

1 **LRRC8A critically regulates myofibroblast phenotypes and fibrotic remodeling**  
2 **following myocardial infarction**

3 **Running title: *LRRC8A Modulates Post-MI Fibrosis***

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## 21 Supplemental materials and methods

### 22 Animals and myocardial infarction models

23 The animal experiment procedures in the present study were approved by the  
24 Animal Care and Use Committee of the Fourth Military Medical University and  
25 adhered to the National Institutes of Health Guidelines for the Use of Laboratory  
26 Animals. *Lrrc8a*<sup>flox/flox</sup> mice were established by the clustered regularly interspaced  
27 short palindromic repeats (CRISPR)/Cas9 (Cyagen Biosciences, Guangzhou, China)  
28 and crossed with tamoxifen-inducible periostin-Cre (PostnMCM) mice (Jackson  
29 Laboratory, Maine, US). At 8-12 wk of age, mice were given a tamoxifen -contained  
30 chow diet (400 mg/kg, Harlan Teklad, US) for 28 d followed by a regular chow diet  
31 for another 15 d to ensure the clearance of tamoxifen from the animals [1]. The  
32 *Lrrc8a*<sup>flox/flox</sup>/PostnMCM/tamoxifen mice were conditionally knockout (CF-KO),  
33 whereas the *Lrrc8a*<sup>flox/flox</sup>/tamoxifen littermates represented wild-type (WT) controls.

34 To establish MI models, age-matched (8 to 12-wk-old) male WT and CF-KO  
35 mice were anesthetized by inhalation of isoflurane (1.5%, YiPin Pharmaceutical,  
36 Shijiazhuang, China) and left anterior descending coronary artery ligation operation  
37 was performed as we previously described [2]. To establish the MI models, the WT  
38 and CF-KO mice (n = 20 per group) were anesthetized by inhaling 1.5% isoflurane. A  
39 purse suture was placed over a tiny skin cut (approximately 1 cm) on the left chest.  
40 The 4th intercostal gap was disclosed when the pectoral major and minor muscles  
41 were dissected and retracted. To open the pleural membrane and pericardium, a small  
42 hole was cut at the 4th intercostal gap with a mosquito clamp. The heart was "popped  
43 out" smoothly and softly with the clamp slightly open. Using a 6-0 silk suture, the left  
44 coronary artery was located, sutured, and ligated at a point ~3 mm from its origin.  
45 When the anterior wall of the LV turned pale, the ligation was pronounced successful.  
46 Following ligation, the heart was immediately returned to the intra-thoracic space,  
47 followed by manual evacuation of air and closure of muscle and skin using the  
48 previously inserted purse-string suture.

49 In the peri-operation period and the following observational period of 28 d  
50 post-MI, the mortality rates were 45% (9/20) in the WT group and 25% (5/20) in the  
51 CF-KO group, respectively. At the indicated time points after MI, the animals were  
52 scarified and the hearts were harvested. The border zone of the MI-operated heart  
53 was defined as the immediate neighboring regions around the infarction as seen under  
54 the stereomicroscope. The remote zone contained areas ~2-3 mm away from the  
55 infarct region and the posterior wall of the left ventricle. Before the tissue collection  
56 and analysis, the border and remote zone were stereologically identified.

### 57 Echocardiography

58 Mice were anesthetized by the inhalation of 1.5% isoflurane. M-mode  
59 echocardiography was performed by a VisualSonics 770 echo system (FUJIFILM  
60 VisualSonics, Toronto, Canada) as we previously described [2]. Left ventricles (LV)  
61 were fully viewed in the short-axis between the two papillary muscles. Measurements  
62 were performed by the results of 3 continuous heartbeats. LV end-systolic diameters

63 (LVESD), LV end-diastolic diameters (LVEDD), LV end-systolic volume (LVESV),  
64 and LV end-distolic volume (LVEDV) were measured. Left ventricular ejection  
65 fractions (LVEF) and LV fractional shortenings (LVFS) were calculated as follows:  
66  $LVEF = [(LVEDV - LVESV) / LVEDV] \times 100\%$  and  $LVFS = [(LVEDD - LVESD) / LVEDD]$   
67  $\times 100\%$ .

#### 68 **Isolation and culture of adult mouse cardiac fibroblasts**

69 Adult cardiac fibroblasts were isolated from 8-12-wk-old WT mice. Hearts were  
70 rapidly excised and washed by ice-cold phosphatase saline buffer (PBS). The tissues  
71 were rinsed in ice-cold Hank's balanced salt solution and digested in type II  
72 collagenase (100 U/ml, Sigma Aldrich, Shanghai, China) at 37 °C for 20 min. The  
73 first digestion was discarded and the medium collected from the second digestion  
74 containing the cardiac fibroblasts was centrifuged for 5 min at 3000 rpm and  
75 resuspended in DMEM (Thermo Fisher, Shanghai, China) supplied with 10 % fetal  
76 bovine serum (FBS, Thermo Fisher, Shanghai, China). This procedure was repeated  
77 until the digestion medium was transparent. Cells were plated in 60 mm dishes and  
78 attached for 1 h before the first media change to remove non-adherent cells, including  
79 myocytes and endothelial cells.

#### 80 **Isolation of adult mouse cardiac myocytes**

81 Adult mouse cardiac myocytes were isolated as we previously described [3]. In  
82 brief, the mice were anesthetized with 1.5% isoflurane and fully heparinized with  
83 heparin. Hearts were carefully removed and placed into ice-cold PBS. Next, the heart  
84 was attached to the Langendorff system via the aorta and was fully perfused with  
85 perfusion solution (126 mmol/L NaCl, 4.4 mmol/L KCl, 18 mmol/L NaHCO<sub>3</sub>, 1  
86 mmol/L MgCl<sub>2</sub>, 11 mmol/L glucose, 10 mmol/L 2,3-butanedione monoxime, 30  
87 mmol/L taurine, and 4 mmol/L HEPES) for 5 min. The heart was then perfused by the  
88 collagenase solution (perfusion solution with 0.1% bovine serum albumin, 0.025  
89 mmol/l CaCl<sub>2</sub> and 0.1% type II collagenase) for another 10 min. After full digestion,  
90 the ventricle was triturated with a 10 ml pipette at a slow speed and filtered through a  
91 100 μm filter.

