Hepatocyte growth factor (HGF) and insulin-like growth factor 1 (IGF-1) improved ventricular remodeling after myocardial infarction through inhibiting MMP8/13 in a rat model

Guangyi Tan^{1#}, Yao Chen^{2#}, Yin Huang³, Jierong Yao⁴, Yongquan Huang⁵, Jian Chen^{3*}, Wei Wu^{3**}

¹Department of Cardiology, Affiliated Nanhai Hospital of Southern Medical University, Foshan, China; ²Department of Intensive Care Unit, Affiliated Nanhai Hospital of Southern Medical University, Foshan, China; ³Department of Cardiology, The Fifth Affiliated Hospital of Sun Yat-sen University, Zhuhai, China; ⁴Department of Cardiology, Shantou Central Hospital (Affiliated Shantou Hospital of Sun Yat-sen University), Shantou, China; ⁵Department of Ultrasound, The Fifth Affiliated Hospital of Sun Yat-sen University, Zhuhai, China

Contributions: (I) Conception and design: W Wu, J Chen; (II) Administrative support: W Wu, J Chen; (III) Provision of study material or patients: G Tan, Y Chen; (IV) Collection and assembly of data: G Tan, Y Chen; (V) Data analysis and interpretation: G Tan, Y Chen, Y Huang, J Yao, Y Huang; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

"These authors contributed equally to this work.

*Second corresponding author.

**First corresponding author.

Correspondence to: Wei Wu, MD; Jian Chen, MD. Department of Cardiology, The Fifth Affiliated Hospital of Sun Yat-Sen University, No. 52 Meihua East Rd., Zhuhai 519000, China. Email: wuwei9@mail.sysu.edu.cn; chenjn@mail.sysu.edu.cn.

Background: Hepatocyte growth factor (HGF) and insulin-like growth factor 1 (IGF-1) have been reported on promoting the recovery after myocardial infarction (MI). We try to investigate the effect of HGF and IGF-1 on ventricular remodeling (VR) after MI and figure out how it works.

Methods: Forty-eight male SD rats were randomized to receive PBS, HGF, IGF-1 and GFs (a mix of HGF and IGF-1) by intramyocardial injection 2 weeks after MI. Echocardiography was performed before and 6 weeks after the injection. Then, histopathological examination performed. The mRNA and protein expression of matrix metalloproteinases 8/13 (MMP8/13), p38 MAPK and PI3K/Akt were measured.

Results: HGF, IGF-1 and GFs group indicated improving cardiac structure and function with dLVIDd, dLVIDs, dIVS and dLVPW decreased while dLVEF and dLVFS increased compared with those in PBS group (P<0.05, respectively). And HE staining showed that the myocardial cells in HGF, IGF-1 and GFs group arranged more regularly compared with PBS group. Masson staining showed that the area of myocardial fibrosis in PBS group expanded compared with HGF, IGF-1 and GFs group. The mRNA and protein expression of MMP8/13 in HGF, IGF-1 and GFs group were lower than PBS group (P<0.05, respectively). Further study showed that mRNA and protein expression of p38 MAPK in GFs and HGF group were higher than PBS group (P<0.05, respectively), while no significant difference was seen between them (P>0.05, respectively). Similarly, mRNA and protein expression of PI3K/Akt in GFs and IGF-1 group increased compared with PBS group (P<0.05, respectively). But there was no significant difference between them (P>0.05, respectively).

Conclusions: HGF and IGF-1 improved VR after MI by suppressing MMP 8/13, which may through p38 MAPK and PI3K/Akt pathway respectively.

Keywords: Hepatocyte growth factor (HGF); insulin-like growth factor 1 (IGF-1); Matrix metalloproteinases 8/13 (MMP8/13); myocardial infarction (MI); ventricular remodeling (VR)

Received: 09 August 2020; Accepted: 28 October 2020; Published: 30 June 2021. doi: 10.21037/jxym-20-95 View this article at: http://dx.doi.org/10.21037/jxym-20-95

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Introduction

Myocardial infarction (MI) is a severe type of coronary artery disease. With the activation of local inflammation and the stimulation of cytokines, the heart changes at the cellular and extracellular matrix (ECM) level, leading to ventricular remodeling (VR), eventually cause heart failure (1,2).

