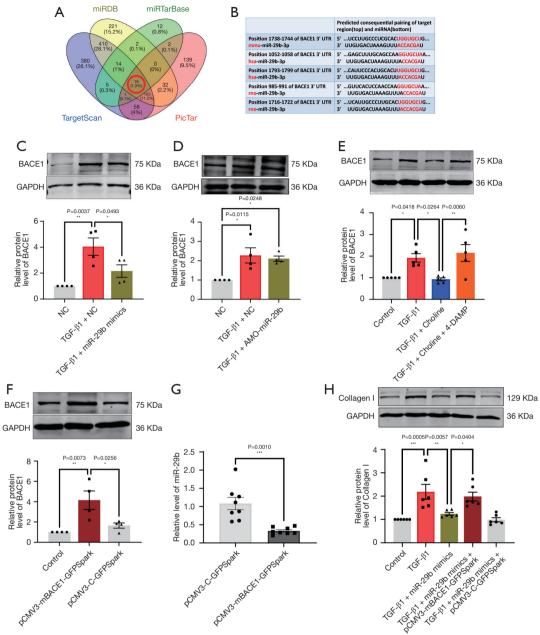


Figure 5 MiR-29b participates in the anti-fibrotic effects after M₃ receptor activation through targeting BACE1. (A) Screening target genes of miR-29b by online prediction websites: miRDB, miRTarBase, TargetScan, and PicTar. (B) The binding sites of miR-29b and BACE1 in different species. (C-F) Relative protein level of BACE1 in cardiac fibroblasts detected by western blot (C, n=4; D, n=4; E, n=5; F, n=4) (G) qRT-PCR assay was used to examine the expression of miR-29b in the presence of BACE1 overexpression, n=8. (H) Relative protein level of collagen I in cardiac fibroblasts detected by western blot, n=6. ***, P<0.001; **, P<0.01; *, P<0.05. The data are expressed as the mean ± SEM. Student's *t*-test was used for two-group comparisons. BACE1, beta-site app cleaving enzyme 1; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; NC, negative control; TGF-β1, transforming growth factor beta 1; miR-29b, microRNA-29b; AMO-miR-29b inhibitor; 4-DAMP, 4-diphenylacetoxy-Nmethylpiperidine methiodide; M₃ receptor, M₃ subtype of muscarinic acetylcholine receptor; qRT-PCR, quantitative real-time reverse transcription polymerase chain reaction.

also one of the limitations of this article. We plan to further explore the relationship between BACE1 and cardiac fibrosis in future experiments. Hopefully, it will provide a new strategy for the treatment of cardiac fibrosis.

Comparison with similar researches

In previous studies, we found that M_3 receptor are not only expressed in cardiomyocytes, but also in cardiac fibroblasts, and exert anti-fibrotic effects when activated by choline. In this study, we further confirmed that the role of M_3 receptor in myocardial fibrosis. But on top of that, we discovered a new member involved in this process, which is miR-29b. Activation of the M_3 receptor by choline could promote the expression of miR-29b, and we also found another member, BACE1, a new target of miR-29b in myocardial fibrosis. These results complement our previous research and provide new insights.

Explanations of findings

MiRNAs regulate a variety of pathological processes and have become a research hotspot in the prevention and treatment of diseases in recent years. We screened the miRNAs related to fibrosis and selected miR-29b as the research object to explore whether miR-29b was involved in the anti-fibrotic effect produced by M₃ receptor activation. We found that increased miR-29b expression exerted an anti-fibrotic effect in cardiac fibroblasts with exposure to TGF-β1, a master fibrosis inducer, whereas inhibition of miR-29b antagonized the anti-fibrotic effect of M₃ receptor. We identified BACE1 as a new target gene for miR-29b in cardiac fibroblasts. The results showed that choline downregulated the protein level of BACE1 by altering the expression of miR-29b, whereas 4-DAMP upregulated its protein level, suggesting that BACE1 was a downstream component of the M3 receptor signaling connecting to cardiac fibrosis. These findings delineated a new molecular/ signaling mechanism underlying the protective actions of the M3 receptor and a new mechanism for the development of cardiac fibrosis with miR-29b/BACE1 as new mediators for fibrogenesis.

