

**Activation of miR-34a-5p/Sirt1/p66shc pathway contributes to
doxorubicin-induced cardiotoxicity**

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Supplementary Data

Table 1 Primers used in real-time qRT-PCR

Gene	Sequence (5'- 3')	Product size (bp)
Sirt1	F, TACCCCATGAAGTGCCTCAA	195
	R, CCTTTTGTGTTCGTGGAGGT	
Bcl-2	F, GGGAGATCGTGATGAAGTACA	218
	R, GCTGAGCGCAGGCCAG	
BAX	F, CCAGCTCTGAACAGATCATG	201
	R, CAATCATCCTCTGCAGCTCC	
NF-κB P65	F, ACCATCATCACGCTGGAAGA	202
	R, TCTTTTGCTGGGGAGAGGAG	
P53	F, GGACAGCTTTGAGGTTTCGTG	202
	R, CCCACGGATCTTAAGGGTGA	
GAPDH	F, CAAGAAGGTGGTGAAGCAGG	200
	R, CCACCCTGTTGCTGTAGCC	

Table.2 Assessment of the cardiac function by echocardiography

Group	Control	Dox-4w	Dox+DEX-4w	Dox-8w	Dox+DEX-8w
LVAWd	2.04±0.14	1.76±0.27	1.85±0.06	1.52±0.29 ^{**}	1.91±0.12 ^Δ
LVAWs	2.96±0.17	2.39±0.43 [*]	2.49±0.18	2.06±0.39 ^{***}	2.60±0.19 ^Δ
LVIDd (mm)	5.53±0.44	7.66±0.62 ^{***}	7.07±0.90	7.63±0.74 ^{***}	7.21±0.82
LVIDs (mm)	3.31±0.48	5.52±0.35 ^{***}	4.74±0.53 [#]	6.02±0.61 ^{***}	4.81±0.37 ^{ΔΔ}
LVPWd	2.07±0.21	1.85±0.23	2.00±0.30	1.63±0.31 [*]	1.87±0.14
LVPWs	2.90±0.27	2.51±0.24 [*]	2.45±0.35	2.06±0.47 ^{**}	2.61±0.28 ^Δ

LVAWd, left ventricular anterior wall end-diastolic thickness, LVAWs, the systolic left ventricular anterior wall thickness, LVIDd, left ventricular internal dimension at end-diastole, LVIDs, left ventricular internal dimension at end-systole, LVPWd, the left ventricular posterior wall end-diastolic thickness, LVPWs, the left ventricular posterior wall end-systolic thickness, EF, ejection fraction. FS, fraction shrinkage. Data represent the mean±SD, ^{*} $p<0.05$, ^{**} $p<0.01$, ^{***} $p<0.001$ vs. control group, [#] $p<0.05$, ^{##} $p<0.01$, ^{###} $p<0.001$ vs. Dox-4w group, ^Δ $p<0.05$, ^{ΔΔ} $p<0.01$, ^{ΔΔΔ} $p<0.001$ vs. Dox-8w group.

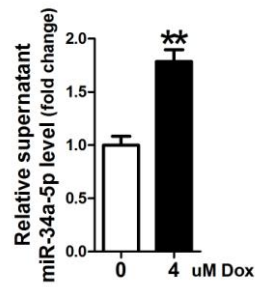


Figure 1. Determination of miR-34a-5p level in the supernatant of Dox-treated H9c2 cells. Level of miR-34a-5p in the supernatant of H9c2 cells exposed to 4 uM Dox was detected by RT-qPCR assay. Data are shown as mean $\pm$ sem. ** $p < 0.01$ vs 0 uM Dox control. $n = 3$.

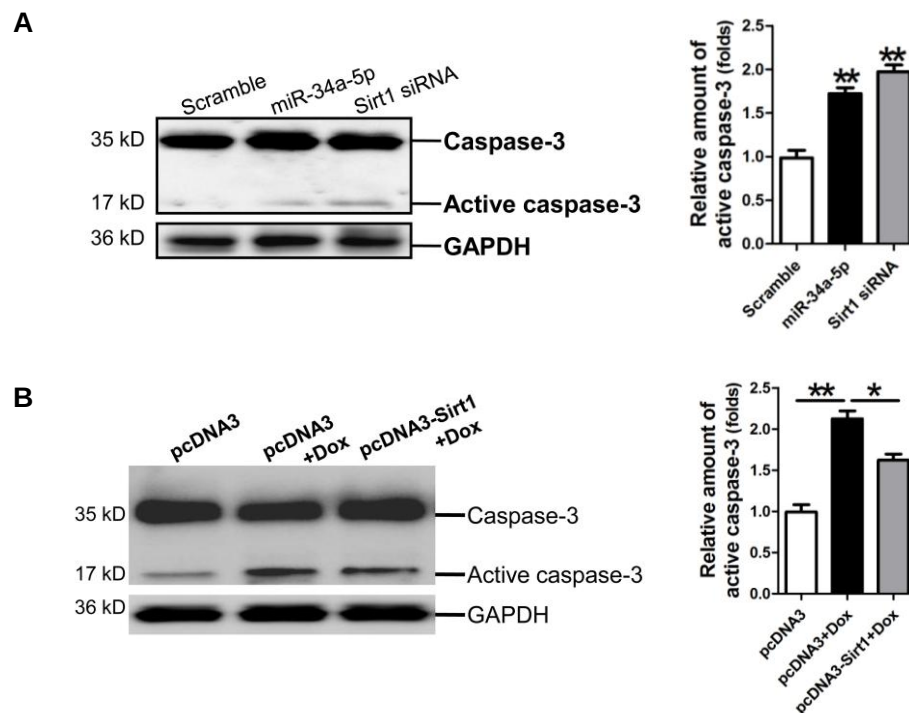


Figure 2. Determination of Caspase-3 protein level in H9C2 by Western blot assay. (A) H9c2 cells were transfected with miR-34a-5p mimic and Sirt1 siRNA, respectively. Data are shown as mean $\pm$ sem. ** $p < 0.01$ vs scramble group. $n = 3$. (B) H9c2 cells were treated by 4 uM Dox after transfection with pcDNA3-Sirt1. Data are shown as mean $\pm$ sem. * $p < 0.05$, ** $p < 0.01$. $n = 3$.