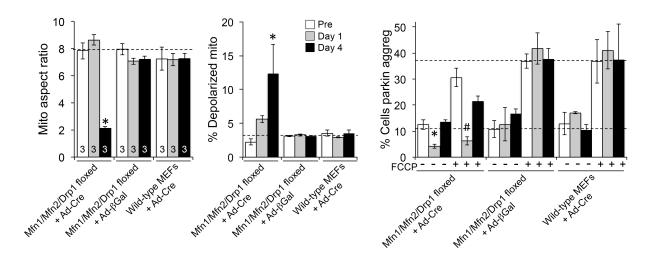
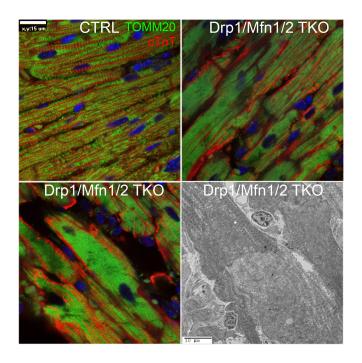


**Figure S1.** Characterization of two lines of cardiac Drp1 transgenic mice, Related to Figure 1. On the left are immunoblots and mitochondrial respiration studies from 10-fold normal expressing TG Drp1-10. On the right are immunoblot data from 25-fold normal expressing TG Drp1-25, which is the same line described in Figure 1. TG Drp1-25 results here are different mice than in Figure 1. TOM20 is translocase of outer mitochondrial membrane 20; multiple bands labeled CI-CV are respiratory complex subunits as annotated. No significant difference (*t*-test).



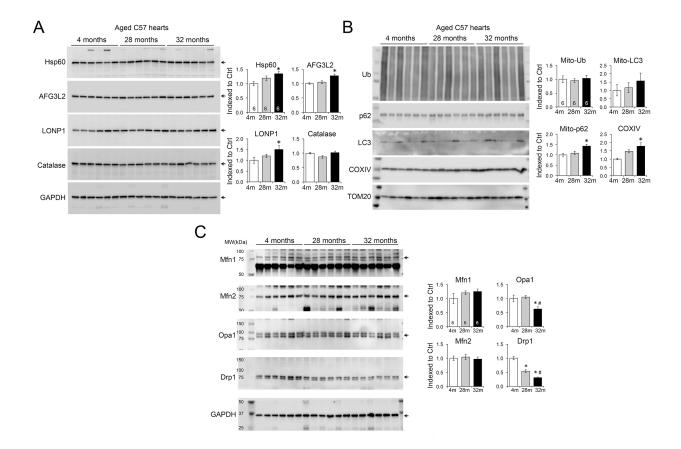
**Figure S2.** Studies to control for non-specific adenoviral effects on mitochondria, Related to Figure 2.

Mfn1/Mfn2/Drp1 triple floxed MEFs or control wild-type MEFs were infected with adenoviri encoding Cre-recombinase (Ad-Cre) or control βGal at 100 MOI. Mitochondrial aspect ratio (left), polarization status (middle) and parkin aggregation (right) were measured before (Pre; white), 1 (grey) and 4 days (black) after Ad-virus application. Mfn1/Mfn2/Drp1 triple floxed MEFs + Ad-Cre data are re-plotted from Figure 2 for comparison. N=3 each treatment condition. Control conditions did not alter any read-out.



