RESEARCH ARTICLE

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SERPINB1 overexpression protects myocardial damage induced by acute myocardial infarction through AMPK/mTOR pathway

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Abstract

Background: SERPINB1 is involved in the development of a variety of diseases. The purpose of this study was to explore the effect of SERPINB1 on acute myocardial infarction (AMI).

Methods: Serum SERPINB1 level of AMI patients was measured for receiver operating characteristic curve analysis. The AMI rat model was constructed to observe myocardial damage, and the H9C2 cell oxygen glucose deprivation (OGD) model was constructed to detect cell viability. Transthoracic echocardiography was used to assess the cardiac function. TTC staining and HE staining were used to detect pathologic changes of myocardial tissues. The apoptosis of myocardial tissues and cells were measured by TUNLE staining and flow cytometry assay. CCK-8 assay to measure cell viability. SERPINB1 expression was measured by qRT-PCR. Protein expression was measured by western blot.

Results: The serum SERPINB1 level was down-regulated in AMI patients. AMI modeling reduced the SERPINB1 expression level, induced inflammatory cells infiltrated, and myocardial apoptosis. OGD treatment inhibited cell viability and promoted apoptosis. The AMPK/mTOR pathway was inhibited in AMI rats and OGD-treated H9C2 cells. Overexpression of SERPINB1 reduced infarct size and myocardial apoptosis of AMI rats, inhibited apoptosis of H9C2 cells, and activated AMPK/mTOR pathway. However, AMPK inhibitor Dorsomorphin reversed the protective effect of SERPINB1 on myocardial cells.

Conclusion: SERPINB1 overexpression relieved myocardial damage induced by AMI via AMPK/mTOR pathway. **Keywords:** SERPINB1, Acute myocardial infarction, AMPK/mTOR pathway, Apoptosis

Background

Acute myocardial infarction (AMI) is myocardial necrosis caused by acute and persistent ischemia and hypoxia in coronary arteries [1]. The main feature of AMI is apoptosis of myocardial cells [2, 3]. AMI is one of the most serious ischemic heart diseases. The morbidity and

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⁴ Department of Clinical Laboratory, Central Hospital of Shengli Oilfield, No. 31 Jinan Road, Dongying 257000, Shandong, People's Republic of China mortality of AMI remains high although the mortality rate of AMI has been declining over the past decades [4]. Therefore, the therapeutic measures involved in the mechanism of myocardial cell apoptosis will provide a new therapeutic method for the treatment of AMI. Serine protease inhibitors (serpins) are a protein super-

Serine protease inhibitors (serpins) are a protein superfamily with conserved tertiary structure [5, 6]. Due to the lack of secretory signal sequence, SERPINB1 belongs to the B serpins, which mainly resides in the cytoplasm of neutrophils and monocytes [7]. El Ouaamari et al. have shown that SERPINB1 promotes pancreatic β cell



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proliferation in multiple species as a circulating factor secreted by the liver [8]. Study of Yao et al. has found that SERPINB1 can alleviate acute lung injury in liver transplantation [9]. Inhibition of SERPINB1 attenuated the protective effects of FoxO1 on diabetic nephropathy in vitro [10]. Besides, serpins may prevent acute cardiovascular syndromes by inhibiting serine proteases [11]. Also, SerpinA1 was shown to directly affect apoptosis by inhibition of caspase-1 and caspase-3 [12, 13]. Alpha-1-antitrypsin (the protein encoded by SerpinA1) treatment reduced infarct sizes in AMI mice [13]. Furthermore, SerpinB9 was demonstrated to inhibit cell apoptosis and protect against atherosclerotic lesion progression [14, 15]. However, the molecular mechanism of SERPINB1 acting on AMI still needs further detailed study.

Myocardial apoptosis induced by AMI may be related to the activation or inhibition of signaling pathways. Na et al. has demonstrated that gambogic acid protects against myocardial damage in MI rats by activating NF-κB and p38 pathways [16]. Liu et al. have found that inhibition of AMPK aggravated cell injury in OGD-mediated cardiomyocytes [17]. Additionally, Hua et al. have found that AMPK/mTOR signaling pathway is involved in AMI process [18]. A study of Wang et al. reported that activation of AMPK attenuated isoprenaline-induced myocardial fibrosis in vivo and decreased collagen deposition [19]. Furthermore, activation of the AMPK/mTOR pathway attenuates myocardial injury and cardiac insufficiency following MI [20]. However, whether SERPINB1 could alleviate myocardial apoptosis induced by AMI by regulating the AMPK/mTOR pathway has not been fully explored.

