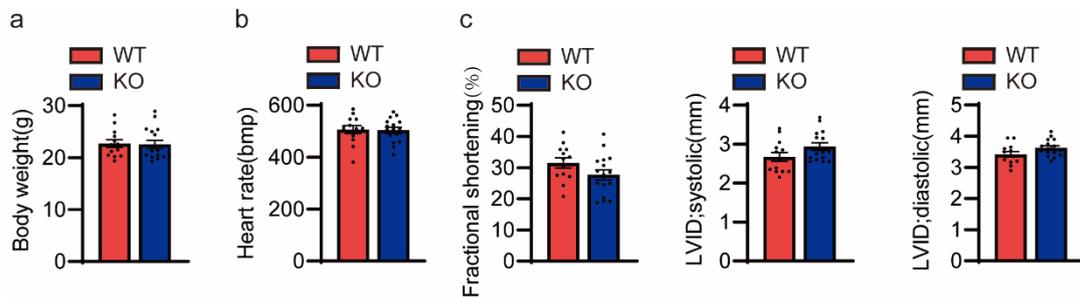
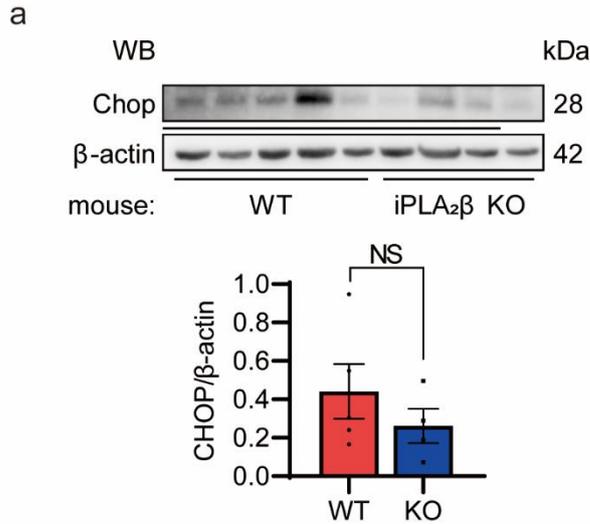


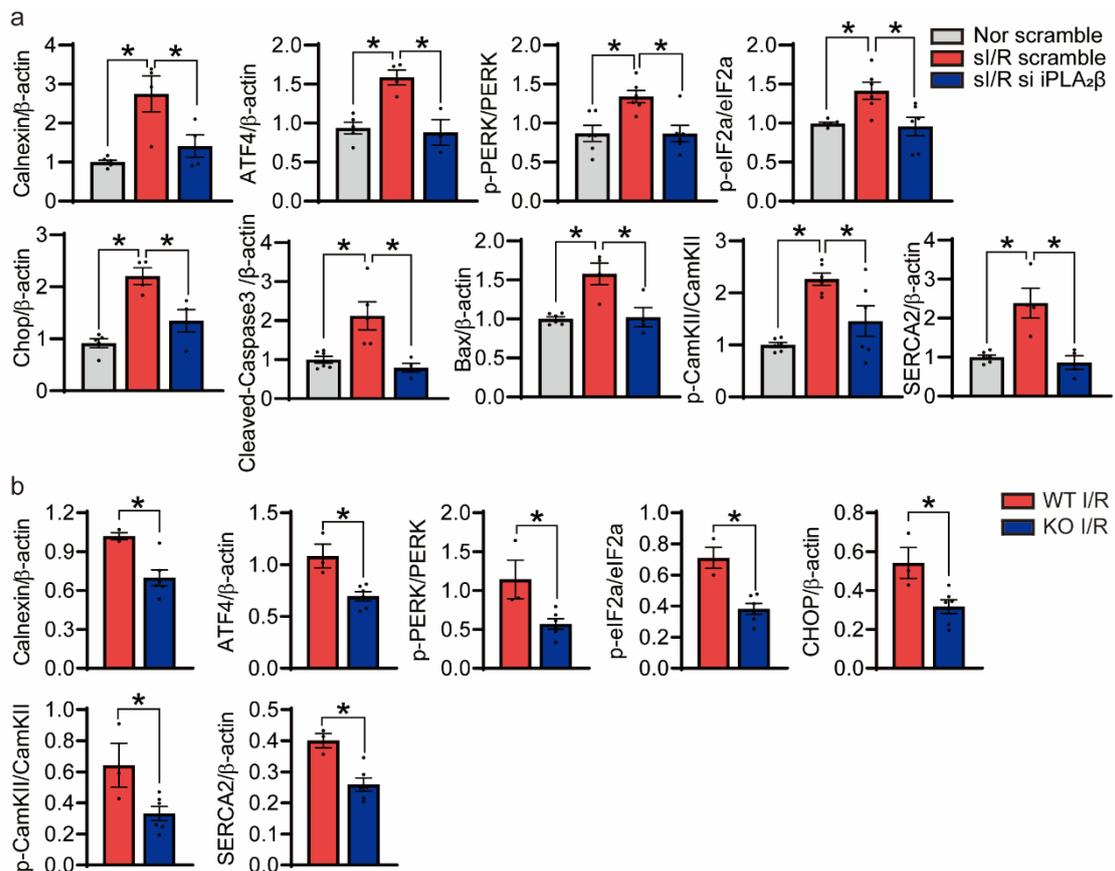
Supplementary Fig S1. Simulated ischemia/reperfusion (sI/R) injury-triggered ER stress in NRVMs. (a) sI/R led to strong induction of the ER resident chaperone GRP78 at mRNA level. NRVMs were subjected to ischemia for 3 h, followed by different times of reperfusion. Quantification real-time PCR was conducted to measure mRNA levels with 18S as an internal control. $n = 3-11$ per group. (b) Ischemia for 3 h followed by reperfusion for 3 h induced the expression of the ER chaperone GRP78 at protein level. Immunoblotting showed that GRP78 was significantly increased at protein level. β -actin was used as a loading control. Quantification is shown below. $n = 5$ per group. (c) Effects of sI/R on XBP1 splicing. NRVMs were subjected to 3 h ischemia and 3 h reperfusion or 0.5 μ M thapsigargin (TG), an ER stress activator, for 6 h as a positive control. XBP1 splicing: RT-PCR was used to assess the relative levels of unspliced XBP1 (290 bp) and spliced XBP1 (264 bp), as described in the Materials and Methods. Data are represented as mean \pm SEM. All experiments were independently replicated in triplicate. * $p < 0.05$.