Beta blocker and angiotensin-converting enzyme inhibitor/angiotensin receptor blocker (ACEI/ARB) have been proved in improving VR and prognosis after MI, but they cannot completely prevent the occurrence of heart failure (3,4). In the last decade, stem cell therapy has become a hot spot in basic and clinical research (1-6). However, there are different conclusions about the effectiveness. Moreover, the potent to arrhythmia partly affect its clinical trials (7-10). It has been suggested that the potential benefits of stem cell therapy after MI may derive from a variety of cytokines secreted by the cells, including hepatocyte growth factor (HGF) and insulin-like growth factor 1 (IGF-1) (11,12). HGF and IGF-1 has been reported on promoting the recovery after MI (13-15). But whether combination of HGF and IGF-1 enhances the effect and how it works is still not fully understood.

In the process of myocardial fibrosis, matrix metalloproteinase (MMPs) play an important role (16). MMPs are a series of zinc ions dependency endogenous enzymes, which degrade the ECM components mainly (17). It has been reported that on the edge of infarcted area, the increasing expression of MMP8/13 had been observed, which indicates it may be related to VR process after MI. But the mechanism stayed unknown (18).

The application of MI SD rat model is an important means to study the scientific problems related to MI, and the conclusions come from which may have a good relevance to human biology. We present the following article in accordance with the ARRIVE reporting checklist (available at http://dx.doi.org/10.21037/jxym-20-95).

Objectives

Here we try to figure out how combination therapy of HGF and IGF-1 influence the expression of MMP8/13 after MI and what the possible mechanism may be.

Methods

Preparation for rat myocardial infarction model

Forty-eight male SD rats weighing 300-350 g were fasted all night except for free access to water. Then the rats were anesthetized with pentobarbital (concentration: 1%, 0.4-0.5 mL/100 g) by intraperitoneal injection. Endotracheal intubation was performed under a 14-gauge cannula mounted on a blunt and hollow needle with a 145° angled tip. The rats were mechanically ventilated with room air at a tidal volume of 0.5 mL/100 g body weight and a frequency of 100 breaths/min. The electrocardiogram Lead II was selected for continuous monitoring (BIOPAC MP150). A thoracotomy was performed via the left fourth intercostal space. Then, the pericardium was incised, and the left atrial appendage was elevated to expose the left anterior descending coronary artery. The left anterior descending coronary artery was ligated 2 mm below the left atrial appendage by using a 5/0 nylon suture. Successful occlusion was confirmed electrocardiographically by ST segment elevation. Drainage tubes were used to withdraw air or blood in the cavity while the chest is closed. The rats recovered from anesthesia and were then placed back into their cages for 2 weeks. Intramuscular injection of penicillin was administered post-operatively for prevention of infection.

Injection of cytokine solution

Two weeks after surgical intervention, the 34 surviving rats were anesthetized and orally intubated as described above. The rats were mechanically ventilated with room air, and a new thoracotomy was performed. The animals were randomized to receive phosphate buffer solution (PBS group, n=8), HGF solution (HGF group, concentration: 100 ng/mL, n=9), IGF-1 solution (IGF-1 group, concentration: 150 ng/mL, n=9) and a mix of HGF and IGF-1 (GFs group, concentration: HGF:100 ng/mL, IGF-1:150 ng/mL, n=8) by injection into the border of infarcted area. Successful injection was confirmed by the formation of a bled covering the infarcted zone. After recovering from anesthesia, the rats were placed back into their cages for 6 weeks. Prevention of postoperative infection was controlled as described above.

Table I List of prin	iers used in qK1-1 CK experiments	
Gene	Forward primer 5'-3'	Reverse primer 5'-3'
MMP8	TCCAGGTTACCCCACTAGCA	AGTGACTCTGCGACTGACAAG
MMP13	TGCATACGAGCATCCATCCC	AGCACTGAGCCTTTTCACCT
PI3K	AACACTTCCTCTGCATCTGGA	CTAGGTGACCTGACACAGCACT
p38 MAPK	CCAGCTTCAGCAGATAATGCG	TCATGGCTTGGCATCCTGTT
ACTB	TCAGCAAGCAGGAGTACGATG	GTGTAAAACGCAGCTCAGTAACA

 Table 1 List of primers used in qRT-PCR experiments

MMP8, matrix metalloproteinase 8; MMP13, matrix metalloproteinase 13; PI3K, phosphatidyl inositol 3-kinase; p38 MAPK, p38 mitogen activated protein kinase.