#### 92 **Isolation and culture of neonatal rat cardiac fibroblasts**

93 Neonatal cardiac fibroblasts were isolated from 1- to 3-d-old Sprague-Dawley  
94 rats. Briefly, the rat's heart was quickly removed and 0.04% collagenase II (Sigma  
95 Aldrich, Shanghai, China) was used to digest the minced ventricles 4 to 6 times. Cells  
96 were harvested and suspended in DMEM medium enriched with 10% FBS. Cardiac  
97 fibroblasts were isolated and collected by using the differential time adherent method.  
98 These cells were cultured in DMEM supplied with 10% FBS. Passage 2 cells were  
99 used for the following experiments.

#### 100 **Adenovirus transfection**

101 For adenoviral infection of cardiac fibroblasts, we employed  
102 replication-defective adenovirus vectors. The adenoviruses were constructed by

103 Hanbio Biotech, Shanghai, China. Adenovirus expressing GFP was used to control for  
104 nonspecific effects of adenoviral infection (Ad-Control). Two independent adenovirus  
105 vectors carrying 2 distinct siRNAs targeting *Lrrc8a* were transfected into fibroblasts  
106 to knock down LRR8A protein expression. The siRNA sequences were as follows:  
107 siRNA#1 5'-GCTGATGATTGCTGTCTTTGG-3'; siRNA#2  
108 5'-GCCTTCATGTTGCACCTCATC-3'. At 24 h after plating, cardiac fibroblasts  
109 were infected with the adenovirus vectors diluted in DMEM medium (MOI=100). The  
110 medium was replaced with virus-free SFM medium 24 h following adenoviral  
111 infection, and cells were cultured for an additional 12 h prior to experiments.

## 112 **Lentivirus transfection**

113 Lentiviruses carrying rat wild-type full-length *Lrrc8a* gene (LV-*Lrrc8a*WT),  
114 mutated rat *Lrrc8a* gene with deletion of its C-terminal LRRD (LRR8A del  
115 91/+26aa, LV-*Lrrc8a*ΔLRRD), and empty control (LV-Control) were constructed by  
116 Genechem Co.,Ltd., Shanghai, China. At 24 h after plating, cardiac fibroblasts were  
117 infected with lentiviruses diluted in DMEM medium (MOI=200). Polybrene (5 μg/ml)  
118 was added to enhance the transfection efficacy. The medium was replaced with  
119 virus-free SFM medium 24 h following adenoviral infection, and cells were cultured  
120 for an additional 48 h prior to experiments.

## 121 **Western blotting**

122 Tissue or cell lysates were collected in the RIPA buffer (P0013B, Beyotime,  
123 Beijing, China) in the presence of 1% protease/phosphatase inhibitor cocktails (#5872,  
124 Cell Signaling Tech, Shanghai, China). Proteins were separated on SDS-PAGE gels  
125 and transferred to polyvinylidene difluoride (PVDF) membranes (Merck Millipore,  
126 Darmstadt, Germany). Membranes were incubated overnight at 4 °C with primary  
127 antibodies. After washing by phosphate saline buffer with 0.1% tween 20, blots were  
128 incubated at room temperature for 1 h with horseradish peroxidase (HRP)-conjugated  
129 goat anti-mouse or goat anti-rabbit secondary antibodies (1:5,000, Cowin  
130 Biotechnology, Beijing, China). Proteins were visualized using an enhanced  
131 chemiluminescence kit (Merck Millipore, Darmstadt, Germany). The blot was  
132 scanned by the ChemiDocXRS equipment (Bio-Rad Laboratories, Shanghai, China),  
133 and densities of the bands were analyzed by Quantity One software (Bio-Rad  
134 Laboratories, Shanghai, China). The detailed information about primary antibodies is  
135 listed in **Table S1**.

## 136 **Real-time quantification PCR**

137 Total RNA was extracted from frozen tissues or cultured cells by the MiniBEST  
138 Universal RNA Extraction Kit (#9767, Takara, Dalian, China). RNA was reversely  
139 transcribed into cDNA using the PrimeScript™ RT reagent kit with gDNA Eraser  
140 (DRR047A, Takara, Dalian, China). RT-PCR was performed in triplicate using a PCR  
141 detection kit (DRR081A, Takara, Dalian, China) and a CFX96 system (Bio-Rad  
142 Laboratories, Shanghai, China). The primer sequences used in the present study are  
143 available in **Table S2**.

## 144 **RNA sequencing**

145 Cardiac fibroblasts were transfected with Ad-Control or Ad-*Lrrc8a* siRNA and  
146 treated with TGF- $\beta$ 1 (10 ng/ml) for 48 h. RNA library preparation, sequencing, and  
147 analysis were performed by Personalbio Tech Ltd. (Shanghai, China). A total of 3  $\mu$ g  
148 of RNA per sample was utilized for RNA library preparation with the NEBNext®  
149 UltraTM RNA Library Prep Kit for Illumina® (NEB, USA). The library was  
150 sequenced on an Illumina NovaSeq6000 platform. After quality control, reads were  
151 aligned to the reference genome using the STAR read mapper. Then, HTSeq v0.6.0  
152 software was used to count the read numbers mapped to each gene. Differentially  
153 expressed genes (DEGs) were identified using the DESeq2 package (Log<sub>2</sub>|fold  
154 change| $\geq$ 1 and *P* adj<0.05). Kyoto Encyclopedia of Genes and Genomes (KEGG)  
155 analysis in DEGs was performed. The detailed information about DEGs in the  
156 JAK-STAT Signaling Pathway is shown in **Table S3**. The whole RNA sequencing  
157 data is available in **File S2**.

