

# MiR-140-5p targets BCL2L1 to promote cardiomyocyte apoptosis

M.-Y. SUN<sup>1</sup>, L.-P. LI<sup>2</sup>

<sup>1</sup>Department of Clinical Laboratory, Jining No. 1 People's Hospital, Jining, China

<sup>2</sup>Department of Cardiology, Jining No. 1 People's Hospital, Jining, China

**Abstract. – OBJECTIVE:** Recovery of blood flow after ischemic cardiomyopathy can lead to aggravation of myocardial injury. This is very detrimental to the patient's prognosis. The purpose of this study was to investigate the effects and mechanisms of microRNA-140-5p (miR-140-5p) on myocardial ischemia-reperfusion injury (IRI).

**MATERIALS AND METHODS:** We made the myocardial IRI model in rats and detected the expression of miR-140-5p. Anta-miR-140-5p was administered intravenously in the tail of rats. Then, we used 2, 3, 5-triphenyl tetrazolium chloride staining, cardiac function test, and histological experiment to observe the changes of myocardial infarct size, cardiac function, and cardiomyocyte apoptosis in rats. In *in vitro* experiments, we induced the damage of H9c2 cells by hypoxia/reoxygenation (H/R) model and detected the effects of miR-140-5p on the proliferation ability and apoptosis level of H9c2 cells. TargetScan database was used to predict the binding target of miR-140-5p and we verified the effect of miR-140-5p on the target through Dual-Luciferase reporter assay.

**RESULTS:** MiR-140-5p expression in myocardial tissue of rat increased after IRI. Anta-miR-140-5p can reduce myocardial infarction area, improve cardiac function, and reduce the number of myocardial cells apoptosis in rats. The expression of miR-140-5p in H9c2 cells was higher than that in the control group. MiR-140-5p inhibition was found to promote the proliferation and increase the apoptosis level of H9c2 cells, while miR-140-5p mimic was the opposite. The TargetScan system predicts the presence of binding sites for miR-140-5p and B-cell lymphoma 2-like 1 (BCL2L1). The Dual-Luciferase reporter assay found that miR-140-5p can bind to BCL2L1 and promote its degradation. In addition, the inhibition of BCL2L1 was found to promote apoptosis of H9c2 cells.

**CONCLUSIONS:** In myocardial IRI, miR-140-5p targets BCL2L1 and promotes its degradation, thereby promoting myocardial apoptosis.

**Key Words:**

miR-140-5p, Myocardial ischemia-reperfusion injury, Apoptosis.

## Introduction

Acute myocardial infarction (AMI) is one of the diseases with high morbidity and mortality worldwide. It is mainly characterized by myocardial ischemic necrosis caused by coronary artery obstruction<sup>1</sup>. Reperfusion therapy is the basis for improving the prognosis of AMI patients. However, reperfusion itself increases irreversible damage to the myocardium and coronary circulation, leading to an increase in the infarction area. This pathological change is called myocardial ischemia-reperfusion injury (IRI)<sup>2</sup>. Wang<sup>3</sup> found that when patients with AMI did not accept reperfusion therapy, the area of myocardial infarction accounted for 70% of the ischemic danger area, and only accepting reperfusion therapy could reduce the area of myocardial infarction to 40%. When patients get both reperfusion therapy and myocardial protection therapy that can prevent myocardial IRI, the area of myocardial infarction will be significantly reduced to 5%. Therefore, to restore effective reperfusion of myocardium while adopting myocardial protection measures that can effectively prevent myocardial IRI will be the key to reduce the mortality and complication rate of AMI patients.

The mechanisms of myocardial IRI are diverse, such as free radical damage, calcium overload, oxidative stress, and myocardial energy metabolism disorders<sup>4</sup>. A variety of factors can jointly lead to the apoptosis of myocardial cells and irreversible damage to the myocardium. Cardiovascular disease and microRNA (miRNA) are closely related. MiRNAs have 20-24 nt in length and do not encode proteins. The biological functional diversity of miRNA is its distinctive feature<sup>5</sup>. MiRNAs can promote mRNA degradation through complementary pairing. In addition, there are many miRNA species, and 1048 species have been found. Among total human genes, miRNAs account for

only 2%, of which 20-30% are related to gene regulation<sup>6</sup>. Although the proportion of miRNAs is low, they play a strong regulatory role, especially in cell growth, cell proliferation, and apoptosis. MiRNAs play an increasingly important role in the occurrence and development of cardiovascular disease<sup>7</sup>. The knockout of some miRNAs often triggers myocardial remodeling and even heart failure<sup>6</sup>. There are many types of miRNA specific to cardiomyocytes. According to statistics, there are more than 800 types, including miR-1, miR-133, miR-24, miR-126, miR-145, miR-30, miR-132, and miR-122. When myocardial ischemia occurs, miRNAs will change significantly, especially in symptoms, such as hypoxia<sup>8</sup>.

MiR-140-5p belongs to the members of the miRNA family and is a common mature miRNA produced by miR-140 precursors. MiR-140-5p can regulate cell growth, tissue differentiation, organ development, and the occurrence of various cancers<sup>9</sup>. MiR-140-5p can act as a tumor suppressor on multiple different target genes and inhibit the invasion of tongue squamous carcinoma cells<sup>10</sup>. In addition, miR-140-5p can inhibit the proliferation and metastasis of hepatoma carcinoma cells, and can also participate in the formation of myelin sheath and the occurrence of obesity<sup>11</sup>. However, the effect of miR-140 on myocardial IRI was rarely studied. Therefore, we made the rat model and observed the role of miRNA-140-5p in myocardial IRI, laying a foundation for the subsequent research on the mechanism of myocardial IRI.

## Material and Methods

### Animals

Sprague Dawley (SD) rats (240±20g, male) was purchased and raised at the Jining No. 1 People's Hospital Animal Center. The rats were placed in an SPF barrier facility (12 hours of alternating light, 20-26 °C, 40-70% relative humidity) and fed with clean water and standard food. This study was approved by the Animal Ethics Committee of Jining No. 1 People's Hospital Animal Center.

### Myocardial IRI Model

The rats were anesthetized with 1.55% sodium pentobarbital (50 mg/kg) based on body weight. The rats were then fixed on the operating table in the supine position. The rat trachea was cut open and connected to a small animal ventilator (CWE SA-800, Orange, CA, USA). The small animal

ventilator is set to a ventilation frequency of 60-80 beats per minute, a tidal volume of 2 mL, and a breathing ratio of 1.5:1. After the skin in the anterior cardiac region of the rat was disinfected by iodophor, we used a sterile scalpel to cut from the rat's left chest and bluntly separated the tissue until the heart was exposed. After the pericardium was gently cut with a scalpel, the left anterior descending branch of the coronary artery of the rat was ligated with a sterile suture. The elevation of the ST segment of the electrocardiogram at this time indicates that the coronary blood flow was successfully blocked. After ligation for 30 minutes, the ligature was loosened for 20 minutes. We, then, took blood from the rat femoral artery and used cardiac gas analyser to measure cardiac function. Finally, we collected the heart part of the rats. Rats in the miR-140-5p antagomir (anta-Con) control (Con) group and miR-140-5p anta-miR-140-5p group were injected with 200 µL of anta-Con and anta-miR-140-5p into the tail vein 7 days before modeling, respectively. Anta-Con and anta-miR-140-5p were constructed in Shanghai GenePharma (Shanghai, China).

