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Supplemental Information

miR-486 attenuates cardiac ischemia/reperfusion

injury and mediates the beneficial effect

of exercise for myocardial protection

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Supplemental Figures









Β

D



С

hiPSC-CM



cTNT+DAPI









0.5

0.0

HC SIRNA

围

SIPTEN 2 SIRMA

SiPTEN

miR-486 inhibitor

siPTEN?

SiPTEN

NC inhibitor



0.0

HC SIRNA

5iPTEN¹

NC inhibitor

SIPTENS

siPTEN.

miR-486 inhibitor

SIPTEN 2 SIRNA













α-SMA

*



α-SMA/EdU/Hoechst

Α

С



NC mimic

Smad1

GAPDH

Smad2

GAPDH

miR-486 mimic

60 KD

36 KD

60 KD

36 KD

Β



D





Smad1 Smad2























0



Supplemental Figure Legends

Figure S1. Preventive intervention by miR-486-AAV9 in acute cardiac ischemia/reperfusion injury. (A) Schematic diagram showing that AAV9 expressing miR-486 (miR-486-AAV9) or AAV9 controls (CTL-AAV9) were injected via tail vein, and 3 weeks later mice were subjected to cardiac ischemia/reperfusion (I/R) injury for 24 hrs. (B) RT-PCR for miR-486 expression in mice heart tissues at 24 hrs post cardiac I/R injury (n=6). (C) Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in mice heart tissues (n=6). (D) RT-PCR for CTGF, Col1a1, Col3a1, and BNP expressions in mice heart tissues (n=6). Data were compared by two-way ANOVA test followed by Tukey post hoc test. *, P<0.05; **, P<0.01; ***, P<0.001.

Figure S2. Preventive intervention by miR-486-AAV9 in chronic cardiac ischemia/reperfusion injury. (A) Schematic diagram showing that AAV9 expressing miR-486 (miR-486-AAV9) or AAV9 controls (CTL-AAV9) were injected via tail vein, and 1 week later mice were subjected to cardiac ischemia/reperfusion (I/R) injury for 3 weeks. (B) RT-PCR for miR-486 expression in mice heart tissues at 3 weeks post cardiac I/R injury (n=9-10). (C) Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in mice heart tissues (n=3). Data were compared by two-way ANOVA test followed by Tukey post hoc test. **, P<0.01; ***, P<0.001.

Figure S3. Long term protective effect of miR-486 overexpression against cardiac ischemia/reperfusion injury and cardiac dysfunction over 6 weeks. (A) Survival rate of mice injected with miR-486-AAV9 or CTL-AAV9 after 6 weeks of cardiac ischemia/reperfusion (I/R) injury (n=8,8,13,12 before I/R injury, n=8,8,10,12 at 6 weeks post I/R injury). (B) RT-PCR for miR-486 expression in mice heart tissues at 6 weeks post cardiac I/R injury (n=8-12). (C) Echocardiography for left ventricular

ejection fraction (EF, %) and fractional shortening (FS, %) in mice at 6 week after I/R injury (n=8-12). (D) Masson Trichrome staining for cardiac fibrosis in mice heart tissues (n=6-7). Scale bar=100 μ m. (E) RT-PCR for CTGF, Col1a1, Col3a1, and BNP expressions in mice heart tissues (n=8-12). Data were compared by two-way ANOVA test followed by Tukey post hoc test. **, *P*<0.01; ***, *P*<0.001.