## 158 **Immunofluorescence staining**

159 Cells were fixed with 4% paraformaldehyde for 10 min and permeabilized with  
160 0.4% Triton 100 (Beyotime, Beijing, China) in phosphate saline buffer (PBS) for 10  
161 min at room temperature. After being rinsed with PBS, cells were blocked with 2%  
162 bovine serum albumin (BSA) in PBS for 1 h at room temperature and incubated  
163 overnight with primary antibodies at 4 °C. Cells were then stained with secondary  
164 antibodies for 1 h following 4',6-diamidino-2-phenylindole (DAPI) staining for  
165 another 3 min at room temperature. Images were acquired with a confocal microscope  
166 (LSM880, Carl Zeiss, Germany) using the same exposure parameters. Negative  
167 controls stained with only secondary antibodies were included in all experiments.  
168 Images were analyzed using ImageJ. Fixed tissues were sectioned at a thickness of 5  
169  $\mu$ m following a series of deparaffinization and dehydration. Slides were then  
170 subjected to antigen retrieval in a hot citric acid buffer. After cooling, slides were  
171 permeabilized, blocked, incubated with primary antibodies, incubated with secondary  
172 antibodies, and stained DAPI as described before. Images were scanned by  
173 Panoramic MIDI (3D HISTECH) and analyzed using ImageJ. The detailed  
174 information of the primary antibodies are listed in **Table S1**.

## 175 **Co-immunoprecipitation**

176 Cardiac fibroblasts were transfected with LV-Control, LV-*Lrrc8a*WT, and  
177 LV-*Lrrc8a* $\Delta$ LRRD. Cells were harvested in a cell lysis buffer. After protein  
178 quantification, 500  $\mu$ g of protein was incubated with 4  $\mu$ g of antibodies against  
179 LRRC8A extracellular epitope (Alomone, Israel) overnight at 4 °C using a rotator  
180 followed by another 2-h incubation with Protein G Plus/Protein A agarose beads (Cell  
181 Signaling Technology, Shanghai, China). Beads were washed 3 times with ice-cold  
182 PBS, protein complexes were released by 5 min of boiling in SDS sample buffer,  
183 resolved by SDS-PAGE, and subjected to immunoblotting.

## 184 **Small interfering RNA (siRNA) transfection**

185 siRNA targeting rat *Grb2* and scramble control were designed, synthesized, and  
186 validated by Genechem Co.,Ltd. (Shanghai, China). *Grb2* siRNA sequence was the  
187 follows. GRB2 siRNA: 5'-GTACAAGGCAGAGCTTAATTT-3'. The siRNA  
188 transfection was performed using Lipofectamine RNAi<sup>MAX</sup> according to the  
189 manufacturer's instructions (Invitrogen) with minor modification. Briefly,  $5 \times 10^6$   
190 cells were transfected in 2 ml of SFM containing 500  $\mu$ l Opti-MEM (Invitrogen,  
191 Shanghai, China), 8  $\mu$ l Lipofectamine RNAi<sup>MAX</sup>, and 100 nmol/L of siRNA.

## 192 **Collagen gel contraction assay**

193 The ability of fibroblasts to contract collagen was assessed using the Cell  
194 Contraction Assay (#CBA-201, Cell Biolabs, San Diego, US) per the manufacturer's  
195 protocol. Fibroblasts were combined with a collagen solution, which was then  
196 allowed to polymerize. The collagen gel was detached from the plate and collagen gel  
197 size change (contraction index) was measured at various times and quantified with the  
198 ImageJ software.

## 199 **Masson trichrome staining**

200 Myocardium tissues were fixed with 4% paraformaldehyde for 24 h, dehydrated  
201 by increasing concentrations of ethanol, and embedded in paraffin. Tissue sections (5  
202  $\mu$ m) were stained with Masson trichrome staining kit (Sigma Aldrich, Shanghai,  
203 China) as the manufacturer's instruction described.

## 204 **Wheat germ agglutinin (WGA) staining**

205 Myocardium tissues were fixed with 4% paraformaldehyde for 24 h, dehydrated  
206 by increasing concentrations of ethanol, and embedded in paraffin. Tissue sections (5  
207  $\mu$ m) were stained with WGA staining kit (ServiceBio, Wuhan, China) as the  
208 manufacturer's instruction described.

## 209 **2,3,5-triphenyltetrazolium chloride (TTC) staining and the infarction size** 210 **measurement**

211 The heart was rapidly removed from the mouse and washed by ice-cold PBS. The  
212 heart was put on the solidified carbon dioxide for 15 min to freeze. The heart was cut  
213 into 1 mm thick slices along the atrioventricular sulcus from the apex to the bottom.  
214 The heart was cut into five pieces. The pieces were quickly placed into 5 ml 1.5%  
215 TTC buffer (pH = 7.4) for 15 min at 37 °C. TTC staining showed that the infarcted  
216 area was white and the non-infarcted area was red. The white/red area ratio at the  
217 papillary muscle surface was calculated by ImageJ software and recognized as the  
218 infarction size.

## 219 **References**

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222 ischemic heart. *Circulation*. 2014;130(5): 419-30.

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224 signaling contributes to cardiac inflammation, dysfunction, and remodeling following  
225 myocardial infarction. *Am J Physiol Heart Circ Physiol.* 2016; 310(2): H250-61.
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227 exacerbate myocardial ischemia/reperfusion vulnerability via enhancing  
228 GCN2/ATF6/PPAR- $\alpha$  pathway-dependent fatty acid oxidation. *Theranostics.* 2020;  
229 10(12):5623-40.

230 **Supplemental tables**231 **Table S1. Primary antibody information**

<b>Target</b>	<b>Supplier</b>	<b>Catalog</b>	<b>Usage</b>	<b>Dilution</b>
LRRC8A	CST	#24979	WB, IF	1:500 (WB), 1:50 (IF)
LRRC8A	Alomone Labs	AAC-001	IP	8 µg/mg protein lysate
α-SMA	CST	#19245	WB, IF	1:1000 (WB), 1:100 (IF)
p-JAK2	Abclonal	AP0531	WB	1:1000
JAK2	Abclonal	A19629	WB	1:1000
p-STAT3	Abclonal	AP0705	WB	1:1000
STAT3	Abclonal	A19566	WB	1:1000
GRB2	Abclonal	A19059	WB	1:1000 (WB)
Vimentin	CST	#5741	WB, IF	1:1000 (WB), 1: 200 (IF)
GAPDH	Abclonal	A19056	WB	1:3000