### Cardiac Function Test

7 days after the rat's coronary arteries were ligated, we used echocardiography to measure the ejection fraction (EF), short-axis contraction fraction (FS), and stroke output (SV) of the rat. EF, FS and SV can reflect the function of rat hearts. A decrease in their value indicates a decrease in cardiac function.

### 2, 3, 5-Triphenyl Tetrazolium Chloride (TTC) Staining

Rat hearts were washed twice with physiological saline and frozen in a -20 °C refrigerator. The hearts were then transected into 2 mm thick slices perpendicular to the long axis. Heart slices were then stained in 2% TTC staining solution (Sigma-Aldrich, St. Louis, MO, USA) at 37°C. 4% paraformaldehyde was used to fix the heart slices. Image J 1.51k (National Institutes of Health, Bethesda, MA, USA) was used to calculate the infarct area.

### Histology and Hematoxylin-Eosin (HE) Staining

Rat hearts were washed twice with normal saline and fixed in 4% paraformaldehyde for 24 hours. Myocardial tissue was washed with phosphate buffered saline (PBS). Myocardial tissue was dehydrated in ethanol solutions of different

concentrations. The dehydrated myocardial tissue was put into xylene and paraffin solution in order, and finally made into paraffin blocks. We then cut the paraffin block into 5 μm thick paraffin sections using a microtome (LEICA RM2235, Koln, Germany). Paraffin sections were placed in a 37°C incubator for 24 hours, and then, stored at room temperature. Before HE staining, paraffin sections were baked in a 55°C incubator for 1 hour. Paraffin sections were then dewaxed in xylene and hydrated in ethanol. The sections were then soaked in hematoxylin staining solution (Sigma-Aldrich, St. Louis, MO, USA) for 3 minutes. After differentiation with hydrochloric acid alcohol, the sections were stained with cytoplasm in eosin staining solution (Sigma-Aldrich, St. Louis, MO, USA). Finally, the sections were dehydrated and sealed with neutral gum. A high-power optical microscope (LEICA, Koln, Germany) is used to observe the morphology of myocardial tissue.

#### Immunohistochemical (IHC) Staining

After dewaxing and hydration, the sections were placed in citrate buffer and heated to 95°C for 20 minutes. After the temperature of citrate buffer was naturally cooled to room temperature, we took the sections and incubated the myocardial tissue with 3% H<sub>2</sub>O<sub>2</sub> for 30 minutes. We, then, washed the sections with PBS and incubated the myocardial tissue with 10% goat serum for 1 hour. After washing the remaining water on the sections, we incubated the myocardial tissue with primary antibody dilution (caspase-3, Abcam, Cambridge, MA, USA; caspase-8, Abcam, Cambridge, MA, USA) at 4°C overnight. The next day, after washing the sections with PBS, we incubated the sections with the secondary antibody dilution of the IHC staining kit (GeneTeck, Shanghai, China) for 1 hour and washed the sections with PBS. Finally, we use the coloring solution of the IHC staining kit for color development. A high-power optical microscope (LEICA, Koln, Germany) was used to observe the results of IHC staining.

#### RNA Isolation and Quantitative Reverse Transcription-Polymerase Chain Reaction (qRT-PCR)

Before extracting RNA from myocardial tissue and H9c2 cells, we immersed consumables, such as Eppendorf (EP; Hamburg, Germany) tubes and pipette tips in diethyl pyrocarbonate (DEPC) water (Beyotime, Shanghai, China) for 12 hours to remove RNase from the consumables. Myocardial tissue was cut and placed in Eppendorf (EP) tubes containing 1 mL of RPI reagent (Invitrogen, Carlsbad, CA, USA). Homogenization was used to grind myocardial tissue. 1 mL of chloroform was then added to the EP tube. After mixing, the EP tube was placed in a centrifuge (12000 rpm, 10 minutes, 4°C) and the supernatant was collected. 500 μL of absolute ethanol was added to the EP tubes and mixed. Spin column tubes were then used to collect the RNA from the solution. 700 μL of Buffer RWT and 500 μL of RPE were used to wash the RNA, respectively. Finally, 1 mL of DEPC water was used to dissolve the RNA in 1.5 mL spin tubes and collected the RNA solution in 1.5 mL eppendorf tubes. The RNA extraction method in H9c2 cells is similar to the above. RNA was first reversed into complementary deoxyribonucleic acid (cDNA). The RT system was 2 μL total RNA + 1 μL 2.5U/μU PolyA Polymerase + 1 μL RTase Mix + 5 μL 5 × Reaction buffer + 16 μL RNase-free water (Invitrogen, Carlsbad, CA, USA). SYBR Green Master Mix (Invitrogen, Carlsbad, CA, USA) was used to amplify cDNA. The primer sequences were shown in Table I. U6 expression was used as a control for miR-140-5p, and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) expression was used as a control for mRNA. The expression amount of RNA was represented by 2<sup>-ΔΔCT</sup>.

#### Detection of Lactate Dehydrogenase (LDH) in Serum

After collecting the blood of rats, we obtained the serum by centrifugation (3000 rpm, 5 minutes, 4°C) and stored it in a -80°C refrigerator. The

Table I. RT-PCR primer sequences.

Name	Sense sequences (5'-3')	Anti-sense sequences (5'-3')
miR-140-5p	ACATGAGCTATCGACGCATGTC	TCGGCTACTGAGGCGCAAA
Caspase-3	GGAACGCGAAGAAAAGTG	ATTTTGAATCCACGGAGGT
Caspase-8	CACATCCCGCAGAAGAAG	GATCCCGCCGACTGATA
BCL2L1	ACCCCTCTCCCTCCAG	CCTCAGCCAACTCTACGC
U6	GCTTCGGCAGCACATATACTAAAAT	CGCTTCACGAATTTGCGTGTGTCAT
GAPDH	ATGGCTACAGCAACAGGGT	TTATGGGGTCTGGGATGG

standards were diluted to different concentrations (1.25, 2.5, 5, 10, 20, 40 IU/mL). Then, 50  $\mu$ L of the test sample and standard were added to a 96-well plate. After the 96-well plate was incubated in a 37°C incubator for 30 minutes, we added 50  $\mu$ L of enzyme-labeled reagent to each well and made it incubated for another 30 minutes. Finally, 50  $\mu$ L of developer was added to each well and we used a microplate reader (Molecular Devices, Santa Clara Valley, MD, USA) to detect the absorbance of each well at 450 nm.