Figure S4. Inhibition of miR-486 does not further aggravate cardiac ischemia/reperfusion injury. (A) The 2,3,5-triphenyltetrazolium chloride (TTC) staining for the infarct size at 24 hrs after cardiac ischemia/reperfusion (I/R) injury as determined by the infarct size/area at risk (INF/AAR) ratio. The area at risk/left ventricle weight (AAR/LV) ratio represents the homogeneity of surgery (n=6). (B) Schematic diagram showing that miR-486 sponge AAV9 or control AAV9 (CTL-AAV9) were injected via tail vein, and 1 week later mice were subjected to cardiac I/R injury for 3 weeks. (C) RT-PCR for miR-486 expression in mice heart tissues at 3 weeks post cardiac I/R injury (n=9-10). (D) Luciferase reporter assays performed in 293T cells co-transfected with negative control (NC mimic), miR-486 mimic or miR-210 mimic and the miR-486 binding site-carrying luciferase reporter plasmids (n=6). (E) Echocardiography for left ventricular ejection fraction (EF, %) and fractional shortening (FS, %) in mice at 3 week after I/R injury (n=9-10). (F) Masson Trichrome staining for cardiac fibrosis in mice heart tissues (n=6-7). Scale bar=100 µm. (G) Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in mice heart tissues (n=6). (H) RT-PCR for CTGF, Col1a1, Col3a1, and BNP expressions in mice heart tissues (n=9-10). Data between 2 groups were compared by unpaired two-tailed Student's t-test. Data among 3 groups were compared by one-way ANOVA test. Data among 4 groups were compared by two-way ANOVA test followed by Tukey post hoc test. *, P<0.05; ***, P<0.001.

Figure S5. Transfection of miR-486 mimic or inhibitor in cardiomyocytes *in vitro*. (A) RT-PCR for miR-486 expression in neonatal rat cardiomyocytes (NRCMs) transfected with miR-486 mimic, inhibitor, or negative controls (NC) (n=6). (B) Representative image of immunofluorescent staining for cardiac Troponin T (cTnT) which ensures the purification of human induced pluripotent stem cell-derived cardiomyocytes (hiPSC-CMs). Scale bar=100 μ m. (C) RT-PCR for miR-486 expression in hiPSC-CMs transfected with miR-486 mimic or negative control (n=4). (D) Relative hiPSC-CM numbers after oxygen glucose deprivation/reperfusion (OGDR) and transfection of miR-486 mimic (n=3). Data between 2 groups were compared by unpaired two-tailed Student's t-test. Data among 3 groups were compared by one-way ANOVA test. *, *P*<0.05; ***, *P*<0.001.

Figure S6. Regulation of PTEN and FoxO1 in human induced pluripotent stem cell-derived cardiomyocytes with miR-486 overexpression. RT-PCR for PTEN and FoxO1 in human induced pluripotent stem cell-derived cardiomyocytes (hiPSC-CMs) transfected with miR-486 mimic or negative control (NC mimic) (n=4). Data were compared by unpaired two-tailed Student's t-test.

Figure S7. Si-RNAs significantly downregulate PTEN or FoxO1 expression in neonatal rat cardiomyocytes. (A,B) RT-PCR for PTEN (A) or FoxO1 (B) expressions in neonatal rat cardiomyocytes (NRCMs) transfected with siRNAs targeting PTEN or FoxO1, respectively (n=4-6). Data were compared between PTEN or FoxO1 siRNA group and NC siRNA group using unpaired two-tailed Student's t-test. **, P<0.01; ***, P<0.001.

Figure S8. Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in oxygen glucose deprivation/reperfusion-treated neonatal rat cardiomyocytes transfected with miR-486 inhibitor and PTEN siRNA. (n=4). Data were compared by two-way ANOVA test followed by Tukey post hoc test. **, P<0.01; ***, P<0.001.

Figure S9. Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in oxygen glucose deprivation/reperfusion-treated neonatal rat cardiomyocytes transfected with miR-486 inhibitor and FoxO1 siRNA. (n=6). Data were compared by two-way ANOVA test followed by Tukey post hoc test. *, *P*<0.05; ***, *P*<0.001.

Figure S10. Overexpression of PTEN or FoxO1 attenuates the protective effect of miR-486 mimic against cardiomyocytes apoptosis. (A,B) RT-PCR for PTEN and FoxO1 in neonatal rat cardiomyocytes (NRCMs) transfected with plasmids expressing PTEN (A) or FoxO1 (B) (n=6). (C,D) TUNEL staining for α -Actinin-labelled NRCMs transfected with miR-486 mimic and plasmids expressing PTEN (C) or FoxO1 (D) in the condition of oxygen glucose deprivation/reperfusion (OGDR) treatment (n=4). Scale bar=100 µm. Data between 2 groups were compared by unpaired two-tailed Student's t-test. Data among 4 groups were compared by two-way ANOVA test followed by Tukey post hoc test. *, P < 0.05; **, P < 0.01; ***, P < 0.001.