In this study, we measured serum SERPINB1 levels in AMI patients and healthy volunteers. Then we established the AMI rat model and cell model by LAD and OGD treatment to detect the effect of SERPINB1 on AMI rats and OGD-treated myocardial cells. The results showed that SERPINB1 in AMI patients was down-regulated. Overexpression of SERPINB1 could significantly alleviate myocardial damage and apoptosis induced by AMI through activating the AMPK/mTOR pathway. The results of our study might provide new ideas for the therapy of AMI.

Methods

Patients and samples

In this prospective study, we analyzed 40 cases AMI patients admitted to the Department of Cardiovasology, First people's Hospital of Jinan and 40 matched healthy volunteers between August, 2017 and November, 2019. All patients received no therapy prior to blood collection, and healthy volunteers had no medical history of

cardiovascular diseases. The diagnosis of AMI is based on the Third Universal Definition of Myocardial Infarction [21]. The inclusion criteria were as follows: Under these conditions any one of the following criteria meets the diagnosis for MI: Detection of a rise and/or fall of cardiac biomarker values [preferably cardiac troponin (cTn)] with at least one value above the 99th percentile upper reference limit (URL) and with at least one of the following: Symptoms of ischaemia; New or presumed new significant ST-segmenteT wave (STeT) changes or new left bundle branch block (LBBB); Development of pathological Q waves in the ECG; Imaging evidence of new loss of viable myocardium or new regional wall motion abnormality; Identification of an intracoronary thrombus by angiography. Venous blood was collected directly from patients with AMI and healthy volunteers upon admission, and serum samples were obtained by centrifugation. SERPINB1 level of serum samples were measured for receiver operating characteristic (ROC) curve analysis. All the experiments were approved by the Ethics Committee of the hospital, and informed consents were received from all participants.

The creation and treatment of AMI rat model

Adult male SD rats (200–220 g) were obtained from Medical Experimental Animal Center of Guangdong Province. The rat model of AMI was established by the left anterior descending (LAD) coronary artery. The rats were anesthetized by intraperitoneal injection of 50 mg/ kg pentobarbital sodium and underwent thoracotomy to expose cardiac tissue. Left main coronary artery was ligated with a 9–0 prolene suture at 1 mm below the ostium. Sham rats underwent the same surgical procedure without ligating LAD coronary artery.

In vivo transfection experiment, AMI rats were transfected with SERPINB1 overexpressed plasmid pcDNA3.1 (Thermo Fisher Scientific, Waltham, MA, USA) SER-PINB1 (AMI+pc- SERPINB1) or its negative control (AMI+pc-NC).

Cell culture and oxygen-glucose deprivation (OGD) treatment

Rat myocardial cell H9C2 was purchased from mibio (Shanghai, China). To simulate AMI, the H9C2 cells were cultured in non-serum and non-glucose medium at 37 °C for 0, 6, 12 or 24 h under 94% $N_2/5\%$ CO₂/1% O₂.

In vitro transfection experiment, H9C2 cells were transfected with pc-SERPINB1 or pc-NC. 48 h after transfection, the cells were exposed to OGD model. In vitro validation experiment, H9C2 cells were treated with pc-SERPINB1 and AMPK inhibitor Dorsomorphin (Dors). 48 h after treatment, the cells were exposed to OGD model.

Transthoracic echocardiography (TTE)

24 h after AMI modeling, rats were anesthetized by pentobarbital sodium (40 mg/kg) intraperitoneally. The TTE was used to assess the cardiac function of rats. The ultrasonic echocardiographic system was used to assess the rats' cardiac function after anesthetization. The left ventricular enddiastolic diameter (LVEDD) and left ventricular endsystolic diameter (LVEDD) were measured from three consecutive cardiac cycles. The ejection fraction (EF) and fractional shortening (FS) were calculated by the following equation: $EF = (LVEDV-LVESV)/LVESV \times 100\%$; $FS = (LVEDD-LVESD)/LVESD \times 100\%$. Then, rats were sacrificed by cervical dislocation to collect myocardial tissue for follow-up experiments.