232

233 **Table S2. Primer sequence information**

<b>Gene name</b>	<b>Forward (5'-3')</b>	<b>Reverse (5'-3')</b>
<i>Lrrc8a</i>	ACTCCTGCAACGACTCCTT C	ACTGGTGTGCGATCTAGGTCA TAC
<i>Coll1a1</i>	AGGGCGAGTGCTGTGCTTT	CCCTCGACTCCTACATCTTC TGA
<i>Coll1a3</i>	TGAAACCCCAGCAAAACAA AA	TCACTTGCACCTGGTTGATAA GATT AA
<i>Col3a1</i>	CCCTGGACCTCAGGGTATC A	GGGTTTCCATCCCTTCCAGG
<i>Postn</i>	TGGTATCAAGGTGCTATCTG CG	AATGCCAGCGTGCCATAA
<i>Fn1</i>	ACCGAAGCCGGGAAGAGC AA	GGTCCGTTCCCCTGCTGAT TTAT C
<i>Ctgf</i>	CTGCGAGGAGTGGGTGTG	ATGTGTCTTCCAGTCGGTAG G
<i>Nppa</i>	GCTTCCAGGCCATATTGGAG	GGGGGCATGACCTCATCTT
<i>Nppb</i>	GAGGTCACCTCCTATCCTCTG G	GCCATTTCCCTCCGACTTTTC TC
<i>Myh7</i>	ACTGTCAACACTAAGAGGG TCA	TTGGATGATTTGATCTTCCA GGG
<i>Gapdh</i>	ACCACAGTCCATGCCATCA C	TCCACCACCCTGTTGCTGTA

234

235 **Table S3. Differentially expressed genes in JAK-STAT signaling pathway**  
 236 **between Ad-Control and Ad-*Lrrc8a* siRNA group**

Gene ID	Ad-Control	Ad- <i>Lrrc8a</i> siRNA	Fold change	P-value	Gene name
ENSRNO G0000000 0145	87.49134118	222.5110921	2.543235 583	6.24414 E-08	<i>Pik3r3</i>
ENSRNO G0000000 0875	577.6936815	1885.237021	3.263385 218	7.50245 E-32	<i>Fhl1</i>
ENSRNO G0000000 2919	3.193384588	14.68520432	4.598633 177	0.003405 065	<i>Gfap</i>
ENSRNO G0000000 5747	29.72161992	66.75308429	2.245943 675	0.000108 012	<i>Il27ra</i>
ENSRNO G0000001 1973	57.71639369	139.2792385	2.413165 993	8.7029E- 05	<i>Il7</i>
ENSRNO G0000001 5654	216.8865848	460.7569129	2.124414 072	3.43351 E-16	<i>Ghr</i>
ENSRNO G0000001 8962	32.92337412	68.94714647	2.094170 124	0.000891 445	<i>Ctfl</i>
ENSRNO G0000000 0187	3957.263326	1258.12006	0.317926 798	7.2251E- 25	<i>Csf2rb</i>
ENSRNO G0000000	8116.416682	3841.068393	0.473246 821	2.84147 E-26	<i>Cdkn1a</i>

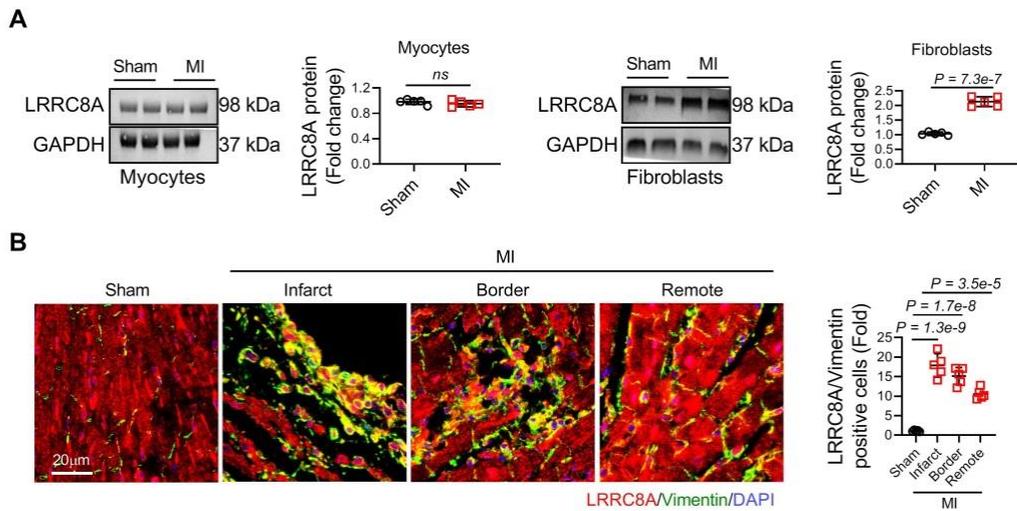
0521						
ENSRNO						
G0000000	1536.524992	484.8748693	0.315565	5.99642		<i>Pim1</i>
			885	E-42		
0529						
ENSRNO						
G0000000	21988.00063	8305.228416	0.377716	1.99581		<i>Pdgfra</i>
			399	E-48		
2244						
ENSRNO						
G0000000	3045.647011	559.0599277	0.183560	2.22442		<i>Socs3</i>
			316	E-61		
2946						
ENSRNO						
G0000000	119.1371115	7.981193771	0.066991	1.00039		<i>Il23a</i>
			668	E-13		
3254						
ENSRNO						
G0000000	469.0472515	180.1116552	0.383994	2.32761		<i>Il2rg</i>
			693	E-14		
3954						
ENSRNO						
G0000000	1207.876953	583.1332589	0.482775	1.7087E-		<i>Egfr</i>
			383	08		
4332						
ENSRNO						
G0000000	11.09136841	0	0	1.04763		<i>Il12b</i>
				E-05		
4380						
ENSRNO						
G0000000	101.1896102	5.054098141	0.049946	1.2434E-		<i>Il10</i>
			809	15		
4647						
ENSRNO						
G0000000	26.07415502	4.22553034	0.162058	4.42664		<i>Ifnb1</i>
			189	E-06		
6268						
ENSRNO						
G0000000	11574.44942	617.2348215	0.053327	1.81801		<i>Lif</i>