Figure S11. Preventive delivery of miR-486-AAV9 regulates PTEN and FoxO1 in mice heart tissues. (A,B) RT-PCR (A) and Western blot (B) for PTEN and FoxO1 expressions in heart tissues from mice injected with miR-486-AAV9 before cardiac ischemia/reperfusion (I/R) injury. Heart tissues were harvested at 24 hrs post I/R injury (n=6). (C) Western blot for AKT phosphorylation levels in heart tissues from mice injected with miR-486-AAV9 before cardiac I/R injury. Heart tissues were harvested at 24 hrs post I/R injected with miR-486-AAV9 before cardiac I/R injury. Heart tissues were harvested at 24 hrs post I/R injected with miR-486-AAV9 before cardiac I/R injected with miR-486-AAV9 before cardiac I/R injected by two-way ANOVA test followed by Tukey post hoc test. *, P < 0.05; **, P < 0.01; ***, P < 0.001.

Figure S12. MiR-486 inhibits cardiomyocyte apoptosis through activating AKT and mTOR. (A,B) Western blot for Bax/Bcl-2 ratio and cleaved-Caspase3/Caspase3 ratio in oxygen glucose deprivation/reperfusion (OGDR)-treated neonatal rat cardiomyocytes transfected with miR-486 mimic in the presence or absence of AKT inhibitor MK2206 (A) or mTOR inhibitor Rapamycin (B) (n=6). Data were compared by two-way ANOVA test followed by Tukey post hoc test. *, P<0.05; **, P<0.01; ***, P<0.001.

Figure S13. MiR-486 inhibits cardiac fibroblast proliferation and activation. (A) RT-PCR for miR-486 in neonatal rat cardiac fibroblasts (NRCFs) transfected with miR-486 mimic, inhibitor, or negative controls (NC) (n=6). (B,C) Immunofluorescent staining for α -SMA/EdU in NRCFs transfected with miR-486 mimic (B), inhibitor (C), or NC (n=5). Scale bar=100 µm. Data between 2 groups were compared by unpaired two-tailed Student's t-test. Data among 4 groups were compared by two-way ANOVA test followed by Tukey post hoc test. *, *P*<0.05; **, *P*<0.01; ***, *P*<0.001.

Figure S14. MiR-486 negatively regulates Smad1 and Smad2 in cardiac fibroblasts. (A,B) RT-PCR for Smad1 and Smad2 in neonatal rat cardiac fibroblasts (NRCFs) transfected with miR-486 mimic (A), inhibitor (B), or negative controls (NC) (n=6). (C,D) Western blot for Smad1 and Smad2 in NRCFs transfected with miR-486 mimic (C), inhibitor (D), or NC (n=3). Data were compared by unpaired two-tailed Student's t-test. *, P < 0.05; **, P < 0.01; ***, P < 0.001.

Figure S15. Therapeutic delivery of cTnT-miR-486 AAV9 regulates PTEN and FoxO1 in mice heart tissues. (A) Immunofluorescent imaging for co-localization of ZsGreen (indicative of AAV9) and α -Actinin staining in heart tissues from mice injected with cTnT-miR-486 AAV9 or cTnT-control AAV9. Scale bar=50 µm. (B,C) RT-PCR (B, n=7-10) and Western blot (C, n=6) for PTEN and FoxO1 expressions in heart tissues from mice treated with cTnT-miR-486 AAV9 within 30 min post myocardial reperfusion. Heart tissues were harvested at 3 weeks post cardiac ischemia/reperfusion (I/R) injury. (D) Western blot for AKT phosphorylation levels in heart tissues from mice treated with cTnT-miR-486 AAV9 within 30 min post myocardial reperfusion. Heart tissues were harvested at 3 weeks post cardiac I/R injury (n=6). Data were compared by two-way ANOVA test followed by Tukey post hoc test. *, P<0.05; **, P<0.01; ***, P<0.001.