Previous reports suggested that the M_3 receptor is expressed in cardiomyocytes and that the activation of the M_3 receptor had a protective effect on arrhythmia (12), myocardial hypertrophy (13), and sudden cardiac death (23). We demonstrated that the M_3 receptor level in

cardiac fibroblasts was as high as in cardiomyocytes and M₃ receptor activation by choline suppressed cardiac fibrosis in mice by inhibiting the P38 MAPK pathway (14). Choline is a precursor for the synthesis of acetylcholine, a non-specific agonist of the M3 receptor. Since 4-DAMP is a specific antagonist of M₃ receptor subtype and it counteracts the effect of choline, the effects of choline observed in our experiments can be attributed to M₃ receptor activation. However, pharmacological studies suggest that a highcholine diet aggravates cardiac dysfunction, inflammation, and fibrosis in a mouse model of heart failure (26). We consider that such different effects of choline are related to the concentration under study. As our previous results, the anti-fibrotic effect of choline decreased while its concentration increased, suggesting that higher concentrations of choline may bring about the opposite effect, which deserves further investigation.

In this study, we found that miR-29b negatively regulated BACE1 in cardiac fibroblasts and might mediate the anti-fibrotic effect of M3 receptor. BACE1 is a candidate biomarker for Alzheimer's disease (27). It has been reported that the expression of BACE1 is elevated in both patients and mice with ischemic heart failure and promotes the expression of its antisense transcript BACE1-AS to increase the endothelial cell apoptosis (28). BACE1 can also play a role in ischemic heart disease (29). Three types of myocardial fibrosis have been identified: replacement fibrosis from tissue necrosis, reactive fibrosis from myocardial stress, and infiltrative interstitial fibrosis from progressive deposition of non-degradable material such as amyloid (30). This provides a conjecture for the role of BACE1 in myocardial fibrosis. Our experimental results in this article showed that the expression of BACE1 was increased in cardiac fibroblasts with TGF-β1, which was reversed after transfection with miR-29b mimics. Some studies have reported the targeting relationship between miR-29b and BACE1 in Alzheimer's disease, thus, in the present study we test the relationship in myocardial fibrosis by western blot rather than dual-luciferase reporter assay. We identified the involvement of the miR-29b/BACE1 axis in the anti-fibrotic effect of the activation of M₃ receptor, which is a new molecular mechanism of the activation of M₃ receptor to inhibit myocardial fibrosis. However, the role of BACE1 in cardiac fibrosis remains to be further investigated. ST β-galactoside alpha-2,6-sialyltransferase 1 (ST6GAL1) is a type II membrane protein that is commonly localized in the Golgi apparatus, and has been shown to regulate matrix metalloproteinases (MMPs) and vascular endothelial growth

factor (VEGF) in lung cancer (31). BACE1 can cleave and release ST6GAL1 from the trans-Golgi into the secretory pathway to regulate the secretion of inflammatory factors in chronic obstructive pulmonary disease (32,33). Moreover, we reviewed the literature and found that NKX-COUP-TFII composite elements (NCCE) in regulatory regions of Madcam1 and St6gal1 are involved in morphogenesis of the heart (34). Meprin β contributes to collagen deposition in lung fibrosis (35), and meprin β complements the role of BACE1 in Alzheimer's disease (36). These have the potential to play a role in the heart, but follow-up experiments are needed to confirm this.

