Supplementary materials

The long non-coding RNA lncCIRBIL disrupts the nuclear translocation of Bclaf1 alleviating cardiac ischemia/reperfusion injury

Short title: LncCIRBIL regulates cardiac I/R injury via Bclaf1

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Supplementary Figure 1 The genomic location and sequence of lncRNA

NONMMUT058343



Sequence of NONMMUT058343.2 Chr6(108918953~108938686) 862nts TGCTTAACCAGTGAGCATAGAAATGGACGTGCTCTGAGTAAGTGTTTAACA GTCCTCTATCTAAGCTGCTTCTTCTGTGGCATGCATTACACAGGCACTGTAC TGAATGTCAGAGAGCAGGCATTTCGAACTGGAATTTCATGGTGTAAAAATA AAGGCCTTGTTATTGACTCTGAGAACCATGGGAACAAATTAAACACAGCTT AATAGCACAAAGGGGAATGATGGGGGACATTCGGAGCCTAATAAGGTGATCC ATGGGGTTCATTAGCACAGGCAGAGCTTATCAGAAGAGGCCTCAGAACCTA ATTAGCTGAACCCCCATCAAATAGATGCATTTGATTAATATGACTCCATTCTG AGTTTAAACATCAACATTAATTAGCCCCCATAATTGATAGATTTCCCAGCCATG TCAAGACAGGTTCTCTCTGGGTAGTCCAGACTGACCTCAAACTAACACTCT TCCAGCCTTGCTTTCAGGGATGCTGGGATTATAAGTATGTGCCACCATCTAC AACAAGATCTGTCTTAATATGTAAAGCCTTTGAAAACTGAAAACCTTAGGC CAATGAGACAGCTCCCTGGGCTCTTGGCATGGAAATCTGAAATCCTTAGAA ACACACACACACACACACACACACACACAATGAACAAGAAAAAGAAG AAGAAGGAGGAGGACGAGGAGGAGGACGAGGAGGATGAGGACAAGGAG

GAAGAAGAAGAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAA

Supplementary Figure 2. The conservative sequence of LncCIRBIL in human genome

A. The sequence in red is the longest one obtained when blasted LncCIRBIL sequence with human genome



C. Conservative sequence of human CIRBIL

AGTGCTTAACAGTCTAGCTAGGCAGCTTCTTCTATGACATGCATTATGCAGG CACTGTACTGAATTAGAGAGTGAATATTCCTAACTCCCACGACATGGTGTAA AAATAAAGGCCTAGTTATTGACTCTGAGAACCATGTGAACAAATTAAACAT GGCCATAATAGTGAAGGGGAAAGGTGGCAACAGTGAGATCTTAACACAGT GATCCATGGGGCTCATTAGCACGGAGTGGAGCTTATCAGAAGAGGTTTTCA GAACCTAATTAATTAGGCCCCCCATCAAATAGATGTATTCAATTAATATGACTC CATTCTGAGTTTGAACATCAACATTAATTAACCCTATAATTGATACATTTCCC ATCCATGTAAGAAACATTCATCACATTAATCAAAAAGGATC 407 nts



Supplementary Figure 3. The cardiac-specific lncCIRBIL Tg mice. The sequence of lncCIRBIL was cloned into the murine α -MHC promoter expression vector and the obtained DNA fragment containing lncCIRBIL driven by α -MHC promoter was microinjected into fertilized eggs.