Hematoxylin-eosin (H&E) staining

The pathological changes of myocardial tissues were detected by H&E staining. The rats were deeply anesthetized with an overdose of pentobarbital (100 mg/kg) intraperitoneally and then sacrificed via dislocation. The hearts of rats were harvested and rinsed with ice-cold PBS. The myocardial tissue was fixed with 4% paraformaldehyde and embedded in paraffin. Then the myocardial tissue was cut into 4 μ m sections. The sectioning was performed with H&E to evaluate the pathological changes of myocardial tissue. The stained sections were visualized using a light microscope at × 200 magnification.

TUNEL staining

The TUNEL staining was used to evaluate apoptosis of myocardial tissues. Myocardial tissue was immobilized with 4% paraformaldehyde and embedded in paraffin. Then tissues were sectioned into 4 μ m slices. After dewaxing and rehydration, TUNEL kit (Shanghai Ruisai Biotechnology Co., Ltd., Shanghai, China) was used strictly followed the method of the kit. Finally, hematoxylin was used for restaining, and the results were observed in five randomly selected visual fields under a light microscope at \times 200 magnification.

Flow cytometry assay

The cell apoptosis was detected by the FITC-Annexin V Apoptosis Detection Kit (Sigma Aldrich, St. Louis, MO, USA).H9C2 cells were washed with cold PBS and resuspended with 500 μ L binding buffer. Then the cells were stained with 5 μ L Annexin V-FITC and 10 μ L PI for 15 min in the dark. The cell apoptosis was measured on a flow cytometer (BD, Franklin Lakes, NJ, USA).

qRT-PCR

The expression level of SERPINB1 in myocardial tissues or cells was detected by qRT-PCR. TRIZOL regent was used to extract the total RNA of myocardial tissues and cells (Invitrogen, Carlsbad, CA, USA). The sequences were as follows: SERPINB1, 5'-CGGCCTGTCGGT TTTCAC-3' (forward) and 5'-TCTCACTCAACGCCA GGAAC-3' (reverse); GAPDH, 5'-GAAGGTCGGAGT CAACGGATT-3' (forward) and 5'-TTCCCGTTCTCA GCCATGT-3' (reverse). SuperScriptTM IV first-strand Synthesi System (Invitrogen, Carlsbad, CA, USA) was used to synthesize cDNA. The expression of SERPINB1 was detected by using the SYBR Green kit (Invitrogen, USA). PCR amplification was implemented as follows: 95 °C for 5 min, 40 cycles of 95 °C for 5 s, and 61 °C for 30 s. GAPDH was used as a reference gene for SER-PINB1 expression calculation. RNA expression was calculated by the $2^{-\Delta\Delta Ct}$ method.

Western blot

To obtain the total proteins, myocardial tissues or cells were lysed with RIPA lysate containing protease inhibitors (Thermo Fisher Scientific, Waltham, MA, USA). The protein was subjected to SDS-PAGE, and then was transferred onto PVDF membrane. The membranes were blocked by nonfat milk and then incubated at the temperature of 4 °C overnight with primary antibodies (Bax, 1:1000, 14,796; Bcl-2, 1:1000, 4228 s, Cell Signaling Technology, Danvers, MA, USA; SERPINB1, 1:1000, sc-293462; caspase-3, 1:1000, sc-271759, Santa Cruz Biotechnology, Santa Cruz, CA, USA; AMPK, 1:1000, SAB4502329, p-AMPK, 1:1000, SAB4503754; mTOR, 1:1000, T2949, p-mTOR, 1:1000, SAB4504476, Sigma Aldrich, St. Louis, MO, USA). Subsequently, membranes were incubated with secondary antibodies (Sigma Aldrich, St. Louis, MO, USA) properly under room temperature. The western blots were visualized with ECL detection reagents (GE Healthcare, Little Chalfont, Buckinghamshire, UK).

Statistical analysis

Data were shown as means \pm SD. Statistical analysis was performed with SPSS 23.0 (SPSS, Chicago, USA) and GraphPad Prism 7.0 (GraphPad Prism Software, Inc., San Diego, US). One-way analysis of variance test was used for comparison among groups, and unpaired Student's t-test was used for comparison between groups. The receiver operating characteristic (ROC) curve was used to analyze the clinical diagnostic value of the detection of serum SERPINB1. A P-value of<0.05 was considered statistically significant.

