NADPH oxidase 2 mediates cardiac sympathetic denervation and myocyte autophagy, resulting in cardiac atrophy and dysfunction in doxorubicin-induced cardiomyopathy

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Supplementary Fig. 1. Changes in myocardial PGP9.5, GAP43, tyrosine hydroxylase (TH) and noradrenaline transporter (NET) proteins in WT saline and Nox2 KO saline groups. A, B, E, F. Representative Western blots of PGP9.5, GAP43, TH and NET proteins, respectively. Equal loading of proteins is illustrated by GAPDH bands. C, D, G, H. The graphs show the relative expression of PGP9.5, GAP43, TH and NET in the two groups, respectively. Values are presented as means \pm S.E.M.; n = 5 for C; n = 6-7 for D; n = 6-8 for G; n = 5-6 for H. Comparisons between two groups were used unpaired Student's t-test.



Supplementary Fig. 2. Changes in myocardial ERK, p-ERK, Akt, p-Akt, S6 and p-S6 proteins in WT saline and Nox2 KO saline groups. A. Representative Western blots of ERK, p-ERK, Akt, p-Akt, S6 and p-S6 proteins. Equal loading of proteins is illustrated by GAPDH bands. B-G. The graphs show the relative expression of ERK, p-ERK, Akt, p-Akt, S6 and p-S6 proteins in the two groups, respectively. Values are presented as means \pm S.E.M.; n = 5. Comparisons between two groups were used unpaired Student's t-test.



Supplementary Fig. 3. Changes in myocardial LC3 II and Beclin1 proteins in WT saline and Nox2 KO saline groups. A. Representative Western blots of LC3 II and Beclin1 proteins. Equal loading of proteins is illustrated by GAPDH bands. B, C. The graphs show the relative expression of LC3 II and Beclin1 proteins in the two groups, respectively. Values are presented as means \pm S.E.M.; n = 5-6 for C, n = 5 for D. Comparisons between two groups were used unpaired Student's t-test.

Supplementary Methods

Eight-week old male C57BL/6 J mice were randomly divided into 4 experimental groups as follow: mice in control group (n=9) received normal saline (NS, equal volume, i.p.), mice in chloroquine group (n=8) received intraperitoneal injection of chloroquine (Sigma-Aldrich, St. Louis, MO) at a dose of 10 mg/kg (once a day for 7 days, i.p.), mice in doxorubicin group (n=10) received intraperitoneal injection of doxorubicin (Cayman Chemical, Ann Arbor, Michigan) at a single dose of 15 mg/kg, and mice in doxorubicin plus chloroquine group (n=8) received intraperitoneal injection of doxorubicin at a single dose of 15 mg/kg and 10 mg/kg of chloroquine (beginning 1 hour before doxorubicin injection, once a day for 7 days, i.p.). The doses of doxorubicin and chloroquine were based on our pilot studies and previous reports (Fu et al. Circ Res 2016 March 4;118(5):798-809; Kanamori et al. Am J Pathol 2013 March;182(3):701-13).



Supplementary Fig. 4. Changes in myocardial LC3 II protein in mice with saline (NS), chloroquine (CQ), doxorubicin (Dox) and Dox plus CQ treatments. A. Representative Western blot of LC3 II protein. Equal loading of proteins is illustrated by GAPDH bands. B. The graph shows the relative expression of LC3 II protein in the four groups, respectively. Values are presented as means \pm S.E.M.; n = 8-10. *P < 0.05 vs. NS group. #P<0.05 vs. Dox group. Comparisons among groups were performed by one-way ANOVA followed by a Newman-Keuls test for multiple comparisons.



Supplementary Fig. 5. Changes in myocardial Bcl-2, Bax, caspase 3 and cleaved caspase 3 proteins in WT saline, WT doxorubicin (Dox) and Nox2 KO Dox groups. A. Representative Western blots of Bcl-2, Bax, caspase 3 and cleaved caspase 3 proteins. Equal loading of proteins is illustrated by GAPDH bands. B-F. The graphs show the relative expression of Bcl-2, Bax, caspase 3 and cleaved caspase 3 in the three groups, respectively. Values are presented as means \pm S.E.M.; n = 5 for B, C, D; n = 6 for E, E. *P < 0.05 vs. WT saline group. #P<0.05 vs. WT Dox group. Comparisons among groups were performed by one-way ANOVA followed by a Bonferroni post hoc test for multiple comparisons.