Supplementary Figure 4. Overexpression of LncCIRBIL mitigated hypoxia/reoxygenation (H/R) induced injury of cultured cardiomyocytes. (A) The levels of lncCIRBIL in cardiomyocytes exposed to H/R. CTL N=9, H/R N=8 from 3 independent cultures. *P<0.05 versus control (CTL). P-values were determined by unpaired t test. (B) The levels of lncCIRBIL in cardiomyocytes exposed hypoxia. N=6 from 3 independent cultures. *P<0.05 versus control (CTL). P-values were

determined by unpaired t test. (C) The levels of lncCIRBIL in cardiomyocytes exposed to hydrogen peroxide (H₂O₂). CTL N=10, H₂O₂ N=11 from 3 independent cultures. *P<0.05 versus control (CTL). P-values were determined by unpaired t test. (D) The levels of lncCIRBIL in cardiomyocytes exposed to hypoxia with energy deprivation (low glucose). N=9 from 3 independent cultures. *P<0.05 versus control (CTL). P-values were determined by unpaired t test. (E) Verification of the expression of lncCIRBIL in the cardiomyocytes after transfection of lncCIRBIL plasmid. NC N=10, CIRBIL-OE N=12 from 3 independent cultures. *P<0.05 versus NC (negative control, empty plasmid). P-values were determined by unpaired t test. (F, G) Effects of lncCIRBIL on caspase-3 activity and cleaved caspase-3 protein level. N=12 from 3 independent cultures for caspase-3 activity, and N=3 for cleaved caspase-3 protein level. *P<0.05 versus NC; #P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (H) LncCIRBIL overexpression reduced the H/R-induced apoptosis by TUNEL staining (scale bar: 20µm). N=10 from 3 independent cultures. *P<0.05 versus NC; $^{\#}P$ <0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. Data represent the mean \pm SEM, n numbers are given in parentheses.



Supplementary Figure 5. The IncCIRBIL knockout mice. The

chr6_108868947-108888680 gene is located on mouse chromosome 6. Two exons have been identified. Exon 2 was selected as target site (sequences shown on the next page). Cas9 mRNA and gRNA generated by in vitro transcription were then injected into fertilized eggs for KO mouse productions. The founders were genotyped by PCR followed by DNA sequencing analysis. The positive founders were breeding to the next generation which was genotyped by PCR and DNA sequencing analysis.



Supplementary Figure 6. Downregulation of LncCIRBIL exacerbated hypoxia/reoxygenation induced cardiac myocyte injury. (A) LncCIRBIL in the cardiomyocytes after transfection lncCIRBIL siRNA. NC N=10, CIRBIL-SI N=12 from 3 independent cultures. *P<0.05 versus NC group. P-values were determined by unpaired t test. (**B**, **C**) Caspase-3 activity and cleaved caspase-3 protein level. N=12 from 3 independent cultures for caspase-3 activity, and N=3 for cleaved caspase-3 protein level. *P<0.05 versus NC; *P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (**D**) Apoptosis evaluated by TUNEL staining (scale bar: 20µm). N=10 from 3 independent cultures. *P<0.05 versus NC; *P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (**D**) Apoptosis



Supplementary Figure 7. The cardiac-specific Bclaf1 transgenic mice. A

transgene containing Bclaf1 was cloned downstream of the a-MHC promoter. A fragment containing the promoter and transgene was agarose gel-purified and used in a microinjection of the pronucleus of one-cell mouse embryos of C57BL/6 mice.



Supplementary Figure 8. Overexpression of Bclaf1 promoted apoptosis of cardiomyocytes subjected to hypoxia/reoxygenation. (A) Bclaf1 protein level in cardiomyocytes transfected with Bclaf1 overexpressing plasmids. N=3. *P<0.05 versus NC (negative control, empty plasmid). P-values were determined by unpaired t test. (B, C) Caspase-3 activity and cleaved caspase-3 protein level. N=12 from 3 independent cultures for caspase-3 activity, and N=3 for cleaved caspase-3 protein level. *P<0.05 versus NC, #P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (D) Cardiomyocyte apoptosis by TUNEL staining (scale bar: 20µm). N=10 form 3 independent cultures. *P<0.05 versus NC, #P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (E, F) The mRNA and

protein levels of p53 and Bax. N=9 for PCR and N=3 for Western blot, from 3 independent cultures. *P<0.05 versus NC, *P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. Data represent the mean ± SEM, n numbers are given in parentheses.