Ultrasonic cardiogram detection

Two weeks after the surgical intervention and 6 weeks after the PBS or cytokine solution injection, we performed ultrasonic cardiogram under 1% pentobarbital anesthesia (0.4–0.5 mL/100 g body weight) to detect the cardiac structure and function including left ventricular diastolic diameter (LVIDd), left ventricular systolic diameter (LVIDs), interventricular septum (IVS), left ventricular posterior wall (LVPW), left ventricular ejection fraction (LVEF) and left ventricular fractional shortening (LVFS). Then, the difference of each index between before and after cytokine injection treatment was assessed (dLVIDd, dLVIDs, dLVPW, dLVEF, dLVFS).

Tissue collection

After ultrasonic examination, the rats were killed by cervical dislocation and hearts were harvested. The left ventricular myocardium was cut in half through the center of the infarcted area. One half was immediately stored in -80 °C for further processing, and the other half for histopathological examination of hematoxylin and eosin (HE) staining and Masson staining. The infarcted region was visually identified by a mottled and pale discoloration. The border zone was myocardium which extended 1.0–2.0 mm from the infarcted area.

Western blotting analysis

Frozen border-zone samples were analyzed by Western blotting using the specific antibodies against MMP8 (1:1,000, Invitrogen, USA), MMP13 (1:1,000, Invitrogen, USA), p38 mitogen activated protein kinase (p38 MAPK, 1:1,000, Invitrogen, USA) and phosphatidyl inositol 3-kinase (PI3K, 1:1,000, Invitrogen, USA). To ensure equal loading, the same membranes were probed with GAPDH. Bands were visualized by enhanced chemiluminescence and quantified by the Quantity One Software (BioRad).

Real-time quantitative reverse transcription polymerase chain reaction (qRT-PCR)

Real-time qRT-PCR was performed according to the manufacturer's protocol. Total RNA was extracted from the border-zone with TRIzol reagent (Invitrogen, USA), and reversely transcribed to cDNA using M-MLV Reverse Transcriptase (Promega, USA). qRT-PCR was performed to measure the expression levels of candidate genes using a SYBR Green PCR Master mix (Toyobo, Japan). As an internal control, the expression of *ACTB* mRNA was estimated with the SYBR green system. The specific primer sequences of the selected genes are listed in *Table 1*.

HE staining

Samples taken from the border zone were routinely processed. Three sections of each heart were used for HE staining study. We examined the morphology of the myocardial cell, the inflammatory infiltration and the arrangement of cardiac muscle fibers. Each slide was examined under a microscope (Axio Scope A1, Zeiss).

Masson staining

Samples taken from the border zone were routinely processed. Three sections of each heart were used for Masson staining study. The fibrous tissue was stained blue. The area of stained fiber was determined by Image-Pro Plus 6.0 analysis system (Media Cybemetics, USA). Each slide was examined under a microscope (Axio Scope A1, Zeiss) and fibrous area was calculated by scoring 5 randomly-