G0000000			359	E-41	
7002					
ENSRNO			0.013087	1.13335	
G0000000	1802.371939	23.58828411			<i>Csf3</i>
8525			356	E-32	
ENSRNO			0.285346	0.002514	
G0000000	20.19603388	5.762863427			<i>Il12a</i>
9468			294	319	
ENSRNO			0.045073	2.41099	
G0000001	3796.099222	171.1052725			<i>Il6</i>
0278			973	E-20	
ENSRNO			0.385796	0.019596	
G0000001	18.95411663	7.312426777			<i>Epor</i>
2619			232	396	
ENSRNO			0.482354	6.73212	
G0000001	5708.615595	2753.578143			<i>Il4r</i>
5441			802	E-11	
ENSRNO			0.427644	2.58507	
G0000001	1367.852512	584.9542406			<i>Pik3cb</i>
6384			235	E-35	
ENSRNO			0.495331	0.002762	
G0000001	57.12642453	28.29652209			<i>Pdgfb</i>
7197			579	768	
ENSRNO			0.032089	1.34293	
G0000001	515.1947199	16.53241114			<i>Il11</i>
7386			636	E-10	
ENSRNO			0.405628	0.000184	
G0000001	105.9141616	42.96179204			<i>Fmo5</i>
8076			401	539	

ENSRNO			0.401291	1.48365	
G0000001	1222.655804	490.6409729	166	E-17	<i>Stat5a</i>
9496					
ENSRNO			0.188857	1.00823	
G0000002	310.0408005	58.55363315	831	E-14	<i>Osm</i>
4390					
ENSRNO			0.054621	0.000758	
G0000002	16.97702344	0.927311193	542	054	<i>Il19</i>
5571					
ENSRNO			0.017602	3.53806	
G0000002	542.6096614	9.551301953	528	E-49	<i>Csf2</i>
6805					
ENSRNO			0.236180	6.47685	
G0000002	563.7920962	133.1564555	068	E-53	<i>Cish</i>
9543					
ENSRNO			0.449169	3.6108E-	
G0000003	9922.580262	4456.923336	794	26	<i>Osmr</i>
3192					
ENSRNO			0.436397	1.0512E-	
G0000003	660.5642148	288.2687792	814	23	<i>Ifnlr1</i>
3984					
ENSRNO			0.047333	4.36347	
G0000004	71.23140172	3.371603003	099	E-23	<i>Il2ra</i>
7647					
ENSRNO			0.363941	5.43669	
G0000005	21623.96397	7869.865721	862	E-57	<i>Jak2</i>
9968					

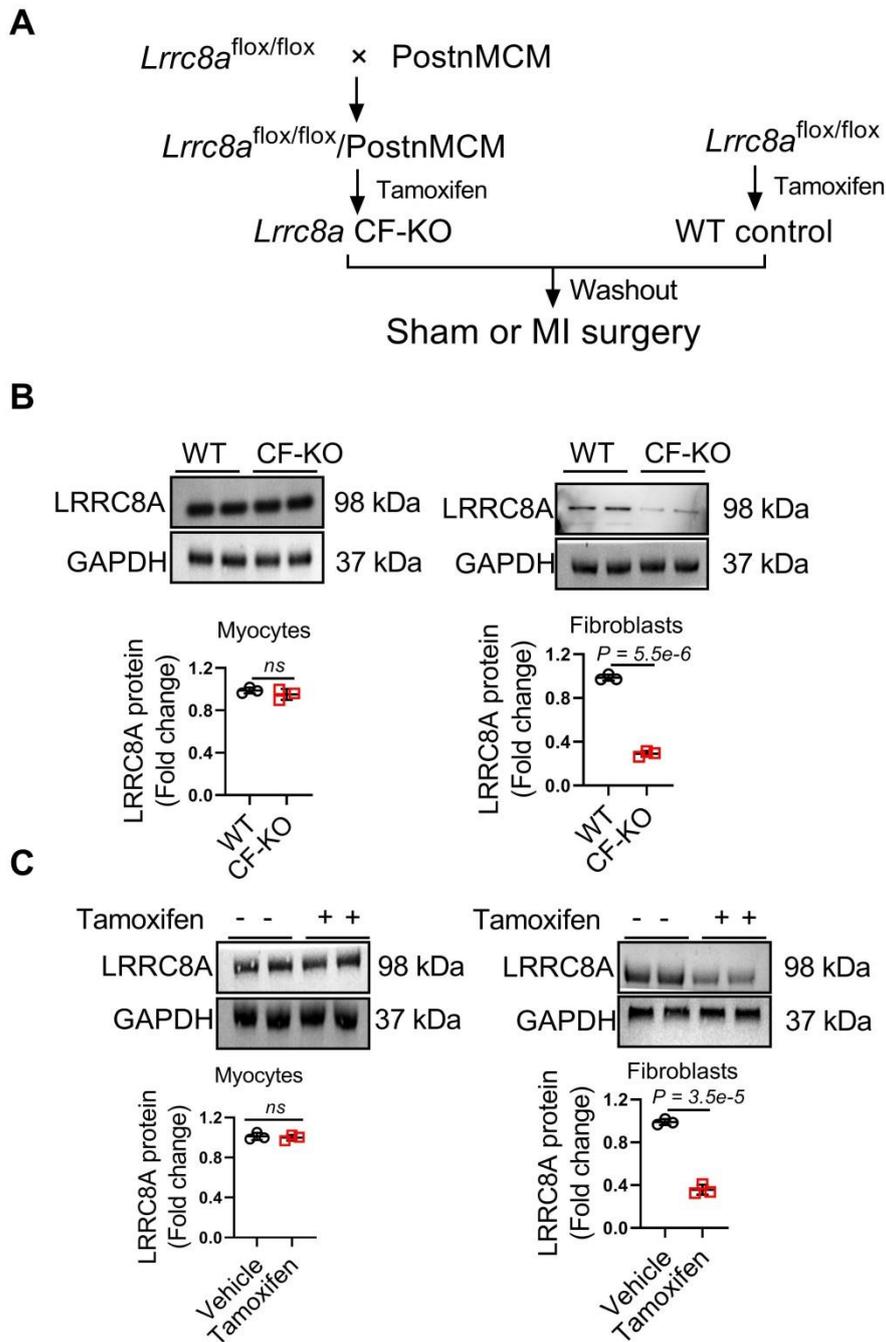
238 Supplemental figures and figure legends