Supplementary Figure 9. Knockdown of Bclaf1 alleviated apoptosis of cardiac myocytes subjected to hypoxia/reoxygenation. (A) Bclaf1 protein level in cardiomyocytes transfected with its siRNA. N=3. *P<0.05 versus NC group. P-values were determined by unpaired t test. (B, C) Caspase-3 activity and cleaved caspase-3 protein level. N=12 from 3 independent cultures for caspase-3 activity, and N=3 for cleaved caspase-3 protein level. *P<0.05 versus NC, *P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (D) Cardiomyocyte apoptosis by TUNEL assay (scale bar: 20µm). N=10, from 3 independent cultures. *P<0.05 versus NC, *P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. (E, F) The mRNA and protein levels of p53 and Bax. N=9 for mRNA by qRT-PCR, and N=3 for protein expression by Western blot, from 3 independent cultures.

*P<0.05 versus NC, #P<0.05 versus NC-HR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc analysis. Data represent the mean ± SEM, n numbers are given in parentheses.



Supplementary Figure 10. p53 regulates the expression of IncCIRBIL. (A) The levels of IncCIRBIL in cardiomyocytes with p53 overexpression. N=9 from 3 independent cultures. *P<0.05 versus negative control (NC, empty plasmid). P-values were determined by unpaired t test. (**B**, **C**) The levels of IncCIRBIL in cardiomyocytes with p53 knockdown under normal and H/R conditions. N=6 from 3 independent cultures. *P<0.05 versus negative control (NC). (**D**) Effects of bclaf1 partial knockdown on the expression of LncCIRBIL. NC N=7, Bclaf1-pKO N=8, NC-I/R N=8, Bclaf1-pKO-I/R N=7. *P<0.05 versus NC; *P<0.05 versus NC-IR. P-values were determined by one-way ANOVA test followed by Bonferroni post hoc

analysis. Data represent the mean \pm SEM, n numbers are given in parentheses.

Supplementary Tables

Primer name	Primer sequence	
Bclaf1(mouse)	Forward: 5'- TTGGATGTGACAACTGCAACG-3'	
	Reward: 5'- GTGGCGGTAACGAATCTCCAG-3'	
Bax (mouse)	Forward: 5'- TGGAAGAAGATGGGCTGAGG-3'	
	Reward: 5'- TTCCCACCCCTCCCAATAAT-3'	
p53 (mouse)	Forward: 5'-TGGAGGAGTCACAGTCGGAT-3'	
	Reward: 5'- CAGTGAGGTGATGGCAGGAT -3'	
LncCIRBIL(mouse)	Forward: 5'-AAGAAGGTGGTGAAGCAGGC-3'	
	Reward: 5'-TCCACCACCCAGTTGCTGTA-3'	
LncCIRBIL(human)	Forward: 5'- TAGTGAAGGGGAAAGGTGGC-3'	
	Reward: 5'- AAGCTCCACTCCGTGCTAAT-3'	
Actin (mouse)	Forward: 5'-GACAGCAGTTGGTTGGAGCA-3'	
	Reward: 5'-TTGGGAGGGTGAGGGACTTC-3'	

Supplementary Table 1. Real-Time PCR sequence of primers

Antibody	Dilution	Company	Catalog Number
Bclaf1	WB 1:1000	Abcam,	Ab181240
	IF 1:500	UK	
Bax	WB 1:1000	Proteintech Beijing	Cat50599-2-1g
		China	
β-actin	WB 1:1000	Cell Signaling	# 4970
		Technology Boston USA	
Lamin-b	WB 1:500	Wanleibio Shenyang	WL01775
		China	
p53	WB 1:1000	Cell Signaling	# 2524
		Technology Boston USA	
Tublin	WB 1:1000	Absin	Abs830032
		China	

Supplementary Table 2. Information of antibodies used in the study