Results

The expression of SERPINB1 was down-regulated in AMI patients

We measured the expression of serum SERPINB1 in 40 AMI patients as well as 40 health controls by

qRT-PCR. The data suggested that SERPINB1 expression in serum of AMI patients was evidently lower than that of control (Fig. 1A, P < 0.05). Besides, we evaluated the diagnostic efficacy of SERPINB1 by ROC curve. The result of ROC curve indicated that the area under the curve (AUC) was 0.8644 (sensitivity = 85%, specificity = 72.5%, P < 0.01). Serum SERPINB1 might be a biomarker to predict AMI in a way.

Overexpression of SERPINB1 alleviated the impairment of myocardial function caused by AMI

We measured the expression of SERPINB1 after the establishment the AMI rat model by LAD. The data of gRT-PCR and western blot indicated that AMI modeling markedly decreased the expression of SERPINB1 both in serum and myocardial tissues of rats (Fig. 2A, B, P < 0.05). To explore the role of SERPINB1 on AMI, the SERPINB1 overexpressed plasmid pcDNA3.1 SER-PINB1 (pc-SERPINB1) was transfected into AMI rats. The data suggested that the SERPINB1 in myocardial tissues was significantly enhanced in AMI+pc-SERPINB1 group in comparison with AMI+pc-NC group (Fig. 2C, P < 0.05). The analysis results of TTE suggested that the myocardial function of rats in AMI group was remarkably destroyed compared with Sham group through the analysis of LVEDD, LVESD, FS, and EF (Fig. 2D, E, P < 0.05). The LVEDD and LVESD were remarkably increased by AMI modeling, while the FS and EF were significantly decreased (Fig. 2D, E, P < 0.05). Distinctly, overexpression of SERPINB1 reduced LVEDD and LVESD and enhanced FS and EF (Fig. 2D, E, *P* < 0.05).

Overexpression of SERPINB1 inhibited AMPK/mTOR pathway and relieved myocardial injury induced by AMI modeling

The H&E staining was used to detect the pathological changes of myocardial tissue after transfection with pc-SERPINB1. The results indicated that AMI modeling caused visibly myocardial injury, and the myocardial injury of AMI rats was markedly remitted by pc-SER-PINB1 (Fig. 3A). We detected the apoptosis of myocardial tissues by TUNEL staining. The TUNEL data suggested that the TUNEL-positive cells was markedly increased in AMI group in comparison of Sham group (Fig. 3B, P < 0.05). Overexpression of SERPINB1 could markedly reduce the number of TUNRL-positive cells compared with AMI + pc-NC group (Fig. 3B, P < 0.05). Besides, we measured the protein expression of Bax, Bcl-2, and caspase-3 by western blot. The results suggested that the protein expression of Bax and caspase-3 was remarkably higher and the Bcl-2 protein expression was lower in AMI group than that in Sham group (Fig. 3C, P < 0.05). SERPINB1 overexpression significantly reduced the protein expression of Bax and caspase-3, while enhanced Bcl-2 protein expression (Fig. 3C, P < 0.05).

OGD inhibited SERPINB1 expression and induced apoptosis of myocardial cells

The OGD cell model was constructed to inquire the influence of SERPINB1 on myocardial cells. The expression of SERPINB1 in H9C2 cells was detected by western blot after OGD treatment at 0, 6, 12 or 24 h. The data showed that OGD treatment reduced SERPINB1 expression and the inhibition of OGD on SERPINB1 expression was enhanced with the extension of treatment time (Fig. 4A, P < 0.05). As shown in Fig. 4B, the expression of SERPINB1 was significantly enhanced by pc-SERPINB1



SERPINB1. *P < 0.05



(P < 0.05). The CCK-8 assay was used to measured cell viability of H9C2 after pc-SERPINB1 transfection. The result suggested that OGD treatment suppressed cell viability of H9C2 cells, and the inhibiting effect of OGD on H9C2 cells could be remitted by SERPINB1 overexpression (Fig. 4C, P < 0.05). In addition, the cell apoptosis of H9C2 cells was detected by flow cytometry analysis. We found that the apoptosis of H9C2 cells was markedly increased by OGD treatment (Fig. 4D, E, P < 0.05). Overexpression of SERPINB1 inhibited cell apoptosis compared with OGD+pc-NC group (Fig. 4D, E, P < 0.05). The western blot result suggested that OGD treatment remarkably reduced protein expression of Bax and caspase-3 and enhanced Bcl-2 protein expression (Fig. 4F, P < 0.05). The protein expression of Bax and caspase-3 was markedly lower and the protein expression of Bcl-2 was significantly higher in OGD+pc-SERPINB1 group than that in OGD + pc-NC group (Fig. 4F, P < 0.05).