239 Figure S1



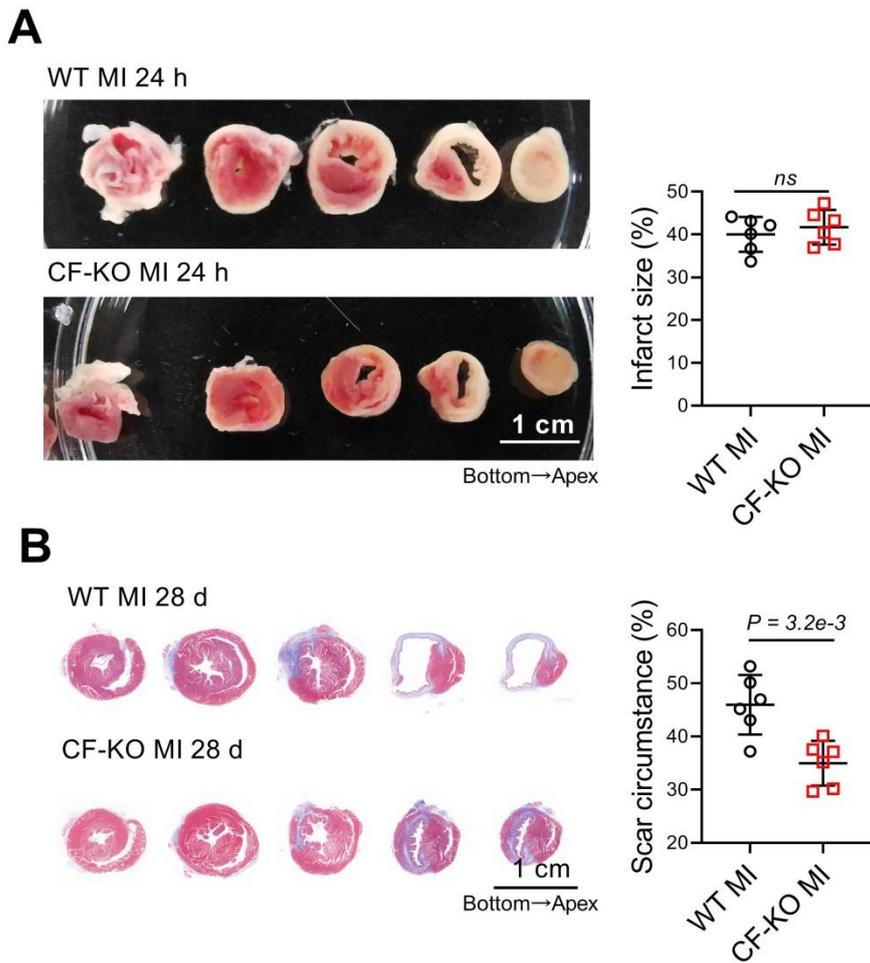
240

241 **Figure S1. LRRC8A expression was upregulated in cardiac fibroblasts in the**  
242 **infarcted heart.** (A) LRRC8A protein expression was measured in cardiac myocytes  
243 and fibroblasts isolated from the sham or MI heart on 28 d after the surgery. n= 5 mice  
244 per group. Data were analyzed by unpaired student's t test. (B) Representative images  
245 and quantification of LRRC8A (red) and vimentin (green) co-immunofluorescent  
246 staining in the infarct, border, and remote area of the infarcted heart. n = 5 mice per  
247 group. Data were analyzed by 1-way ANOVA followed by Boferroni post hoc test.



249

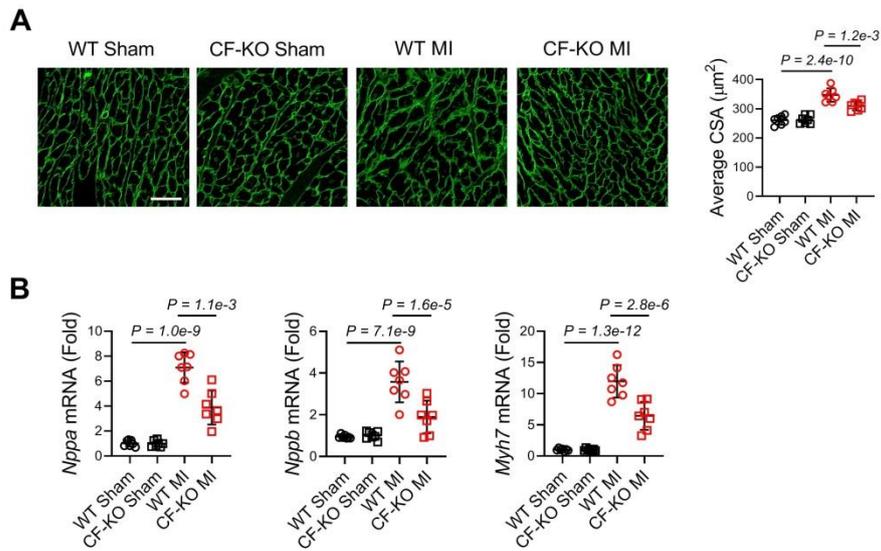
250 **Figure S2. Establishment and validation of the myofibroblast-specific LRRC8A**  
 251 **knockout mouse.** (A) Illustration of conditional myofibroblast-specific LRRC8A  
 252 knockout (CF-KO) mouse model establishment. (B) LRRC8A protein was measured  
 253 in cardiac myocytes and fibroblasts isolated from adult CF-KO or WT mouse hearts  
 254 after MI. (C) LRRC8A protein was measured in cardiac myocytes and fibroblasts  
 255 isolated from *Lrrc8a*<sup>flox/flox</sup>/PostnMCM mice with vehicle or tamoxifen treatment. n =  
 256 3 mice per group. Data were analyzed by unpaired student's t test.