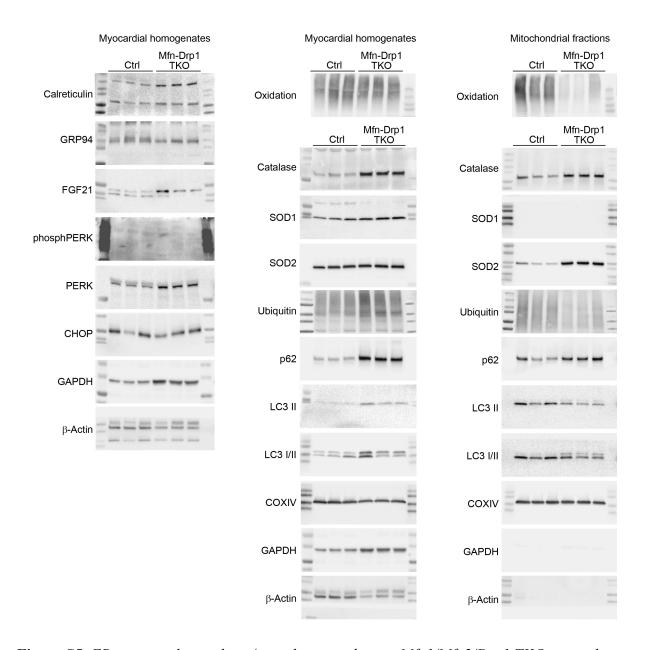
**Figure S3.** Sarcomere and mitochondrial distribution in Mfn1/Mfn2/Drp1 TKO mouse hearts, Related to Figure 5.

Fluorescence microscopy of cardiac Troponin T (red) and TOMM20 (green) stained Ctrl (left upper) or Mfn1/Mfn2/Drp1 TKO (right upper and left bottom) mouse hearts. Cardiac troponin T is an essential component of sarcomere; TOMM20 is a mitochondrial outer membrane protein. Right bottom, representative transmission electron micrograph of Mfn1/Mfn2/Drp1 TKO mouse hearts.



**Figure S4.** Proteostasis, mitophagy and abundance of mitochondrial dynamics proteins in genetically normal aged mouse hearts, Related to Figure 7.

- (A) Induction of the mitochondrial unfolded protein response. Samples are myocardial homogenates; GAPDH is loading control. \* is p<0.05 vs 4 month hearts (ANOVA).
- (B) Mitophagy proteins in cardiac mitochondrial fractions. TOM20 is loading control. \* is p<0.05 vs 4 month hearts (ANOVA).
- (C) Mitochondrial dynamics proteins in myocardial homogenates. GAPDH is loading control. \* is p<0.05 vs 4 month hearts; # is p<0.05 vs 28 month hearts (ANOVA).



**Figure S5.** ER stress and autophagy/mitophagy markers in Mfn1/Mfn2/Drp1 TKO mouse hearts, Related to Figure 7.

(*Left*) Protein levels of ER stress markers in Mfn1/Mfn2/Drp1 TKO mouse hearts 16 weeks after gene ablation. (*Middle and right*) Autophagy and ROS markers in myocardial homogenates (middle) and mitochondria-enriched fractions (right) of Mfn1/Mfn2/Drp1 TKO mouse hearts 16 weeks after gene ablation. These independent studies reproduce and extend findings in the time course results presented in Figure 7.

 Table S1. qPCR primer sequences, Related to Figures 3 and 6.

Oligonucleotides	SOURCE	IDENTIFIER
mtDNA forward primer: CCTATCACCCTTGCCATCAT	Chen et al., 2010	N/A
mtDNA Reverse primer: GAGGCTGTTGCTTGTGAC	Chen et al., 2010	N/A
Nuclear DNA forward primer: ATGGAAAGCCTGCCATCATG	Chen et al., 2010	N/A
Nuclear DNA Reverse primer: TCCTTGTTGTTCAGCATCAC	Chen et al., 2010	N/A
Tfam (Mm00447485_m1)	Thermo Fisher Scientific	Cat#4331182
PGC-1α (Mm01208835_m1)	Thermo Fisher Scientific	Cat#4331182
PGC-1β (Mm00504720_m1)	Thermo Fisher Scientific	Cat#4331182
PPARγ (Mm01184322_m1)	Thermo Fisher Scientific	Cat#4331182
ANP (Mm01255747_g1)	Thermo Fisher Scientific	Cat#4331182
BNP (Mm01255770_g1)	Thermo Fisher Scientific	Cat#4331182
SERCA (Mm01201431_m1)	Thermo Fisher Scientific	Cat#4331182
GAPDH (Mm03302249_g1)	Thermo Fisher Scientific	Cat#4331182