### Cell Culture

Rat myocardial cell line, H9c2 cells, were used in this study. We used 445 mL of Dulbecco's Modified Eagle's Medium (DMEM; Gibco, Rockville, MD, USA), 50 ml of fetal bovine serum (FBS; Gibco, Rockville, MD, USA), and 5 mL of double-antibody (Gibco, Rockville, MD, USA) configured as a complete medium to culture H9c2 cells. Cells were cultured in a 37°C incubator and all cell experiments were performed in a sterile cell clean bench.

### Procedure of Hypoxia/Reoxygenation (H/R) Model

H9c2 cells were seeded in Petri dishes. After the cell density reached 80%, we replaced the culture medium with hypoxia solution for 12 hours. Then, 99.9% nitrogen was passed into the incubator. The volume fraction of oxygen was controlled to less than 1% for 4 hours. Then, we returned the normal culture conditions and the cells continued to culture for 3 hours.

### Transfection

NC mimic, miR-140-5p mimic, miR-140-5p inhibitor, siRNA-BCL2L1 and lymphoma-2 like 1 (BCL2L1) were used to transfect H9c2 cells. The above mimics, inhibitors and siRNAs were all constructed by Shanghai GenePharm (Shanghai, China). After the cell growth density reached 80%, we cultured the cells for 24 hours in serum-free DMEM medium. Then, we dissolved mimics, inhibitors and siRNA in 50  $\mu$ L DMEM medium and mixed with Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA). The mixture was then added to the cell culture medium for 20 minutes. Finally, we re-cultured the cells with complete medium and detected the transfection efficiency by RT-PCR. NC mimic: 5'-AGUCUACGAGCGUAUAA-3'; NC mimic: 5'-GACGUAUUCGAGCCAUA; NC inhibitor: 5'-GUAUUCGAGCGUAUAA-3'; miR-

140-5p inhibitor: 5'-UGUACACACGAUUGACGUG-3'; siRNA-BCL2L1-sense: AUGGUAUUCGAGCCAUA and siRNA-BCL2L1-antisense: UGCUAUACGAGCAGUU.

### Cell Counting Kit-8 (CCK8)

H9c2 cells were seeded in 96-well plates. After the cell growth density reached 50%, we treated the cells differently and added 10  $\mu$ L of CCK8 reagent (Yifeixue, Nanjing, China) to each well. The cells were then cultured in an incubator for 2 hours. Finally, we used the microplate reader to detect the absorbance of each well at 450 nm.

### Immunofluorescence (IF) Staining

H9c2 cells were seeded in 24-well plates. After the cell growth density reached 50%, we treated the cells differently. We then washed the cells with PBS. Cells were immersed in 4% paraformaldehyde for 20 minutes and then immersed them in 0.5% Triton-100 for 15 minutes. After washing the cells with PBS, 10% goat serum was used to block non-specific antigens for 2 hours. We then incubated the cells at 4°C overnight using primary antibody dilution (caspase-3, Abcam, Cambridge, MA, USA; caspase-9, Abcam, Cambridge, MA, USA). The next day, we washed away the excess primary antibody with PBS and incubated the cells with the secondary antibody dilution (Abcam, Cambridge, MA, USA) for 1 hour. Then, we stained the nucleus with DAPI (Sigma-Aldrich, St. Louis, MO, USA). Finally, we used a fluorescence microscope (LEICA, Koln, Germany) to observe the staining results.

### MiRNA Target Prediction and Dual-Luciferase Reporter Assay

We searched the TargetScan database ([http://www.targetscan.org/vert\\_72/](http://www.targetscan.org/vert_72/)) for target genes that could bind to miR-140-5p. After obtaining the possible target gene of miR-140-5p, we verified it by Dual-Luciferase reporter assay. H9c2 cells were seeded in large petri dishes. After the cell growth density reached 85%, we collected H9c2 cells and washed them with PBS. The cells were divided into WT-BCL2L1 + NC mimic group, WT-BCL2L1 + miR-140-5p mimic group, MUT-BCL2L1 + NC mimic group and MUT-BCL2L1 + miR-140-5p mimic group. After the transfected cells were cultured in an incubator for 48 hours, the cells were added to a 96-well plate and 75  $\mu$ L of Dual-Glo Luciferase Reagent (Millipore, Billerica, MA, USA) was added to each



well. We, then, used a fluorophotometer (Thermo Fisher Scientific, Waltham, MA, USA) to detect the Luciferase content.

### Flow Cytometry

We collected H9c2 cells and washed them with PBS. After centrifugation (1000 rpm, 5 minutes), 5  $\mu$ L of AnnexinV-FICT (Sigma-Aldrich, St. Louis, MO, USA) was added to the centrifuge tubes and mixed. After 10 minutes, 1  $\mu$ L of 100  $\mu$ g/ $\mu$ L PI staining solution was added to a centrifuge tube and incubated for 5 minutes. Finally, 400  $\mu$ L of each tube was detected by flow cytometry.

### Statistical Analysis

All data in this study were entered into the EXCEL form for storage and analysis using Statistical Product and Service Solutions (SPSS) 20.0 statistical software (IBM Corp., Armonk, NY, USA). The results of the analysis are expressed as mean  $\pm$  standard deviation (SD). The differences between the two groups were analyzed by using the Student's *t*-test. Comparison between multiple groups was done using One-way ANOVA test followed by post-hoc test (Least Significant Difference). All experiments were repeated more than 3 times. *p*<0.05 indicates that the difference is statistically significant.

## Results

### MiR-140-5p Was Highly Expressed in Myocardial Tissue of IRI Rats and Anta-MiR-140-5p Can Inhibit Myocardial IRI

The rats were divided into Sham group, IRI group, IRI + anta-Con group and IRI + anta-miR-140-5p group. The expression of miR-140-5p in the myocardial tissue of the four groups of rats was detected and the expression of miR-140-5p in the IRI and IRI + anta-Con groups was higher than that in the Sham group and the IRI + anta-miR-140-5p group (Figure 1A). We measured the area of myocardial infarction in rats by the TTC staining. The area of myocardial infarction in IRI and IRI + anta-Con groups was larger than that in Sham group and IRI + anta-miR-140-5p group (Figure 1B). In addition, we measured the concentration of LDH in the serum compared with the Sham group, the LDH concentration of the rats in the IRI group and the IRI + anta-Con group was significantly increased. Anta-miR-140-5p can reduce the concentration of LDH (Figure 1C). Myocardial IRI

also caused a decrease in cardiac function in rats, manifested as a decrease in EF (Figure 1D), SV (Figure 1E), and LV (Figure 1F), while anta-miR-140-5p could partially restore cardiac function in rats. The results of RT-PCR showed that anta-miR-140-5p can reduce the caspase3/8 mRNA (Figure 1G). We examined the pathological changes of myocardial tissue by HE staining (Figure 1H). The cardiomyocytes of the rats in the IRI group and IRI + anta-Con group were significantly disturbed and there was inflammatory cell infiltration among the cardiomyocytes, while the cardiomyocytes of the rats in the IRI + anta-miR-140-5p group were significantly improved. We detected the expression of apoptosis-related molecule caspase3/8 in rat myocardial tissue by IF staining and the results showed that the apoptosis level of myocardial cells in IRI rats was higher than that in the Sham group and anta-miR-140-5p could reduce the level of apoptosis (Figure 1H).