					2										
	ū	roup PBS (n₌	=6)		G	roup HGF (n:	=5)		ũ	oup IGF-1 (n=	=6)		Group GFs ((u=6)	
raiailleters	Pre	Post	t	٩	Pre	Post	t	٩	Pre	Post	t P	Pre	Post	t	Ъ
LVIDd (mm)	8.38±0.61	8.80±0.72	2.68	0.044	8.72±0.85	8.00±0.67	-4.06	0.010	8.00±1.28	7.45±1.29	-3.74 0.01	3 9.05±0.73	3 7.36±0.72	-16.30	<0.001
LVIDs (mm)	5.79±0.44	6.45±0.75	2.79	0.038	6.61±0.74	5.74±0.50	-6.29	0.001	5.78±1.01	5.22±1.00	-3.37 0.02	0 6.81±0.74	l 5.34±0.67	-18.41	<0.001
IVS (mm)	1.32±0.08	1.42±0.09	3.31	0.021	1.37±0.06	1.18±0.07	-4.12	0.009	1.39±0.08	1.35±0.06	-0.86 0.42	9 1.38±0.05	i 1.25±0.07	-4.47	0.007
LVPW (mm)	1.42 ± 0.04	1.46±0.10	3.25	0.023	1.45±0.14	1.20±0.09	-7.11	0.001	1.43±0.13	1.39±0.07	-0.62 0.56	1 1.46±0.12	2 1.30±0.04	-4.63	0.006
LVEF (%)	63.50±2.26	58.50±6.89	-1.63	0.164	53.50±3.50	60.00±4.56	5.81	0.002	58.70±8.48	61.20±6.59	0.75 0.48	5 54.50±5.4	3 59.30±3.72	3.02	0.029
LVFS (%)	30.70±1.37	27.20±4.17	-1.82	0.129	24.00±2.00	30.80±5.67	3.62	0.015	27.80±3.89	30.20±3.76	1.28 0.25	6 25.00±3.2	9 27.80±2.71	2.43	0.059
MI, myocarc left ventricu fractional sh	Jial infarction; lar internal di ortening; Pre,	HGF, hepato mension sys , 2 weeks aft	ocyte g stole; N ter the s	growth fa VS, inter surgical	actor; IGF-1, rventricular s intervention;	insulin-like g eptum; LVP\ Post, 6 weel	rowth f N, left v <s after<="" td=""><td>actor 1; ventricu the PB5</td><td>GFs, a mix of lar posterior S or cytokine</td><td>of HGF and IC wall; LVEF, le solution injec</td><td>aF-1; LVIDc oft ventricul ction.</td><td>, left ventricu ar ejection fra</td><td>ar diastolic di action; LVFS,</td><td>iameter; left ven</td><td>LVIDs, tricular</td></s>	actor 1; ventricu the PB5	GFs, a mix of lar posterior S or cytokine	of HGF and IC wall; LVEF, le solution injec	aF-1; LVIDc oft ventricul ction.	, left ventricu ar ejection fra	ar diastolic di action; LVFS,	iameter; left ven	LVIDs, tricular

selected microscopic fields.

Statistical analysis

All quantitative data was described as mean \pm standard deviation (SD). If data met normal distribution and homogeneity of variance, one-way ANOVA test was performed in comparisons between groups. Otherwise, we chose Kruskal-Wallis instead. A value of P<0.05 was consider statistically significant. The tests were performed with SPSS 20.0 software.

Ethical statement

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). All experimental procedures were performed under licensed by the Animal Ethics Committee of the Fifth Affiliated Hospital of Sun Yat-sen University, in compliance with National Research Council guidelines for the care and use of animals (publication No. 85-23 revised 2011). No consent was required from patients since no patients were involved in this study.

Results

Six weeks after the PBS or cytokine solution injection, all surviving rats in each group {PBS group [6/8], HGF group [5/9], IGF-1 group [6/9] and GFs group [6/8]} were included in the statistical analysis.

GFs improved cardiac structure and function

Compared with PBS group, HGF, IGF-1 and GFs group indicated improving cardiac structure and function with dLVIDd, dLVIDs and dIVS decreased while the dLVEF and dLVFS increased (P<0.05, respectively). Besides, the decrease of dLVIDd and dLVIDs in GFs group was more significant compared with HGF group and IGF-1 group (P<0.05, respectively) (*Table 2* and *Figure 1*).

GFs decreased inflammatory infiltration and reduced infarcted area

The myocardial cells in the PBS group were swollen and disordered, with numerous infiltrating inflammatory cells. In HGF, IGF-1 and GFs group, myocardial cell arranged regularly, cytoplasm swelled lighter and inflammatory cell



Figure 1 Comparison of cardiac structure and cardiac function. (A,B) HGF, IGF-1 and GFs group showed decrease in the LVIDd and LVIDs after the treatment while PBS group increase. GFs group showed a more significant decrease while compared with HGF and IGF-1 group. (C,D) HGF, IGF-1 and GFs group showed decrease in the LVPW and IVS after the treatment while PBS group increase. No significant difference was seen among them. (E,F) HGF, IGF-1 and GFs group showed increase in the LVEF and LVFS after the treatment while PBS group decrease. No significant difference was seen among them. (E,F) HGF, IGF-1 and GFs group showed increase in the LVEF and LVFS after the treatment while PBS group decrease. No significant difference was seen among them. *, P<0.05. HGF, hepatocyte growth factor; IGF-1, insulin-like growth factor 1; GFs, a mix of HGF and IGF-1; LVIDd, left ventricular diastolic diameter; LVIDs, left ventricular systolic diameter; LVPW, left ventricular posterior wall; IVS, interventricular septum; LVEF, left ventricular ejection fraction; LVFS, left ventricular fractional shortening.

infiltration decreased compared with PBS group. The effect was more obvious in GFs group (*Figure 2A,B,C,D*).

