SUPPLEMENTARY FIGURES







Supplementary Figure 2. Aged mice show increased histone 3 acetylation and increased NF- κ B target gene activation: (A) Relative acetylated histone H3K9 levels and (B) relative II-6 and (C) p21 mRNA levels in the heart of young and old mice. Values are mean ± SE, n = 5-6, * P < 0.05.



Supplementary Figure 3. Quantification of the western blot in Figure 1F: (A-E). Quantification of relative Sirt1, Sirt3, OPA1, MFN1 and DRP1 levels in the heart of young and old mice. Values are mean ± SE, n = 6 * P < 0.05, ** P < 0.01 (NS=not significant).





WT

Sirt6.KO



Supplementary Figure 4. Quantification of the western blot in Figure 3F: (A-E). Quantification of relative MFN1, OPA1, acetylated histone H3K9, Sirt1 and Sirt3 levels in the heart of Wild type and Sirt6.KO mice. Values are mean ± SE, n = 6, * P < 0.01 (NS=not significant).

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0



Supplementary Figure 5. Quantification of the western blot in Figure 5C: (A–G). Quantification of relative OPA1, MFN1, OGG, Sirt1, Sirt3, p53, and Sirt6 levels in cardiomyocytes treated as indicated. Values are average of three independent experiments, mean \pm SE, * P < 0.01 (NS=not significant).



Supplementary Figure 6. Quantification of the western blot in Figure 6E: (A–F). Quantification of relative MFN1, DRP1, Sirt1, Sirt3, Sirt6, and acetylated histone H3K9 levels in the heart of 24-month-old Wild type and Sirt6.Tg mice. Values are mean \pm SE, n = 6, * P < 0.05, ** P < 0.01 (NS=not significant).