258

259 **Figure S3. Acute infarct size and chronic fibrotic scar formation in WT and**  
 260 **CF-KO mice following MI.** (A) WT and CF-KO mice were subjected to MI  
 261 operation and the infarcted sizes were evaluated by triphenyl tetrazolium chloride  
 262 (TTC) staining 24 h post-MI. n= 6 mice per group. Data were analyzed by unpaired  
 263 student's t test. (B) Representative images of heart sections from the bottom to apex in  
 264 short-axis direction stained with masson trichrome on 28 d after MI. Scar  
 265 circumference was measured and expressed as a percentage of total circumference of  
 266 left ventricle. n = 6 mice per group. Data were analyzed by unpaired student's t test.

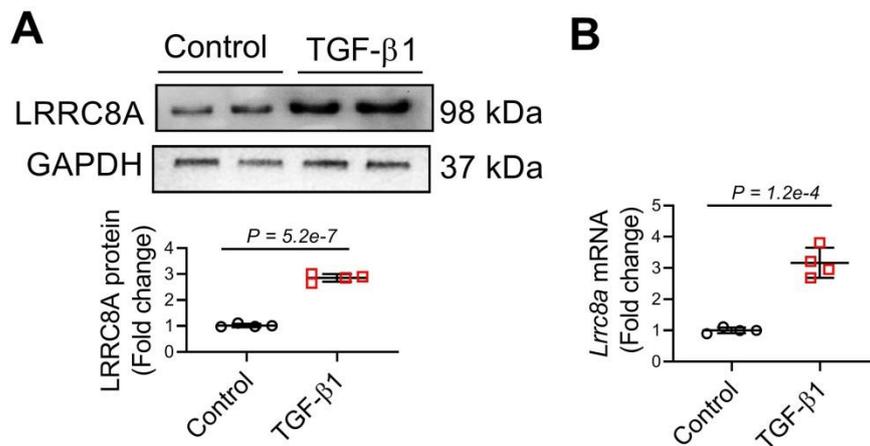
267 **Figure S4**



268

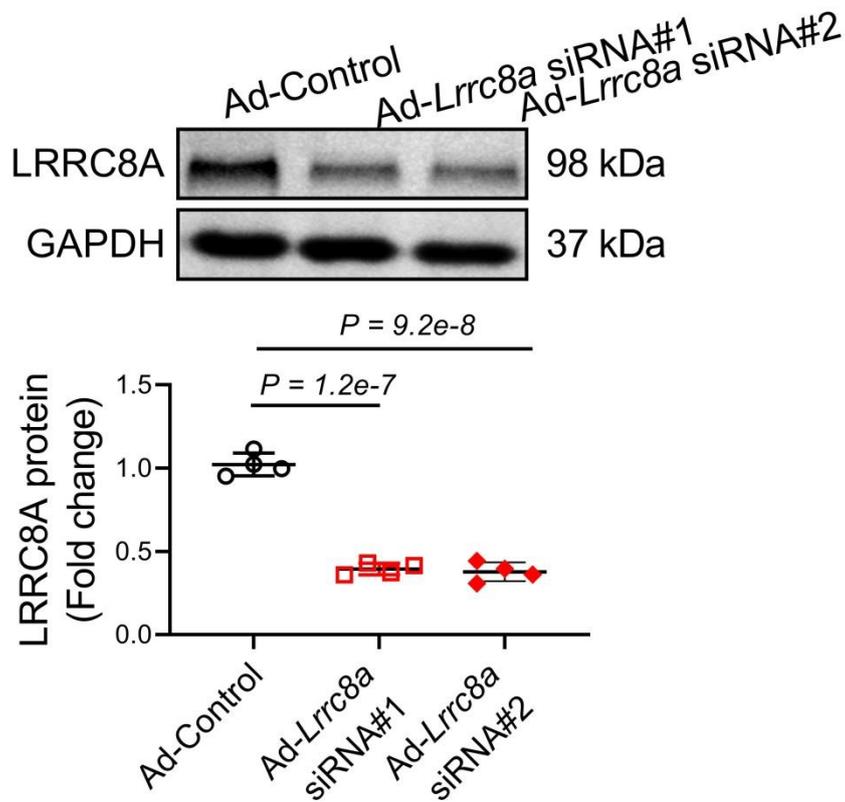
269 **Figure S4. Myocardial hypertrophy in WT and CF-KO mice after MI.** (A)  
 270 Representative images of wheat germ agglutinin (WGA) staining and quantification  
 271 results of cross-sectional areas (CSA) of cardiac myocytes. The average CSA were  
 272 calculated by the CSA of all cardiac myocytes in five randomly selected visual field  
 273 in the border zone on 28 d post-MI. (B) The mRNA levels of hypertrophic marker  
 274 gene including *Nppa*, *Nppb*, and *Myh7* were determined by RT-PCR.  $n = 7$  mice per  
 275 group. Data were analyzed by 1-way ANOVA followed by Bonferroni post hoc test.