### MiR-140-5p Was Highly Expressed in H9c2 Cells and miR-140-5p Mimic Induced H9c2 Cell Apoptosis, while Anta-miR-140-5p Can Reduce H/R-Induced Cell Damage

The proliferation level of H9c2 cells treated with H/R was detected by CCK-8 assay. The proliferation level of cells treated with H/R decreased compared to the Control group, indicating that H/R can successfully induce H9c2 cell damage (Figure 2A). Then, miR-140-5p was found to be highly expressed in H9c2 cells of the H/R group (Figure 2B). NC mimic, miR-140-5p mimic, NC inhibitor and miR-140-5p inhibitor were constructed to detect the effects of miR-140-5p on H9c2 cells. RT-PCR was used to verify the transfection efficiency of miR-140-5p mimic and miR-140-5p inhibitor (Figure 2C). Flow cytometry was used to detect the cell apoptosis rate and it was found that miR-140-5p inhibitor can inhibit the apoptosis of H9c2 cells, while miR-140-5p mimic is the opposite (Figure 2D). In addition, we detected the expressions of caspase3/8 mRNA and protein in H9c2 cells using RT-PCR (Figure 2E) and IF staining (Figure 2F, G), respectively. The results also demonstrated the anti-apoptotic effect of miR-140-5p on H9c2 cells.

### MiR-140-5p Binds BCL2L1 and Induces its Degradation

TargetScan database was used to predict the targets of miR-140-5p. In rats, miR-140-5p was predicted to have a conserved binding site to BCL2L1 (Figure 3A). Therefore, we used the

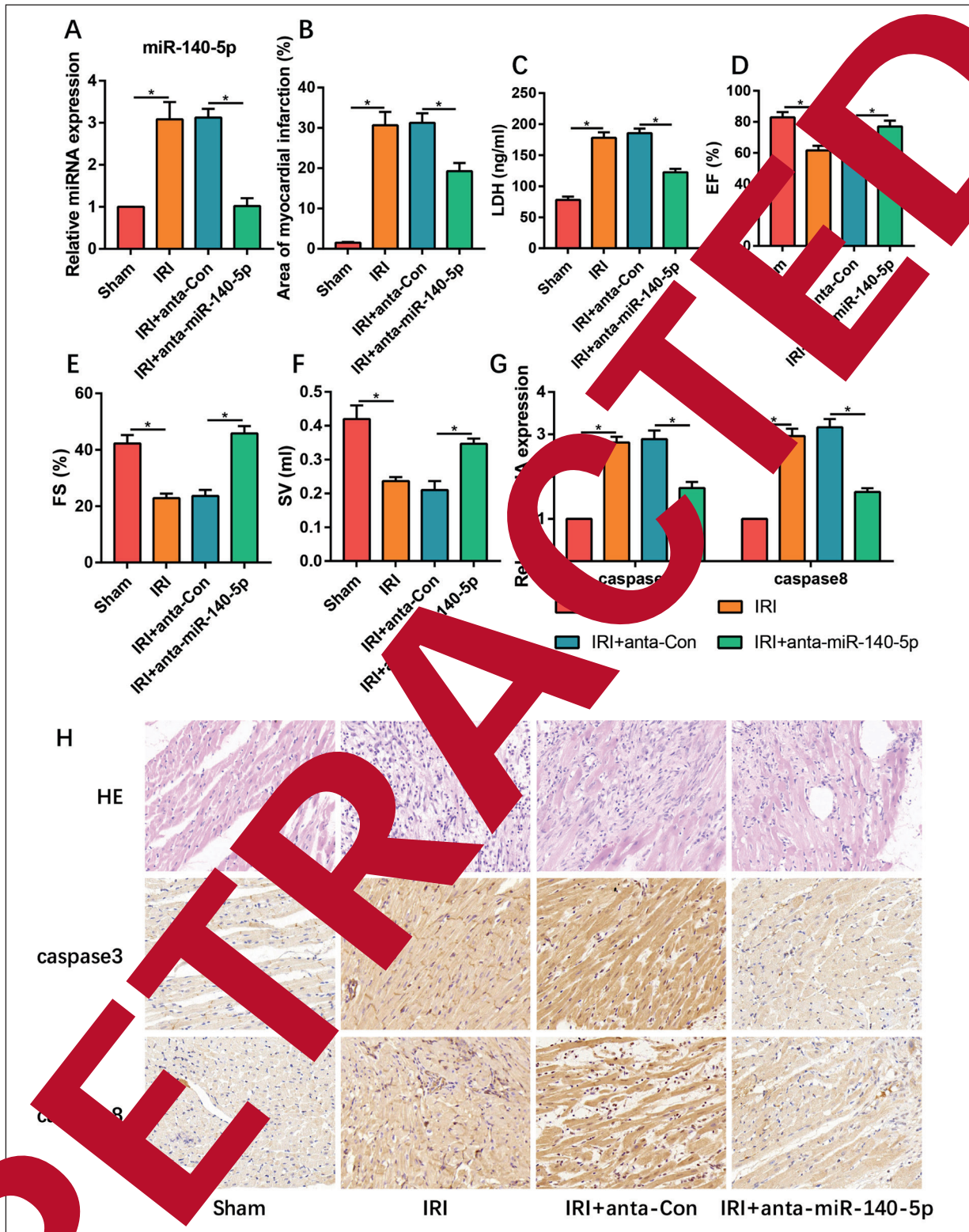
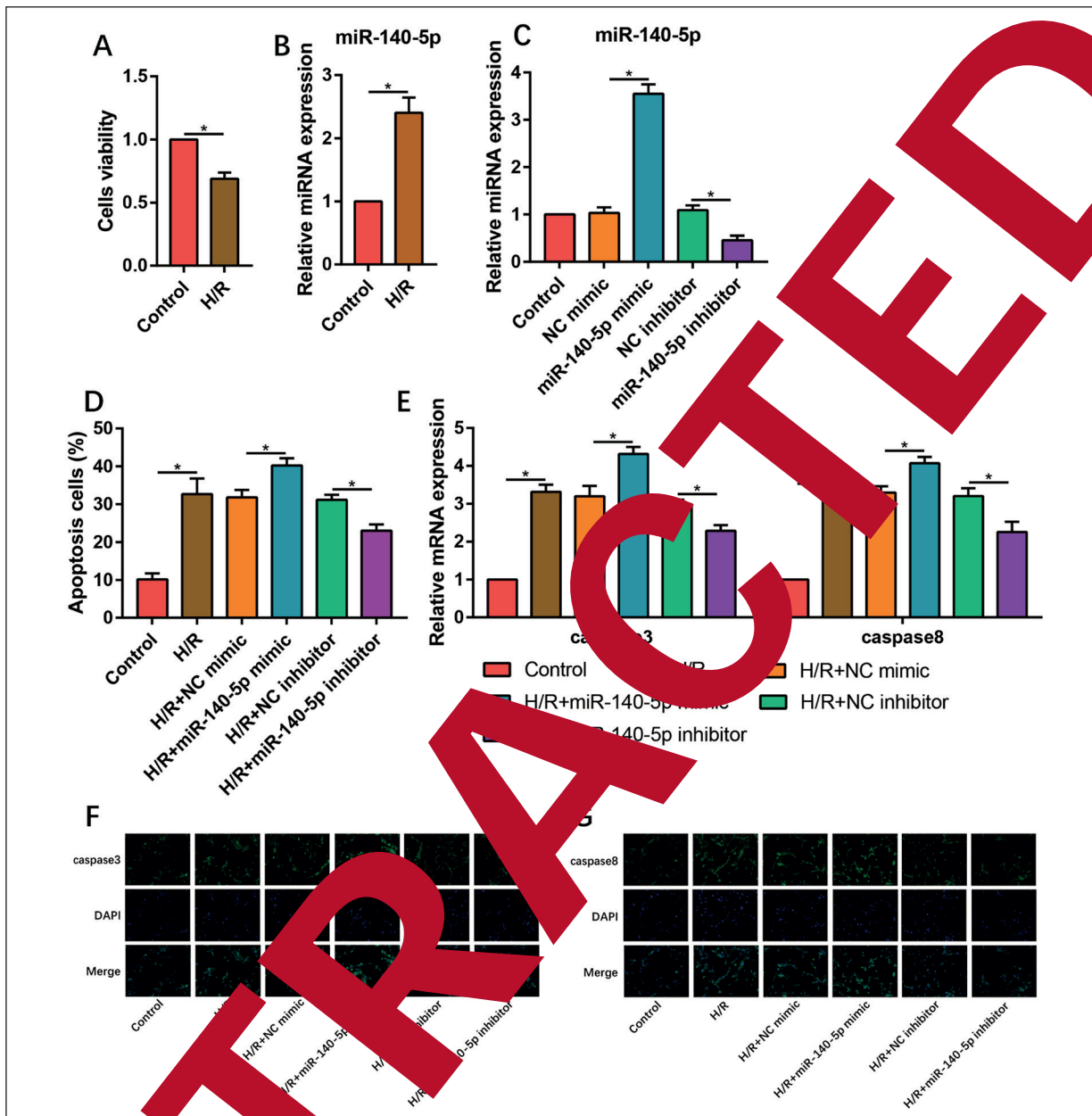