Overexpression of SERPINB1 activated AMPK/mTOR pathway

In this study, we found abnormal protein expression of p-AMPK/AMPK and p-mTOR/mTOR in myocardial cells H9C2 after OGD treatment. The AMPK/mTOR pathway was inhibited by OGD treatment. The western blot data indicated that OGD treatment markedly reduced the protein expression of p-AMPK/AMPK and enhanced the protein expression of p-mTOR/mTOR (Fig. 5, P < 0.05). To ascertain the effect of SERPINB1 on the AMPK/mTOR pathway, we measured the expression of p-AMPK/AMPK and p-mTOR/mTOR after pc-SERPINB1 transfection. The results indicated that overexpression of SERPINB1 activated the AMPK/mTOR pathway. The protein expression of p-AMPK/AMPK was increased and the protein expression of p-mTOR/mTOR was decreased in OGD + pc-SERPINB1 group compared with OGD + pc-NC group (Fig. 5, P < 0.05).



Inhibition of the AMPK/mTOR pathway eliminated the protective effect of SERPINB1 on H9C2 cells

In order to verify the role of AMPK/mTOR pathway on OGD-treated myocardial cells, we cultured H9C2 cells with the AMPK inhibitor Dors before OGD treatment. The data in Fig. 6A suggested that AMPK inhibitor Dors significantly inhibited the protein expression of p-AMPK/AMPK while enhanced p-mTOR/mTOR expression (P<0.05). CCK-8 result showed that Dors treatment reduced cell viability of H9C2 dells (Fig. 6B, P<0.05). Besides, we measured apoptosis of H9C2 cells by flow cytometry. The data suggested that cell apoptosis of OGD+pc-SERPINB1+Dors group was remarkably increased compared with OGD+pc-SERPINB1 group (Fig. 6C, P<0.05). In compaison with OGD+pc-SERPINB1 group, the protein expression of Bax and caspase-3 in OGD+pc-SERPINB1+Dors group was markedly increased, while Bcl-2 protein was decreased (Fig. 6D, P<0.05).

Discussion

AMI is one of the most serious cardiovascular diseases with high mortality despite improvements in treatment strategies over the past decade [4]. In our study, we







found that the SERPINB1 expression level was remarkably down-regulated in AMI patients, AMI rats as well as OGD-treated myocardial cells. AMI modeling resulted in obviously myocardial injury in myocardial tissue. The apoptosis rates of H9C2 cells increased by OGD treatment. Besides, the AMPK/mTOR pathway was inhibited in OGD-treated myocardial cells.

Abnormal SERPINB1 expression affects the progression of many diseases. Benarafa et al. have demonstrated that serpinB1 is vital for the maintaining the health of bone marrow pool in acute lung injury, knockdown of SERPINB1 increased the apoptosis and necrosis of purified bone marrow neutrophil [22]. SERPINB1 can promote the proliferation of porcine pancreatic stem cells (pPSCs) and play an important role in the transformation of pPSCs into insulin secreting cells [23]. Additionally, SERPINB1 gene knockdown aggravated lung injury of mice with orthotopic autologous liver transplantation (OALT), and administration of recombinant SERPINB1 protein attenuated cell apoptosis in the lung after OALT [9]. Moreover, Alpha-1-antitrypsin treatment in AMI mice increased LVEDD and LVESD, and reduced LVEF [24]. In this study, we found that serum SERPINB1 was significantly down-regulated in AMI patients. The result of ROC analysis indicated that the level of serum SER-PINB1 might be a biomarker for the diagnosis of AMI. We constructed AMI rat model and cell model by LAD method and OGD treatment, respectively. The expression levels of SERPINB1 in AMI rat serum and OGDtreated H9C2 cells were both markedly suppressed. These data indicated that SERPINB1 was involved in the procession of AMI. Overexpression of SERPINB1 significantly reduced the myocardial injury caused by AMI modeling. The number of TUNEL-positive cells in myocardial tissue of AMI rats was significantly reduced after overexpression of SERPINB1. Besides, overexpression of SERPINB1 markedly enhanced cell viability and reduced the apoptosis of H9C2 cells treated by OGD. The above results suggested that high expression of SERPINB1 could reduce myocardial apoptosis and alleviate myocardial damage that caused by AMI.