Supplementary Fig S2. Analysis of *Pla2g6*^{-/-} mice and wild-type (WT) mice. (a) iPLA₂ β knockout did not alter the mouse body weight. $n = 13-16$ per group. (b) No changes in heart rate were observed between the *Pla2g6*^{-/-} and wild-type mice. $n = 13-16$ per group. (c) iPLA₂ β knockout did not affect baseline ventricular contractile function, as evidenced by the lack of significant differences in ventricular fractional shortening, left ventricular internal diameter (LVID) systole, and LVID diastole. $n = 13-16$ per group. Data are represented as mean \pm SEM. All experiments were independently replicated in triplicate. * $p < 0.05$.

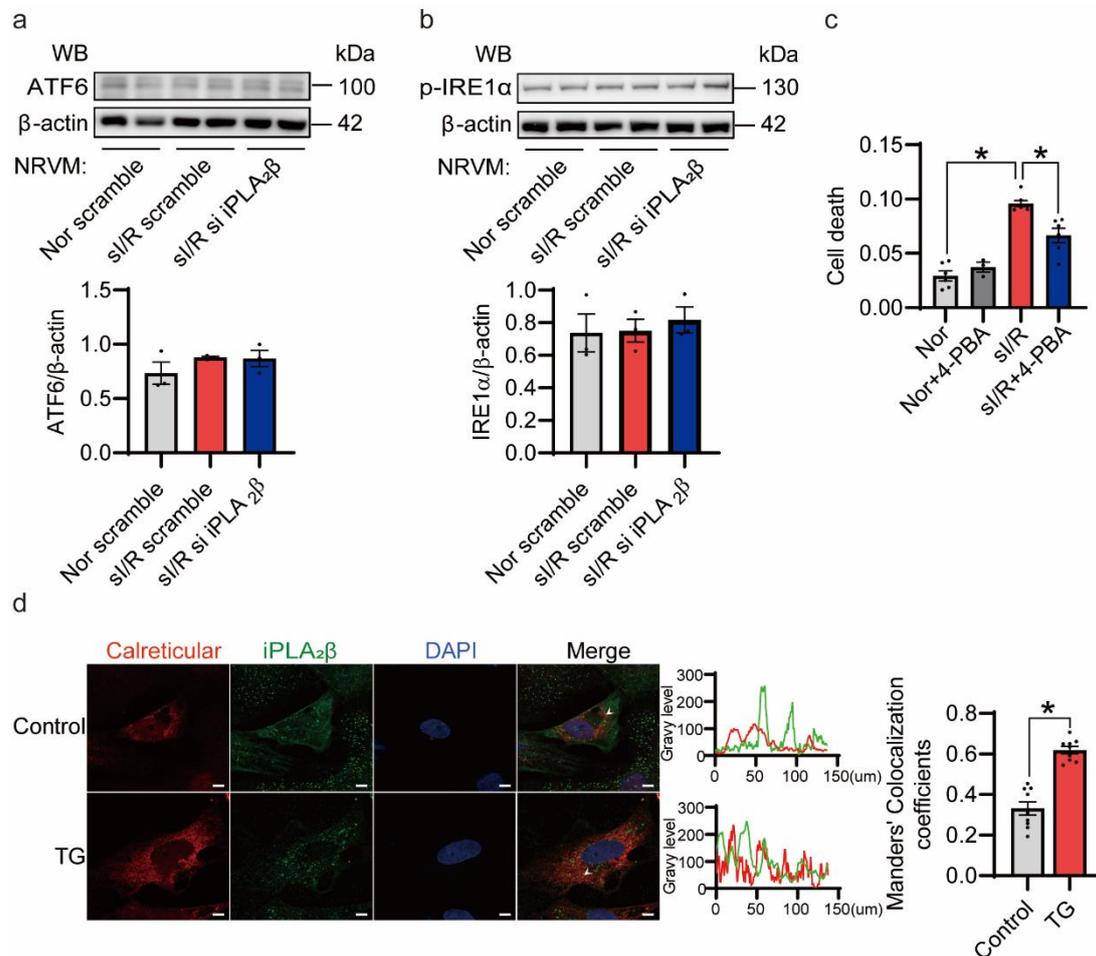


Supplementary Fig S3. No significant difference in the expression of CHOP at basal level between WT and KO mice (a) iPLA₂ β knockout did not affect the expression of basal level of CHOP protein. $n = 4-5$ per group. Data are represented as mean \pm SEM. All experiments were independently replicated in triplicate. * $p < 0.05$.



Supplementary Fig S4. The inhibition of iPLA₂ β *in vitro* and *in vivo* decreased ER stress as well as subsequent apoptosis. (a) Upregulated ER stress, apoptotic markers and Ca²⁺ regulate related proteins caused by si/R decreases with the inhibition of iPLA₂ β in NRVMs. Quantification is shown. $n = 3-6$ per group. (b) Upregulated ER stress, apoptotic markers and Ca²⁺ regulate related proteins

caused by I/R decreases in *Pla2g6*^{-/-} mice hearts. Quantification is shown. *n* = 3–6 per group. Data are represented as mean ± SEM. All experiments were independently replicated in triplicate. **p* < 0.05.



Supplement Fig S5. si/R led to NRVM apoptosis induced by ER stress but not through the ATF6 and *p*-IRE1α pathways. **(a)** There were no significant differences in the ATF6 protein expression in NRVMs subjected to si/R. Quantification is shown below. *n* = 3 per group. **(b)** There were no significant differences in the *p*-IRE1α protein expression in NRVMs subjected to si/R. Quantification is shown below. *n* = 3 per group. **(c)** Pretreatment with 5 mM 4-phenylbutyric acid (4-PBA) for 30 min ameliorated myocardial cell death caused by si/R injury. LDH levels were measured to determine relative cell death. *n* = 7 per group. **(d)** ER stress caused iPLA₂β translocation to ER. NRVMs were treated with 0.5 μM thapsigargin (TG) for 6 h. iPLA₂β was revealed by confocal immunofluorescence staining (green). ER was revealed by confocal immunofluorescence staining for calreticulin (red). Scale bar = 20 μm. The areas of interest are labeled by white arrows. The signaling intensity for both channels was scanned and recorded on the right. Colocalization of iPLA₂β and ER was quantified by Mander's colocalization coefficients (MCCs). *n* = 7–8 per group. Data are represented as mean ± SEM. All experiments were independently replicated in triplicate. * *p* < 0.05.