276 **Figure S5**



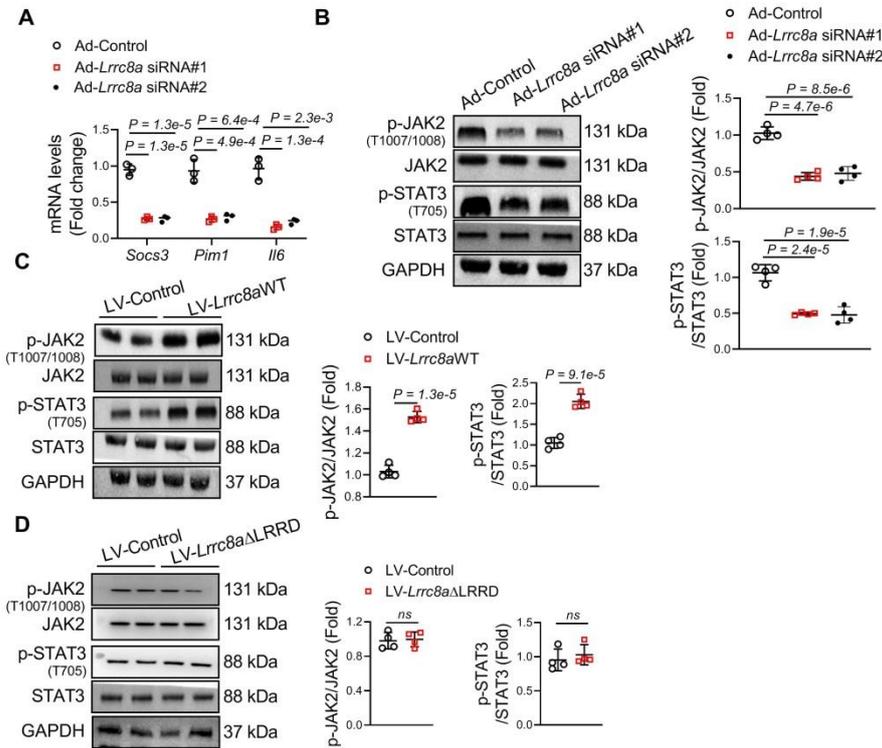
277

278 **Figure S5. TGF-β1 upregulated LRRC8A expression in cardiac fibroblasts.** The  
279 protein (A) and mRNA (B) levels of LRRC8A were determined in primary cardiac  
280 fibroblasts treated with vehicle or TGF-β1 (10 ng/ml) for 48 h. n = 4 individual  
281 experiments. Data were analyzed by unpaired student's t test.



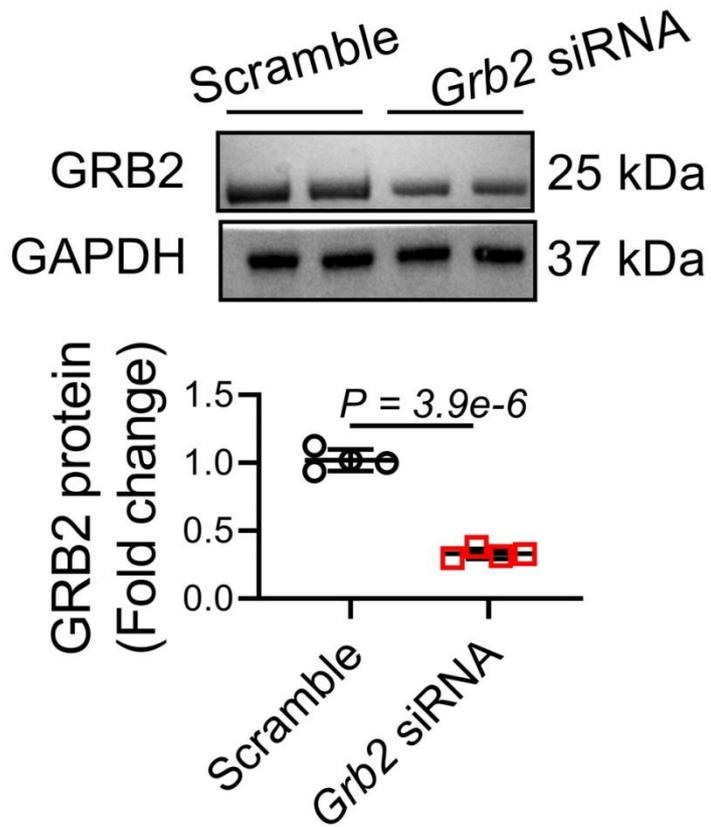
283

284 **Figure S6. Knockdown efficacy of *Lrrc8a* siRNA in cardiac fibroblasts.** Cardiac  
 285 fibroblasts were transfected with adenovirus vectors carrying two independent *Lrrc8a*  
 286 siRNAs (Ad-*Lrrc8a* siRNA#1 and #2) or empty control (Ad-Control). 24 h after the  
 287 transfection, LRRRC8A protein was measured by Western blotting. n = 4 individual  
 288 experiments. Data were analyzed by 1-way ANOVA followed by Bonferroni post hoc  
 289 test.



291

292 **Figure S7. The influence of LRRC8A knockdown or overexpression on the**  
 293 **JAK2-STAT3 signaling pathway in cardiac fibroblasts.** Cardiac fibroblasts were  
 294 transfected with adenovirus vectors carrying *Lrrc8a* siRNA (Ad-*Lrrc8a* siRNA#1 or  
 295 #2) or empty control (Ad-Control). (A) Upon TGF- $\beta$ 1 (10 ng/ml) treatment, mRNA  
 296 levels of STAT3 target genes including *Socs3*, *Pim1*, and *Il6* were measured by  
 297 RT-PCR. (B) Upon TGF- $\beta$ 1 (10 ng/ml) treatment, p-JAK2, JAK2, p-STAT3, STAT3,  
 298 and GAPDH were measured by Western blotting. (C) Cardiac fibroblasts were  
 299 transfected with lentivirus vectors carrying wild-type rat *Lrrc8a* gene (LV-*Lrrc8a*WT)  
 300 or empty control (LV-Control). Upon TGF- $\beta$ 1 (10 ng/ml) treatment, p-JAK2, JAK2,  
 301 p-STAT3, STAT3, and GAPDH were measured by Western blotting. (D) Cardiac  
 302 fibroblasts were transfected with lentivirus vectors carrying mutated *Lrrc8a* gene  
 303 without LRRD (LV-*Lrrc8a* $\Delta$ LRRD) or empty control (LV-Control). Upon TGF- $\beta$ 1 (10  
 304 ng/ml) treatment, p-JAK2, JAK2, p-STAT3, STAT3, and GAPDH were measured by  
 305 Western blotting. n = 3 to 4 individual experiments. Data were analyzed by unpaired  
 306 student's t test or 1-way ANOVA followed by Bonferroni post hoc test.



308

309 **Figure S8. Knockdown efficacy of *Grb2* siRNA in cardiac fibroblasts.** Cardiac  
310 fibroblasts were transfected with scramble or *Grb2* siRNA. 24 h after the transfection,  
311 GRB2 protein was measured by Western blotting. n = 4 individual experiments. Data  
312 were analyzed by unpaired student's t test.