Typical Masson staining results of each group were shown in *Figure 2E,F,G,H*. It can be seen that the infarcted myocardial tissue was stained light blue because collagen replaced normal myocardial cells, and the remaining normal myocardial tissue was stained red. In PBS group, the infarcted area ventricular wall thinned and left ventricular space expanded. Compared with PBS group, the area of myocardial fibrosis decreased in HGF and IGF-1 group, which was more obvious in GFs group (*Figure 2E,F,G,H*).

GFs reduced the expression of MMP8/13 and increased p38 MAPK and PI3K

In HGF, IGF-1 and GFs group, mRNA and protein expression of MMP8/13 significantly decreased on the border-zone as compared with those in PBS group (P<0.05, respectively). And GFs group further decreased protein expression of MMP8/13 compared with HGF and IGF- 1 group while no significant difference was seen between HGF and GFs group in mRNA level (P>0.05). Compared with PBS group, the mRNA and protein expression of p38 MAPK in GFs and HGF group increased (P<0.05, respectively) and no significant difference was seen between HGF and GFs group (P>0.05, respectively). Similarly, compared with PBS group, the mRNA and protein expression of PI3K in GFs and IGF-1 group increased (P<0.05, respectively) and no significant difference was seen between them (P>0.05, respectively) and no significant difference was seen between them (P>0.05, respectively) (*Figures 3,4*).

Discussion

The cardiac protective effects of HGF and IGF-1 have been greatly reported. They can improve cardiac structure and function through promoting angiogenesis and activating cardiac progenitor cells after MI (19-23). But what is the role about HGF and IGF-1 act on the ECM and weather HGF combined with IGF-1 have a synergistic effect are not fully understood. In our experiments, combination therapy



Figure 2 HE staining (A,B,C,D) and Masson staining (E,F,G,H) in each group. (A,E) PBS group showed myocardial swell with nuclear scaled up. A large number of inflammatory cells infiltrated. Hyperplasia of fibers were obvious but the arrangement was irregular. The fibrotic area was the largest in the four groups. (B,F) Compared with PBS group, myocardial swelled lighter, inflammatory cells infiltrated less and fibers arranged more regularly in HGF group. (C,G) IGF-1 group showed lighter myocardial swell than HGF group, but had a similar manifestation in myocardial fibrosis. (D,H) GFs group showed less fibrosis than HGF and IGF-1 group and fiber arranged more regularly. HGF, hepatocyte growth factor; IGF-1, insulin-like growth factor 1; GFs, a mix of HGF and IGF-1.



Figure 3 The mRNA and protein expression of MMP8/13 in each group. (A) Compared with PBS group, the mRNA expression of MMP8 in HGF, IGF-1 and GFs group was less. And GFs group could further decrease the expression than IGF-1 group. No significant difference was seen between HGF and GFs group. (B,C) Compared with PBS group, the protein expression of MMP8 in HGF, IGF-1 and GFs group was less. And GFs group could further decrease the expression than IGF-1 group. (D) Compared with PBS group, the mRNA expression of MMP13 in HGF, IGF-1 and GFs group was less while no significant difference was seen among the three groups. (E,F) Compared with PBS group, the protein expression of MMP13 in HGF and GFs group was less. And GFs group could further decrease the expression than HGF and GFs group was less. And GFs group could further decrease the expression of MMP13 in HGF and GFs group was less. And GFs group could further decrease the expression of MMP13 in HGF and GFs group was less. And GFs group could further decrease the expression of MMP13 in HGF and GFs group was less. And GFs group could further decrease the expression than HGF and IGF-1 group. *, P<0.05. MMP, matrix metalloproteinase; HGF, hepatocyte growth factor; IGF-1, insulin-like growth factor 1; GFs, a mix of HGF and IGF-1.

of HGF and IGF-1 after MI reduced the fiber deposition on the border zone and improved the cardiac function through suppressing the expression of MMP8/13, and the effect may be more remarkable than any single use to some extent.

Collagen is the main component of ECM in heart, among which, type I/III account for up to 90%. They play an important role in maintaining constitutional and functional integrity of heart. Collagen metabolism will be active after MI. In the infarcted zone, collagen degrade with myocytes necrosis, while on the border zone, collagen fibers deposit abnormally, leading to structure network destroy and making cardiac function decrease (24-26).