The AMPK/mTOR pathway is widely taken part in the development of numerous diseases, including ischemic acute kidney injury, diabetes, and cardiac dysfunction [25–27]. Study of Hu et al. has indicated that gAPN prevented cell apoptosis of chondrocytes induced by H2O2 through activation of AMPK/mTOR pathway [28]. Chen et al. have proved that AMPK activator AICAR strengthens the anti-apoptotic effect of Berberine on Müller cells stimulated with high glucose [29]. Overexpression of SIRT3 exhibited a protective role in rotenone-induced Parkinson's disease cell model by activating AMPK/mTOR pathway [30]. A study of Zhou et al. demonstrated

that exedin-4 inhibits cardiac hypertrophy by activating the AMPK/mTOR signaling pathway [31]. Apocynum leaf extract can marked increase the p-AMPK but decrease the mTOR protein expression to reduce blood lipid levels in rats with atherosclerosis and delay atherosclerotic progression [32]. Xu et al. have suggested that leonurine inhibited myocardial apo ptosis and improved myocardial function in rat model by activating the PI3K/ AKT/GSK3β pathway [33]. Yi et al. has found that inhibition of RhoA/ROCK pathway significantly mitigated AMI-induced myocardial apoptosis in mice [34].The analogical results were obtained in our study. We found that AMI modeling inhibited the AMPK/mTOR pathway and induced myocardial apoptosis. The p-AMPK level was remarkably reduced while p-mTOR level was markedly enhanced. SERPINB1 overexpression enhanced p-AMPK level, and increased p-AMPK level significantly reduced myocardial injury and myocardial apoptosis of AMI rats. The cell viability of OGD-treated H9C2 cells was significantly strengthened and the cell apoptosis was suppressed by activating p-AMPK. Besides, the AMPK inhibitor Dors accelerated cell apoptosis of H9C2 cells treated by OGD. These data suggested that activation of AMPK/mTOR pathway might improve myocardial damage by inhibiting myocardial apoptosis.

Conclusion

Overexpression of SERPINB1 alleviated myocardial damage induced by AMI in rats and apoptosis of OGDtreated H9C2 cells through activating AMPK/mTOR pathway. Our findings indicated that SERPINB1 might be a hopeful therapeutic target for AMI. However, we currently only explored the protective effect of SERPINB1 on AMI rats and cardiomyocytes. The role of SERPINB1 in human cardiomyocytes still needs further research. Additionally, extensive in vivo and in vitro studies are required before the results can be applied to clinical studies.

Abbreviations

AMI: Acute myocardial infarction; serpins: Serine protease inhibitors; OGD: Oxygen glucose deprivation; ROC: Receiver operating characteristic; TTE: Transthoracic echocardiography; LAD: Left anterior descending; AUC: Area under the curve; LVEDD: Left ventricular enddiastolic diameter; LVESD: Left ventricular endsystolic diameter; EF: Ejection fraction; FS: Fractional shortening; H&E: Hematoxylin–eosin; pPSCs: Porcine pancreatic stem cells; OALT: Orthotopic autologous liver transplantation.

Acknowledgements

Not applicable.

Authors' contributions

Conception and design: HLW; Perform research: HLW and JH; Data analysis and technical support: YC, SYC and YC; Manuscript writing: HLW. All authors read and approved the final manuscript.

Funding

Not applicable.

Availability of data and materials

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request. (E-mail: yingchencycy@163. com).

Declarations

Ethics approval and consent to participate

The experimental protocol of our study was performed in accordance with the Guide for the Care and Use of Laboratory Animals and approved by First people's Hospital of Jinan.

The protocol of this research has been approved by the Ethics Committee of First people's Hospital of Jinan. All patients have signed written informed consent.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no conflicts of interest.

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Received: 5 August 2021 Accepted: 3 January 2022 Published online: 15 March 2022

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