Figure 1. miR-140-5p was highly expressed in myocardial tissue of IRI rats and anta-miR-140-5p can reduce myocardial IRI. **A**, The changes of miR140-5p expression in IRI rats. **B**, The area of myocardial infarction was detected by TTC staining. **C**, The concentration of LDH in rat serum. **D-F**, EF, FS and SV of rats was detected by echocardiography. **G**, mRNA expression of caspase-3/8 was detected by RT-PCR. **H**, HE staining and IHC staining of caspase3/8 (magnification: 200×). (\*\*\*) means the difference is statistically significant,  $p < 0.05$ .



**Figure 2.** MiR-140-5p was highly expressed in H/R-induced H9c2 cells and miR-140-5p inhibitor can reduce H/R-induced cell damage. **A**, The cells viability of H9c2 cells was detected by CCK8 assay. **B-C**, The changes of miR140-5p expression in H9c2 cells. **D**, The apoptosis rate of H9c2 cells was detected by flow cytometry. **E**, mRNA expression of caspase3/8 was detected by qPCR. **F-G**, The results of caspase3/8 in H9c2 cells (magnification: 200×). (“\*”) means the difference is statistically significant ( $p < 0.05$ ).

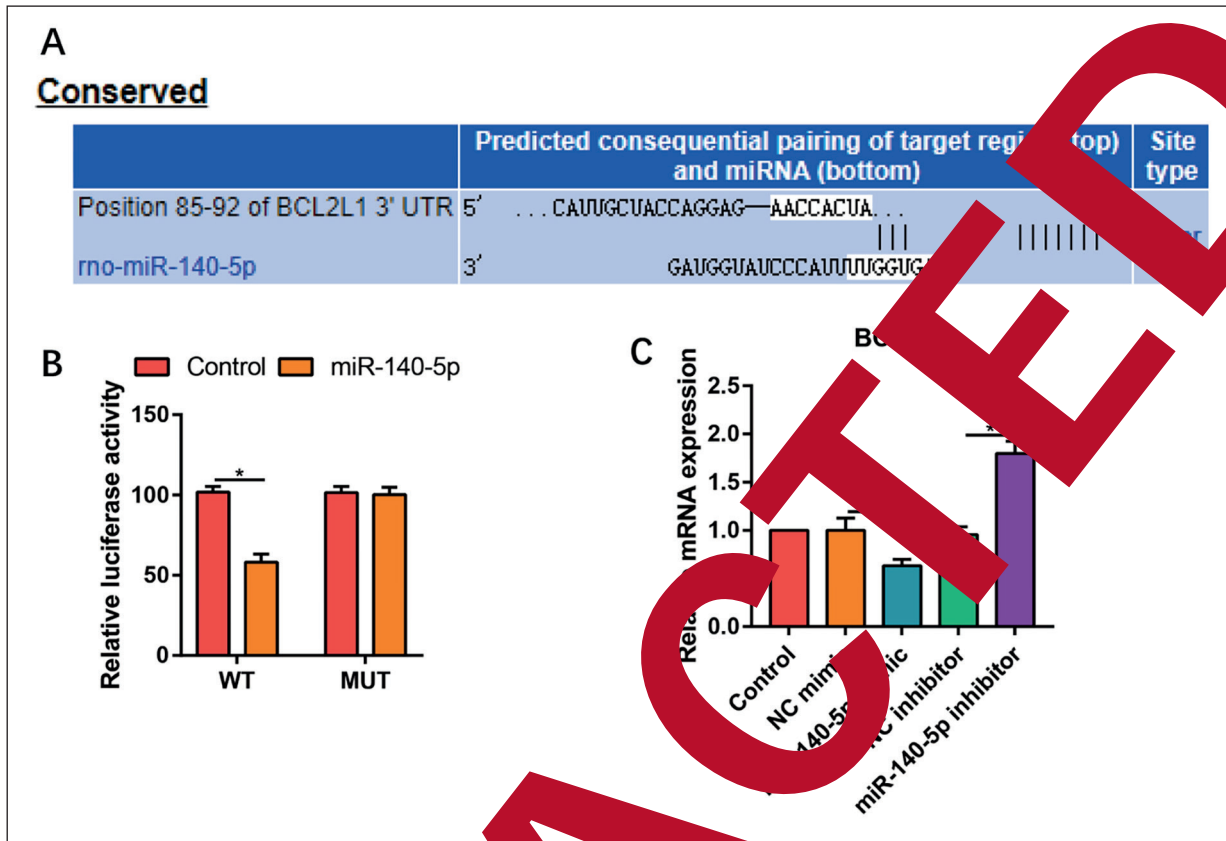
Luciferase reporter assay to detect the interaction between miR-140-5p and BCL2L1. MiR-140-5p mimic and WT-BCL2L1-3'UTR were used in H9c2 cells. The results showed that after the infection of WT-BCL2L1-3'UTR, the level of luciferin in H9c2 cells was significantly reduced, while MUT-BCL2L1-3'UTR had no such effect (Figure 3B). In addition, the results of

RT-PCR also showed that miR-140-5p can act on BCL2L1 and degrade it (Figure 3C).