Decomposition of collagen is modified by MMPs family, a series of zinc irons which mainly participate in the metabolic activity of ECM. MMP8/13 can degrade type I/III collagen specifically (27). It has been reported that expression of MMP8/13 greatly increased on border zone after MI, prompting that it may have some pathologic significance in the remodeling process (28). In this study, we found that HGF and IGF-1 could both decrease the degree of fibrosis and improve myocardial fiber alignment on the border zone accompanied by the expression of MMP8/13 inhibited.

It is reported that MMPs can be regulated by various signal pathways, including MAPK and PI3K/Akt (29-31). MAPK is a highly conservative group of serine/threonine kinases, which plays an important role in the signal transduction process. As a classical pathway of MAPK, p38 MAPK involves in cellular activities of activation, migration and differentiation of HGF (32,33). PI3K consists of p85 and p110 subunits with the activity of serine/threonine kinase and phosphatidyl inositol kinase. It can cause cascade activation of Akt to participate in the cells of proliferation, differentiation and apoptosis. Studies have shown that IGF-1 can increase the expression of effector molecules, such as Bcl-2, α -actin, IGF and connexin-5 to inhibit cell apoptosis,

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Figure 4 The mRNA and protein expression of p38 MAPK and PI3K in each group. (A) Compared with PBS group, the mRNA expression of p38 MAPK in GFs and HGF group was more and no significant difference was seen between them. (B,C) Compared with PBS and IGF-1 group, the protein expression of p38 MAPK in HGF and GFs group was higher. And no significant difference was seen between GFs and HGF group. (D) Compared with PBS group, the mRNA expression of PI3K in IGF-1 and GFs group was more and no significant difference was observed between them. GFs group could further decrease the expression than HGF group. (E,F) Compared with PBS and HGF group, the protein expression of PI3K in GFs and IGF-1 group was higher. And no significant difference was seen between GFs and HGF group, the protein expression of PI3K in GFs and IGF-1 group was higher. And no significant difference was seen between GFs and HGF group, the protein expression of PI3K in GFs and IGF-1 group was higher. And no significant difference was seen between GFs and IGF-1 group, the protein expression of PI3K in GFs and IGF-1 group was higher. And no significant difference was seen between GFs and IGF-1 group. *, P<0.05. HGF, hepatocyte growth factor; IGF-1, insulin-like growth factor 1; GFs, a mix of HGF and IGF-1.

increase myocardial contractility and modify the collagen synthesis activity of fibroblasts (34). In this study, we found that HGF could increase the expression of p38 MAPK while IGF-1 increased the expression of PI3K/Akt. It indicated that HGF and IGF-1 may play a role in inhibiting VR through different pathways.

Conclusions

HGF and IGF-1 improve VR after MI through suppressing the expression of MMP8/13 on the border zone. It suggests that combination therapy of HGF and IGF-1 can play an even greater role than any single treatment to some extent. HGF and IGF-1 may suppress the expression of MMP8/13 through p38 MAPK and PI3K pathway respectively.

Acknowledgments

Funding: This work was supported by the Science and

Technology Planning Project of Guangdong province (grant No. 2014A020212088). The Major Program of Science and Technology Planning Project in Medical and Health of Zhuhai (grant No. 2015B1031).

Footnote

Reporting Checklist: The authors have completed the ARRIVE reporting checklist. Available at http://dx.doi. org/10.21037/jxym-20-95.

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at http://dx.doi. org/10.21037/jxym-20-95). All authors report grants from Science and Technology Planning Project of Guangdong province, grants from The Major Program of Science and Technology Planning Project in Medical and Health of Zhuhai, during the conduct of the study. The authors have no other conflicts of interest to declare.

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 conducted in accordance with the Declaration of Helsinki (as revised in 2013). All experimental procedures were performed under licensed by the Animal Ethics Committee of the Fifth Affiliated Hospital of Sun Yat-sen University, in compliance with National Research Council guidelines for the care and use of animals (publication No. 85-23 revised 2011). No consent was required from patients since no patients were involved in this study.

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doi: 10.21037/jxym-20-95

Cite this article as: Tan G, Chen Y, Huang Y, Yao J, Huang Y, Chen J, Wu W. Hepatocyte growth factor (HGF) and insulinlike growth factor 1 (IGF-1) improved ventricular remodeling after myocardial infarction through inhibiting MMP8/13 in a rat model. J Xiangya Med 2021;6:15. p38 in Il-1alpha induction of MMP-9 and MMP-13 in an established liver myofibroblast cell line. J Biomed Sci 2003;10:757-65.

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