Online Supplementary Data for

Inadequate Ubiquitination-Proteasome Coupling Contributes to Myocardial Ischemia-Reperfusion Injury

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Supplementary Tables 1 and 2 as well as Supplementary Figures 1 through 3.

Genotype	Ubqln1 ^{t/t} , Ubqln1 ^{t/+} , or Myh6-Cre (n=9; 5m+4f)	Ubqln1 ^{f/+} ::Myh6-Cre (n=8; 4m+4f)	Ubqln1 ^{f/f} ::Myh6-Cre (n=8; 4m+4f)
HR (BPM)	474 ± 39	445 ± 65	$460 \pm 48^{\rm ns}$
EF (%)	52.14 ± 5.09	50.90 ± 6.52	52.76 ± 8.54 ^{ns}
FS (%)	26.81 ± 3.19	25.87 ± 4.19	27.08 ± 5.51 ^{ns}
SV (µL)	41.19 ± 6.85	42.36 ± 6.00	41.62 ± 7.86 ^{ns}
LVID;d (mm)	4.18 ± 0.19	4.26 ± 0.13	$4.14 \pm 0.16^{\text{ns}}$
LVID;s (mm)	3.17 ± 0.21	3.34 ± 0.16	3.17 ± 0.28 ^{ns}
LVPW;d (mm)	0.62 ± 0.03	0.62 ± 0.05	0.60 ± 0.00 ^{ns}
LVPW;s (mm)	0.77 ± 0.05	0.75 ± 0.06	0.77 ± 0.06 ^{ns}

Supplementary Table 1. Echocardiographic characteristics of Ubqln1 CKO mice.

Echocardiography was performed on littermate mice at 10 weeks of age. Mice with Ubqln1^{f/f}, Ubqln1^{f/+}, or Myh6-Cre Tg genotype show no significant echocardiographic difference from sexand age-matched WT mice at this age; hence, they are pooled here. Values are mean \pm SD, HR = heart rate; EF = ejection fraction; FS = fractional shortening, SV = stroke volume; LVAW = left ventricular anterior wall thickness at end diastole (d) or end systole (s); LVPW = left ventricular posterior wall thickness; LVID = left ventricular internal dimension; ns p>0.05 among genotypes, 1 way ANOVA.

	Proteins Catalog Number Vendor				
Catalog Number	Vendor				
A7811	Sigma				
8146	Cell Signaling				
10R-G109a	Fitzgerald				
sc-9996	Santa Cruz				
M115-3	MBL				
(N/A)	(custom made)				
BML-PW8305-0025	Enzo Life Sciences				
Sc-55529	Santa Cruz				
ab3341	Abcam				
sc-8017	Santa Cruz				
05-1307	Millipore Sigma				
05-1308	Millipore Sigma				
21052-1-AP	Proteintech				
	8146 10R-G109a sc-9996 M115-3 (N/A) BML-PW8305-0025 Sc-55529 ab3341 sc-8017 05-1307 05-1308				

Supplementary Table 2. Antibodies Information





Supplementary Figure 2. The stain-free total protein image of the SDS-PAGE for the western blot analyses for myocardial Ubqln1 in I/R and sham control WT mice as shown in main text Figure 1C.