#### ***Inhibition of BCL2L1 Attenuated the Protective Effect of MiR-140-5p Inhibitor on H9c2 Cells***

To verify the effect of BCL2L1 in H9c2 cells, we used siRNA-BCL2L1 to reduce the expression





**Figure 3.** MiR-140-5p binds BCL2L1 and induces mRNA degradation. **A**, MiR-140-5p was predicted to have a conserved binding site to BCL2L1. **B**, Dual-Luciferase reporter assay showed that miR-140-5p promoted the degradation of BCL2L1 mRNA. **C**, mRNA expression of BCL2L1 was detected by RT-PCR. \* means the difference is statistically significant,  $p < 0.05$ .

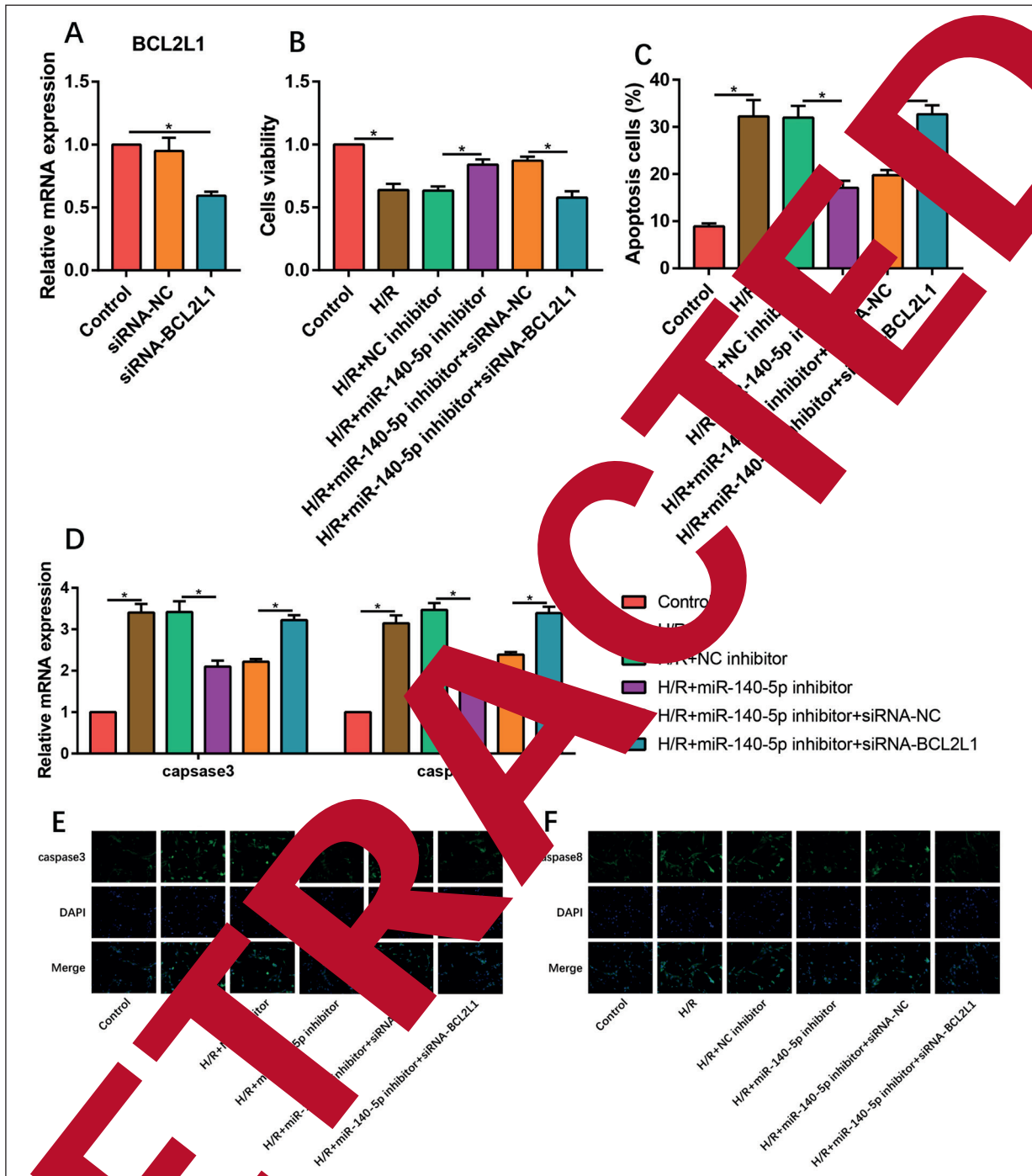
of BCL2L1 in the cells. RT-PCR was used to verify its transfection efficiency. CCK-8 assay detected the viability of cells. MiR-140-5p inhibitor can partially restore IRI-induced decrease in cell viability, while siRNA-BCL2L1 inhibits the effect of miR-140-5p inhibitor (Figure 4B). Flow cytometry also showed that siRNA-BCL2L1 reduced the inhibitory effect of miR-140-5p inhibitor on apoptosis (Figure 4C). RT-PCR (Figure 4D) and IF staining (Figure 4E, 4F) detected the expression of caspase3/8 mRNA and protein in H9c2 cells, respectively. We found that inhibition of BCL2L1 increased the expression of caspase3/8 mRNA and protein.

### Discussion

Myocardial ischemia is a common clinical disease that occurs in many tissues and organs. Diseases caused by tissue ischemia are collectively referred to as ischemic diseases<sup>12</sup>. The main cause of

myocardial ischemia is coronary artery stenosis, which generally exceeds 70%. Decreased myocardial blood flow perfusion at this time often induces myocardial hypoxia and causes metabolic disorders<sup>13</sup>. When the ischemic myocardium is reperfused, the myocardium is often damaged to different degrees, which induces arrhythmia and other symptoms. Severe myocardial IRI can even cause ventricular contraction and diastolic dysfunction, and cause myocardial cell necrosis, cardiomyocyte autophagy, and cardiomyocyte apoptosis<sup>14</sup>. MiRNA, as a medical research hotspot in recent years, has been found to be a therapeutic target for many diseases. In order to study the effect of miR-140-5p on myocardial IRI, we made a model of myocardial IRI using rats and detected the high expression of miR-140-5p in injured myocardial tissue. We determined the effect of IRI model by TTC staining and myocardial injury marker detection. The results showed that the myocardial infarction of the rats in the Sham group did not appear, while