Figure S16. Expression of miR-486 in a mouse model of swimming exercise. (A) RT-PCR for the known factors in response to swimming exercise in heart tissues (n=6). (B) RT-PCR for circulating miR-486 levels in the serum from swimmed mice and sedentary mice (n=7-8). (C,D) RT-PCR for miR-486 expression in the gastrocnemius (C) and anterior tibialis (D) from swimmed mice and sedentary mice (n=3-6). Data were compared by unpaired two-tailed Student's t-test. *, P<0.05; **, P<0.01.

Figure S17. MiR-486 inhibition regulates fibrosis-associated gene markers and its downstream targets in swimmed mice upon cardiac ischemia/reperfusion injury. (A,B) RT-PCR for fibrosis-associated gene markers (A) and PTEN and FoxO1 (B) in mice heart tissues (n=6 for sedentary mice, n=11-12 for swimmed mice). Data were compared by two-way ANOVA test followed by Tukey post hoc test. ***, P<0.001.

Figure S18. The beneficial effect of swimming exercise against cardiac ischemia/reperfusion injury is attenuated in miR-486 knockout mice. (A) Schematic diagram showing that miR-486 knockout (KO) mice and wild type (WT) littermates were subjected to swimming exercise for 3 weeks before cardiac ischemia/reperfusion (I/R) injury for 24 hrs. (B) The 2,3,5-triphenyltetrazolium chloride (TTC) staining for the infarct size at 24 hrs after I/R injury as determined by the infarct size/area at risk (INF/AAR) ratio. The area at risk/left ventricle weight (AAR/LV) ratio represents the homogeneity of surgery (n=5). Data were compared by

unpaired two-tailed Student's t-test. **, P<0.01.

Supplemental Table

Gene	Primer sequence
rno_Pten forward	ATTGCCTGTGTGTGGTGA
rno_Pten reverse	TCCTCTGGTCCTGGTATGA
rno_FoxO1 forward	TGGGGCAACCTGTCGTA
rno_FoxO1 reverse	GGGCACACTCTTCACCATC
mmu_Pten forward	AGCCCTAACCCCAAGAAC
mmu_Pten reverse	ACAAGTCCCGATGAAACCT
mmu_FoxO1 forward	GTACAGCGCATAGCACCA
mmu_FoxO1 reverse	GCGACAGACAGAGTTCCC
hsa_Pten forward	CAGTCAGAGGCGCTATGTGT
hsa_Pten reverse	CACCTTTAGCTGGCAGACCA
hsa_FoxO1 forward	GGATGTGCATTCTATGGTGTACC
hsa_FoxO1 reverse	TTTCGGGATTGCTTATCTCAGAC
mmu_C/EBPβ forward	GGGGTTGTTGATGTTTTTGGT
mmu_C/EBPβ reverse	TCGAAACGGAAAAGGTTCTCA
mmu_CITED4 forward	CCTGGCATACGGCTCCTTC
mmu_CITED4 reverse	AGACTGCAGGTGCGTGCTAC
mmu_CPhar forward	CATGGATTTCTGGACCTCCTA
mmu_CPhar reverse	TTCATGGCTTTACAGCGT
rno_Smad1 forward	CGTGTTGGTGGATGGTTT
rno_Smad1 reverse	TGTGTCGCCTGGTATTTTC
rno_Smad2 forward	GTCAGTGCGATGCTCAAG
rno_Smad2 reverse	CTCAAGTGCTGTTTTCGCT
mmu_CTGF forward	TAAGACCTGTGGGATGGG
mmu_CTGF reverse	GCAGCCAGAAAGCTCAA
mmu_Col1a1 forward	TAAGGGTCCCCAATGGTGAGA
mmu_Col1a1 reverse	GGGTCCCTCGACTCCTACAT
mmu_Col3a1 forward	CTGTAACATGGAAACTGGGGAAA
mmu_Col3a1 reverse	CCATAGCTGAACTGAAAACCACC
mmu_BNP forward	GAGTCCTTCGGTCTCAAGGC
mmu_BNP reverse	TACAGCCCAAACGACTGACG