Supplementary Fig. 6. Changes in cardiac fibrosis in WT and Nox2 KO mice with saline or Doxorubicin (Dox) treatment. A. The representative photomicrographs of left ventricular myocardium with Masson trichrome staining. Bar = 50 μ m. B. The graph shows the quantification of cardiac fibrosis as measured using ImageJ. Values are presented as means ± S.E.M.; n = 5. *P < 0.05 vs. WT saline group. #P<0.05 vs. WT Dox group. Comparisons among groups were performed by one-way ANOVA followed by a Bonferroni post hoc test for multiple comparisons.

Full-length blots for Figure 4 A



Full-length blots for Figure 4 B

GAP43 (sc33705) GAP43 43KD Lane # 23 456789 1 A55NN A55LL A66RL A64NN A63LU A56RU A55RL A55RL Animal ID WT Dox WT saline Nox2 KO Dox GAPDH (ab8245)

GAPDH ——		36
Lane #	1 2 3 4 5 6 7 8 9	
Animal ID	A55NN A55LL A66RL A63LU A63LU A55RL A55RL	
	WT Dox	
	Doy	

Full-length blots used for Figure 4 B.

Area shown in figures are indicated by boxes.

Full-length blots for Figure 4 E

Tyrosine hydroxylase (TH)





Full-length blots used for Figure 4 E. Area shown in figures are indicated by boxes.

Full-length blots for Figure 4 F

Noradrenaline transporter (NET)



Full-length blots used for Figure 4 F. Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 A

ERK (sc1647)



Full-length blots used for Figure 5A. Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 B





Full-length blots used for Figure 5B. Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 E



Full-length blots used for Figure 5E. Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 F



Full-length blots used for Figure 5F.

Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 I



Full-length blots used for Figure 5I. Area shown in figures are indicated by boxes.

Full-length blots for Figure 5 J



Full-length blots used for Figure 5J. Area shown in figures are indicated by boxes.

Full-length blots for Figure 6 C

LC3A/B (CST4108)



Full-length blots used for Figure 6C. Area shown in figures are indicated by boxes.

Full-length blots for Figure 6 D



Full-length blots used for Figure 6D. Area shown in figures are indicated by boxes.

Full-length blots for Figure 7 A and E

Atg5 (sc-133158) Atg5-Atg12 50 KD Atg5 32 KD 1 2 3 4 5 6 7 8 9 10 11 12 Lane # A89RU A79RU A62NN A89RL A59NN A96LU A67LR A92RU A96LL A96RU A100RL A100RU Animal ID Nox2 KO WT saline Nox2 KO saline WT Dox GAPDH (ab8245) 36 KD GAPDH 1 2 3 4 5 6 7 8 9 10 11 12 Lane # A89RU A79RU A62NN A59NN A89RL A96LL A96LU A92RU A67LR A96RU A100RU A100RL Animal ID Nox2 KO Dox WT saline WT Dox Nox2 KO saline

Full-length blots used for Figure 7 A and E. Area shown in figures are indicated by boxes.

Full-length blots for Figure 7 B



Full-length blots used for Figure 7 B. Area shown in figures are indicated by boxes.



Full-length blots for Supplementary Figure 1 B





Full-length blots for Supplementary Figure 1 F



Full-length blots for Supplementary Figure 2 A

ERK (sc-1647)



p-ERK (sc-7383)



Full-length blots for Supplementary Figure 2 A



Full-length blots for Supplementary Figure 2 A

S6 (sc-74459)



GAPDH (ab8245)



Full-length blots for Supplementary Figure 3 A



Full-length blots for Supplementary Figure 3 A

GAPDH (ab8245)



Full-length blots for Supplementary Figure 4 A

LC3 (ab192890)



Full-length blots for Supplementary Figure 4 A

GAPDH (ab8245)



Full-length blots for Supplementary Figure 5 A

Bcl-2 (sc-7382)					
E	Bcl-2		·		26 KD
Lane #		123	456	789	
Animal ID		A79RU A62NN A59NN	A96LL A96LU A67LR	A100RL A100RU A96RU	
		WT saline	WT Dox	Nox2 KO Dox	
Bax (sc-7480)					
		144	DES.	-	
	Bax				23 KD
Lane #		123	456	789	
Animal ID		A62NN A59NN	A96LL A96LU A67LR	A100RU A96RU	A1000
		WT saline	WT Dox	Nox2 KO Dox	

Full-length blots for Supplementary Figure 5 A



Full-length blots for Supplementary Figure 5 A