**Figure 4.** Effect of BCL2L1 attenuated the protective effect of miR-140-5p inhibitor on H9c2 cells. **A**, qRT-PCR determined the relative expression of BCL2L1. **B**, Efficiency of siRNA-BCL2L1 was determined by qRT-PCR. **C**, Cells viability of H9c2 cells was detected by CCK8 assay. **C**, the apoptosis rate of H9c2 cells was detected by flow cytometry. **D-F**, Expression of caspase3/8 was determined by qRT-PCR and immunofluorescence (magnification: 200 $\times$ ). (\*\*\*) means the difference is statistically significant,  $p < 0.05$ .

area of myocardial infarction in the rats with blocked coronary artery increased significantly, indicating that the rat IRI model was successfully established. Compared with IRI group and

IRI + anti-Con group, the area of myocardial infarction in anti-miR-140-5p treated rats was improved. In addition, the cardiac function test in rats also found that EF, FS, and SV in IRI rats

were significantly reduced, indicating that the contractile and diastolic functions of myocardium in IRI rats were decreased. After treating rats with anta-miR-140-5p, the EF, FS and SV of rats all increased, which also confirmed the myocardial protective effect of anta-miR-140-5p. In the experiments on pathological staining and IHC staining of rat myocardial tissue, we also found that myocardial injury and apoptosis in rats treated with anta-miR-140-5p were significantly improved. *In vitro* experiments, we make cell models of IRI by H/R. After H/R treatment, the proliferation level of H9c2 cells decreased significantly, indicating that H/R successfully induced damage to H9c2 cells. The expression of miR-140-5p in H/R-induced cells was significantly higher than that in the Control group. This is similar to the results of animal experiments. In addition, miR-140-5p inhibitor was found to reduce the apoptosis rate and apoptosis molecule (caspase3/8) in H9c2 cells. Caspase3 is the main terminal cleaving enzyme in the process of apoptosis. Caspase3 is activated in the early stages of apoptosis, and is the last known execution factor of apoptosis<sup>15</sup>. MiR-140-5p was found to have possible binding sites with BCL2L1. The results of the Dual-Luciferase reporter experiment also confirmed that miR-140-5p can bind and degrade BCL2L1 mRNA. The BCL2L1 family also plays an important role in the process of apoptosis<sup>16</sup>. The BCL2 family is divided into three different subfamilies according to the differences in function and structure, among which the BCL2 subfamily plays a major role in the process of apoptosis, mainly composed of BCL2, BCL-XL, BCL-W, and BCL1L. As a member of the BCL2 family, BCL2L1 is involved in the development of many diseases. Wang et al<sup>18</sup> studied mesenchymal stem cells (MSCs) that showed a decrease in BCL2L1 under heat stress, leading to the apoptosis of MSCs. They found that epigallocatechin gallate increased BCL2L1 and thus protects cells from heat stress. Ghaemi et al<sup>19</sup> also found that miR-142 can induce glioblastoma cell apoptosis by inhibiting BCL2L1 and MCL1. To verify the effect of BCL2L1 in cardiomyocytes, we used siRNA to reduce BCL2L1 expression in H9c2 cells. The results of CCK-8 and flow cytometry revealed that after BCL2L1 was downregulated in H9c2 cells, the cell proliferation level decreased and the apoptosis rate increased. This indicates that BCL2L1 is a protective gene of H9c2 cells. RT-PCR and IF staining also confirmed this. In the study of myocardial fibrosis,

Schaefer<sup>20</sup> found that BCL2L1 plays an important role in stabilizing the mitochondrial membrane potential of cardiomyocytes. The protein kinase CK2 can regulate the redox balance in the cell by affecting the expression of BCL2L1, thereby protecting the myocardial cells from oxidative stress.

MiR-140-5p was originally discovered as a cartilage-specific miRNA in zebrafish. It is a miRNA that regulates cartilage development and proliferation, and maintains the homeostasis of the cartilage. Its low expression is related to the occurrence of osteoarthritis<sup>21</sup>. MiR-140-5p expression is induced in human vascular smooth muscle cells treated with serum-free medium. This suggests that miR-140-5p may play a key role in vascular smooth muscle phenotypic transformation<sup>22</sup>. MiR-140-5p can also regulate cell cycle and cell proliferation. By targeting the transcription factor Sn1, miR-140-5p can promote the proliferation of chondrocytes and suppress the cell cycle<sup>23</sup>. In breast cancer, miR-140-5p inhibits the occurrence and development of breast cancer by inhibiting the ERK1/2, SOX2 and SOX9 signaling pathways<sup>24</sup>. Xing et al<sup>25</sup> found that miR-140-5p can aggravate the damage of H9c2 cells by regulating the expression of BCL2L1. In addition, miR-140-5p is also involved in the occurrence and development of various inflammatory diseases. After myocardial injury induced by doxorubicin, miR-140-5p increased and then induced apoptosis of cardiomyocytes *in vivo* via regulating Nrf2 and VEGFA/14-3-3 $\gamma$  signaling pathways<sup>26,27</sup>. In a hypoxic-induced pulmonary hypertension model, upregulation of miR-140-5p promoted the apoptosis of pulmonary artery smooth muscle cells by regulating SOD2 and Dnmt1<sup>28</sup>. These results indicated that miR-140-5p plays an important role in the occurrence and development of multiple diseases. We also demonstrated the effect of miR-140-5p on cardiomyocytes in animal experiments and found another regulatory way for miR-140-5p. These results indicated that miR-140-5p can influence the process of myocardial IRI in various ways. Therefore, miR-140-5p may be a new target for the treatment of myocardial IRI.

To sum up, miR-140-5p, as an important regulatory factor regulating a variety of biological activities *in vivo*, has an important effect on myocardial IRI. To our knowledge, this is the first study to investigate the effect of miR-140-5p on myocardial IRI. We believed that this study can provide a new target and theoretical basis for the clinical treatment of myocardial IRI.

## Conclusions

Altogether, this study showed that, in myocardial IRI, miR-140-5p can target the anti-apoptotic molecule BCL2L1 in cardiomyocytes, leading to the degradation of BCL2L1, thereby promoting the apoptosis of cardiomyocytes. Therefore, the inhibition of miR-140-5p can be used as a treatment for myocardial IRI.

## Conflict of Interest

The Authors declare that they have no conflict of interests.

