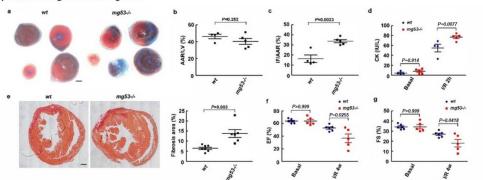
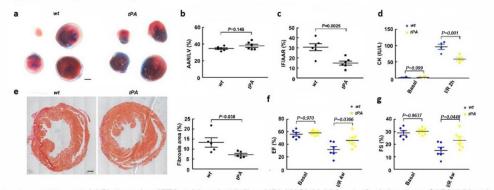
## Supplemental Figure s1. Wang et al

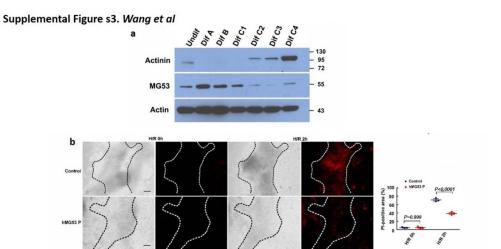


Supplemental Figure s1. (a) Photographs of TTC staining of wt (left) and mg53-/- (right) hearts. Scale bar: 1mm. Area at risk (AAR)/ left ventricle (LV) ratio (b) and infarct size (IF)/ AAR ratio (c) of hearts derived from wt and mg53-/- following I/R injury mice. n=4 for wt and 5 for mg53-/-. (d) Plasma CK level from wt and mg53-/- mice was measured at indicated time points. n=4 for wt and 5 for mg53-/-. (e) Photographs of Picrosirius red staining of wt (left) and mg53-/- (middle) hearts on 4 weeks after I/R injury. Fibrosis area was quantified (right) by ImageJ. n=7 for wt and 6 for mg53-/-; scale bar; 500 µm. Ejection fraction (EF) (f) and fractional shortening (FS) (g) of wt and mg53-/- hearts were measured by echocardiogram. n=7 for wt and 6 for mg53-/-.

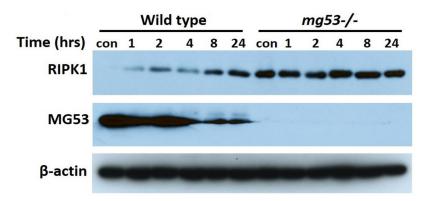
## Supplemental Figure s2. Wang et al



Supplemental Figure s2. (a) Photographs of TTC staining of wt (left) and tPA-MG53 (right) hearts. Scale bar: 1mm. AAR/LV (b) and IF/AAR (c) of wt and tPA-MG53 were quantified. (d) CK release in wt and tPA-MG53 was measured at basal and 2 hr after reperfusion, n=6 mice per group. Four week after I/R injury, cardiac fibrosis in wt and tPA-MG53 hearts was determined by Picrosirius red staining (scale bar: 500 µm) and quantified by ImageJ (n=5 per group) (e). Echocardiogram was used to determine EF (f) and FS (g) of wt and tPA-MG53 hearts at basal and 4 weeks after I/R injury (n=6 for wt and 7 for tPA-MG53).



Supplemental figure s3. (a) Dynamic expression of MG53 was determined by Western blot. Undif: human iPSC, Dif A: iPSC in differentiation medium A; Dif B: iPSC in differentiation medium B; Dif C1 (C2, C3, and C4): iPSC in differentiation medium C for 1day (2 days, 3 days and 4 days), The beating iPSC derived cardiomyocytes are usually observed on day 3 in differentiation medium C. (b) PI staining and quantification of human iPS derived cardiomyocyte following H/R stress with or without rhMG53 treatment. Scale bar,  $100 \ \mu m$ . Data are mean  $\pm$  SEM.



**Figure s4** Dynamic expression of RIPK1 and MG53 was determined by Western blot at indicated time points in wt C2C12 cells (wild type) and *mg53-/*-C2C12 cells (*mg53-/*-) following 12 hr hypoxia and reperfusion (1, 2, 4, 8 and 24 hrs). While expression of MG53 declined and RIPK1 expression increased following reperfusion in C2C12 cells, expression of RIPK1 remained high throughout all reperfusion time points in *mg53-/*-C2C12 cells.