Implications and actions needed

The experimental results in this study, as well as previous studies of M_3 receptors in the heart, suggest that overexpression of M_3 receptors may rescue a number of cardiac diseases, such as myocardial hypertrophy, ischemic heart disease, and myocardial fibrosis. In clinical trials, it is also possible to try to improve the level of M_3 receptor to treat heart diseases. However, myocardial fibrosis is usually a progressive disease accompanied by other diseases, to achieve the therapeutic effect of overexpression of M_3 receptor, it is necessary to take into account the effect of overexpression M_3 receptor on other diseases. And there are multiple subtypes of M muscarinic receptors, which also need to be considered when performing therapeutic administration.

The ultimate goal of studying miRNAs is to translate them into clinical applications. MiRNAs can be stabilized in serum and plasma of humans and animals (37). Thus, serum miRNAs can be used as markers for disease diagnosis. In our study, changes of miR-29b in human serum might serve as a marker of myocardial fibrosis disease from progression to remission. In recent years, RNA interference (RNAi) therapeutic technology has emerged based on the study of miRNAs, and some studies have shown that plants can be carriers of certain therapeutic miRNAs (38). MiR-29b could likewise be further investigated in myocardial fibrosis by drawing on these studies. These are all elements that deserve to be explored further.

Conclusions

This study revealed the mechanism by which choline produces the anti-fibrotic action through activating the M_3 receptor to upregulate miR-29b, thereby inhibiting the

expression of the downstream target gene BACE1.

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Footnote

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was reviewed and approved by the Ethics Committee of the Second Affiliated Hospital of Harbin Medical University (Approval number: Ky2016-103. Approval time: 9 March 2016), in compliance with the "Laboratory animal—Guideline for ethical review of animal welfare" (GB/T 35892-2018) national guidelines for the care and use of animals.

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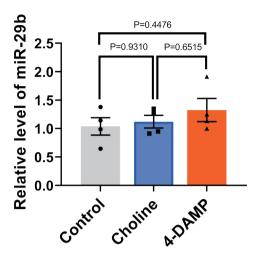


Figure S1 Addition of choline and 4-DAMP alone does not affect the expression of miR-29b. qRT-PCR was performed to detect the relative expression of miR-29b in cardiac fibroblasts, n=4. The data are expressed as the mean ± SEM. 4-DAMP, 4-diphenylacetoxy-N-methylpiperidine methiodide; miR-29b, microRNA-29b; qRT-PCR, quantitative real-time reverse transcription polymerase chain reaction.

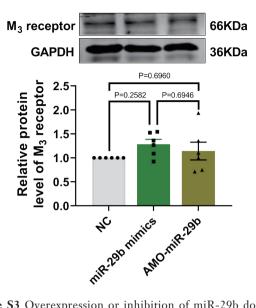


Figure S3 Overexpression or inhibition of miR-29b does not affect the protein level of M_3 receptor. Relative protein level of M_3 receptor in cardiac fibroblasts detected by western blot, n=6. GAPDH was included as a control. The data are expressed as the mean \pm SEM. NC, negative control; AMO-miR-29b, miR-29b inhibitor; miR-29b, microRNA-29b; M_3 receptor, M_3 subtype of muscarinic acetylcholine receptor; GAPDH, glyceraldehyde-3-phosphate dehydrogenase.

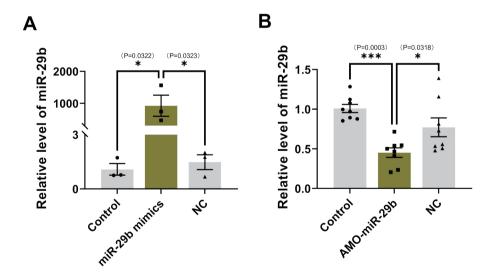


Figure S2 Transfection efficiency of miR-29b. (A) Transfection efficiency of miR-29b mimics detected by qRT-PCR, n=3. (B) Transfection efficiency of miR-29b inhibitor detected by qRT-PCR, n=8. ***, P<0.001; *, P<0.05. The data are expressed as the mean ± SEM. NC, negative control; AMO-miR-29b, miR-29b inhibitor; miR-29b, microRNA-29b; qRT-PCR, quantitative real-time reverse transcription polymerase chain reaction.