## References

- MEFFORD MT, LI BH, QIAN L, READING SR, HARRISON TN, SCOTT RD, CAVENDISH JJ, JACOBSEN SJ, KANTER MH, WOODWARD M, REYNOLDS K. Sex-specific trends in acute myocardial infarction within an integrated healthcare network, 2000 through 2014. *Circulation* 2020; 141: 509-519.
- CHI HJ, CHEN ML, YANG XC, LIN XM, SUN H, ZHAO WS, QI D, DONG JL, CAI J. Progress in therapy of myocardial ischemia reperfusion injury. *Cell Targets* 2017; 18: 1712-1721.
- YELLON DM, HAUSENLOY DJ. Realizing the clinical potential of ischemic preconditioning and postconditioning. *Nat Clin Pract Cardiovasc Med* 2005; 5: 568-575.
- KRZYWONOS-ZAWADZKA A, FRONCZYK M, WAWICKI G, WOZNAK M, BIL-LULA I. Multifaceted prevention or therapy of ischemia-reperfusion injury of the heart-Mini-review. *Environ Toxicol Pharmacol* 2017; 55: 55-59.
- ONG SB, KATWADI S, KAWEK XY, CHAN NI, CHINDA K, ONG SG, HAUSENLOY DJ. Non-coding RNAs as therapeutic targets for preventing myocardial ischemia-reperfusion injury. *Expert Opin Ther Targets* 2018; 22: 247-257.
- MAKHDOUMI P, ROOHBAKHSHI M, KARIMI G. MicroRNAs regulate mitochondrial apoptotic pathway in myocardial ischemia-reperfusion-injury. *Biomed Pharmacother* 2016; 84: 1635-1644.
- LIU A, RAI A, ORTICELLO M, LUCARELLI M. The complex interplay between DNA methylation and miRNAs in gene expression regulation. *Biochimie* 2020. Epub ahead of print. doi:10.1016/j.biochi.2020.02.006.
- FRANQUINOS J, GENSCHEL C, VIERECK J, RUMPF YONGYOSI M, MAXLER D, RIESENHUBER M, SPANNBAUER A, LUKOVIC D, WEBER N, ZLABINGER K, HASIMBEGOVIC M, FIEDLER J, DANGWAL S, FISCHER M, DE LA FUENTE J, JUCIECHOWSKI D, KRAFT T, GARAMVOLGYI R, NEITZEL S, CHATTERJEE S, YIN X, BAR C, MAYR M, XIAO Y, THUM T. Preclinical development of a miR-132 inhibitor for heart failure treatment. *Nat Commun* 2020; 11: 633.
- DUALE N, EIDE DM, AMBERGER ML, GRAUBNER A, BREDE DA, OLSEN AK. Using prediction algorithms to identify miRNA-based markers for low-dose acute rate chronic stress. *Sci Total Environ* 2020; 717: 137068.
- XUE J, ZHANG Z, LI X, REN Y, LIANG Q. Long non-coding RNA TTN-AS1 promotes breast cancer cell migration and invasion via sponging miR-140-5p. *Oncol Lett* 2020; 19: 1255-1260.
- OSÉS M, MARGARETO S, PORTILLO MP, AGUILERA I, LABAYEN I. Circulating miRNAs as biomarkers of obesity and obesity-associated comorbidities in children and adolescents: a systematic review. *Nutrients* 2020; 11: 2833.
- XIA H, LIU Y, WANG W, ZENG X, LIU Y, LIU P, ZHONG Z, YE Q. miR-140-5p overexpression attenuates hepatic ischemia/reperfusion injury by regulating glutathione production and transformation. *Oxid Med Cell Longev* 2020; 2020: 1079129.
- LIU B, HEUSCH G, VAN DE WERF F. Evolutionary therapies for myocardial ischemia/reperfusion injury. *J Am Coll Cardiol* 2015; 65: 1454-1471.
- ALDIWANI H, ZHANG M, SUPPOGU N, QUESADA O, JOHNSON BD, MEHTA PK, WU J, WELT C, PETERSEN J, AZARBAL B, SAMUELSON B, ANDERSON RD, SHAW LJ, KAR S, HANDBERG E, WILSON JF, BAIREY MC. Angina hospitalization in women with signs and symptoms of ischemia but no obstructive coronary artery disease: a report from the WISE (Women's Ischemia Syndrome Evaluation) Study. *J Am Heart Assoc* 2017; 6: e13168.
- GUPTA S, KNOWLTON AA. HSP60, Bax, apoptosis and the heart. *J Cell Mol Med* 2005; 9: 51-58.
- VOGLER M, WALTER HS, DYER M. Targeting anti-apoptotic BCL2 family proteins in haematological malignancies - from pathogenesis to treatment. *Br J Haematol* 2017; 178: 364-379.
- POLCIC P, MENDEL M. Reconstituting the mammalian apoptotic switch in yeast. *Genes (Basel)* 2020; 11: 145.
- BUTT H, MEHMOOD A, EJAZ A, HUMAYUN S, RIAZUDDIN S. Epigallocatechin-3-gallate protects Wharton's jelly derived mesenchymal stem cells against in vitro heat stress. *Eur J Pharmacol* 2020; 872: 172958.
- GHAEMI S, AREFIAN E, REZAZADEH VR, SOLEIMANI M, MORADIMOTLAGH A, JAMSHIDI AF. Inhibiting the expression of anti-apoptotic genes BCL2L1 and MCL1, and apoptosis induction in glioblastoma cells by microRNA-342. *Biomed Pharmacother* 2020; 121: 109641.
- SCHAEFER S, GUERRA B. Protein kinase CK2 regulates redox homeostasis through NF- $\kappa$ B and Bcl-xL in cardiomyoblasts. *Mol Cell Biochem* 2017; 436: 137-150.
- PAPATHANASIOU I, BALIS C, TRACHANA V, MOURMOURA E, TSEZOU A. The synergistic function of miR-140-5p and miR-146a on TLR4-mediated cytokine secretion in osteoarthritic chondrocytes. *Biochem Biophys Res Commun* 2020; 522: 783-791.

- 22) ZHU TT, ZHANG WF, YIN YL, LIU YH, SONG P, XU J, ZHANG MX, LI P. MicroRNA-140-5p targeting tumor necrosis factor-alpha prevents pulmonary arterial hypertension. *J Cell Physiol* 2019; 234: 9535-9550.
- 23) WANG Y, SHEN S, LI Z, LI W, WENG X. MIR-140-5p affects chondrocyte proliferation, apoptosis, and inflammation by targeting HMGB1 in osteoarthritis. *Inflamm Res* 2020; 69: 63-73.
- 24) BAI F, YU Z, GAO X, GONG J, FAN L, LIU F. Simvastatin induces breast cancer cell death through oxidative stress up-regulating miR-140-5p. *Aging (Albany NY)* 2019; 11: 3198-3219.
- 25) XING B, LI OJ, LI H, CHEN SS, CUI ZY, MA J, ZHANG ZW. MiR-140-5p aggravates hypoxia-induced cell injury via regulating MLK3 in H9c2 cells. *Biomed Pharmacother* 2018; 103: 1652-1657.
- 26) CHEN XY, HUANG WL, PENG XP, LV YN, LI JH, XIONG JP. MiR-140-5p mediates bevacizumab-induced cytotoxicity to cardiomyocytes by targeting VEGFA/14-3-3gamma signal pathway. *Toxicol Res (Camb)* 2019; 8: 875-884.
- 27) ZHAO L, QI Y, XU L, TAO X, HAN Y, PENG J. MicroRNA-140-5p aggravates doxorubicin-induced cardiotoxicity by promoting myocardial oxidative stress via targeting Nrf2 and Sirt2. *PLoS One* 2018; 15: 284-296.
- 28) ZHANG Y, XU J. MiR-140-5p regulates hypoxia-mediated human pulmonary artery smooth muscle cell proliferation, apoptosis and differentiation by targeting Dcaml and promoting SCML2 expression. *Biochim Biophys Res Commun* 2016; 473: 342-348.