

Supplementary information

The sirtuin SIRT6 blocks IGF-Akt signaling and development of cardiac hypertrophy by targeting c-Jun.

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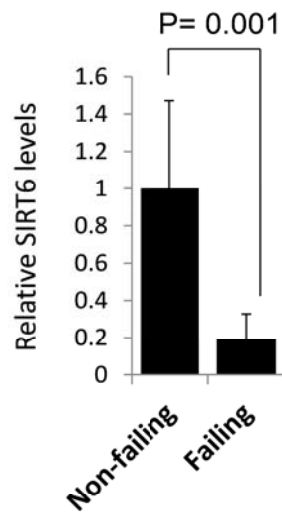
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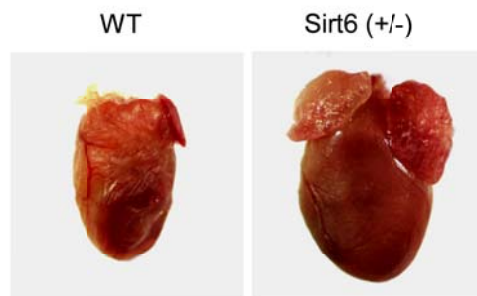
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Supplementary Figure 1

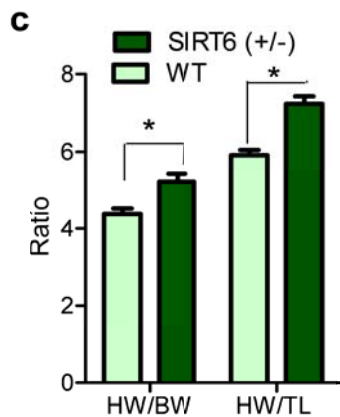
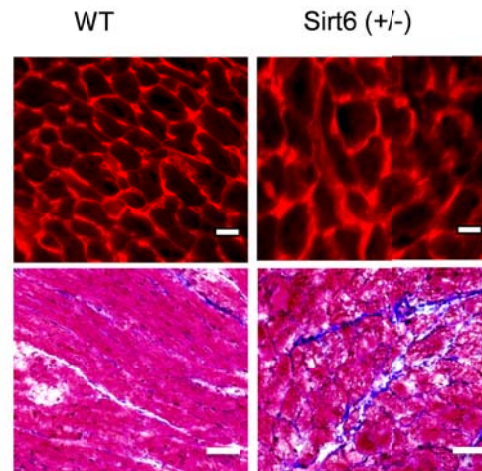
a : human hearts



b : mouse hearts



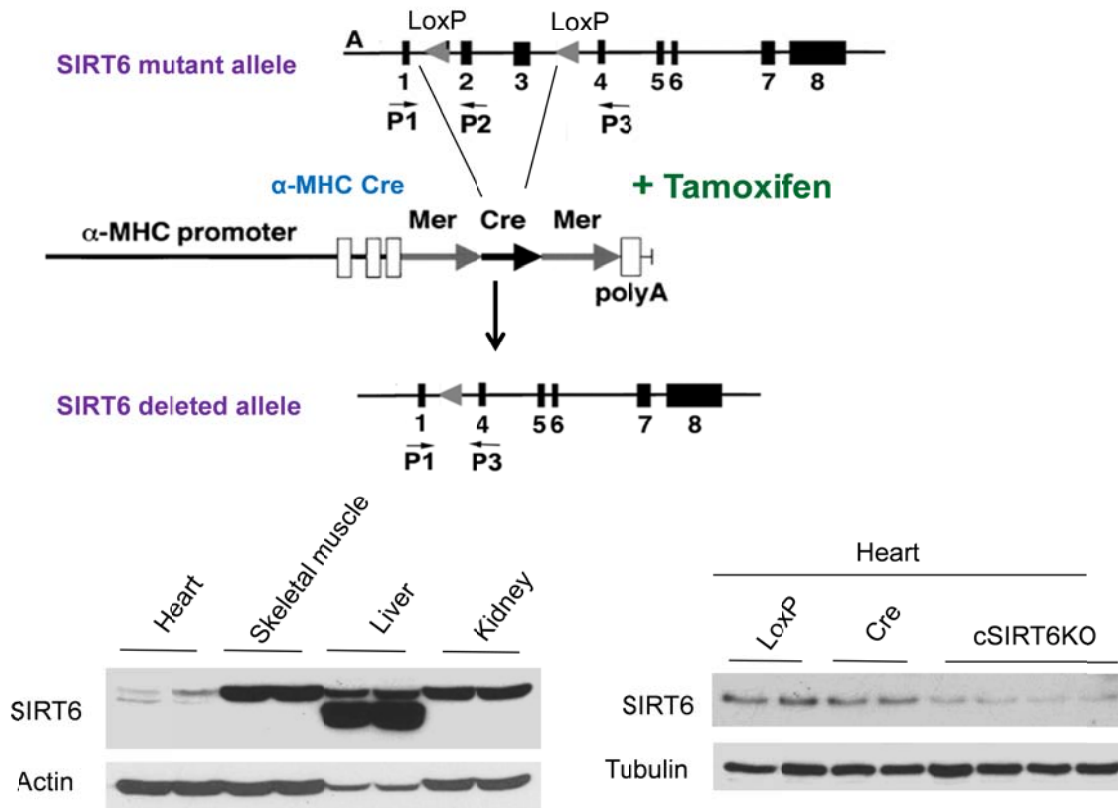
d



Supplementary figure 1: Reduced SIRT6 levels lead to induction of cardiac hypertrophy.

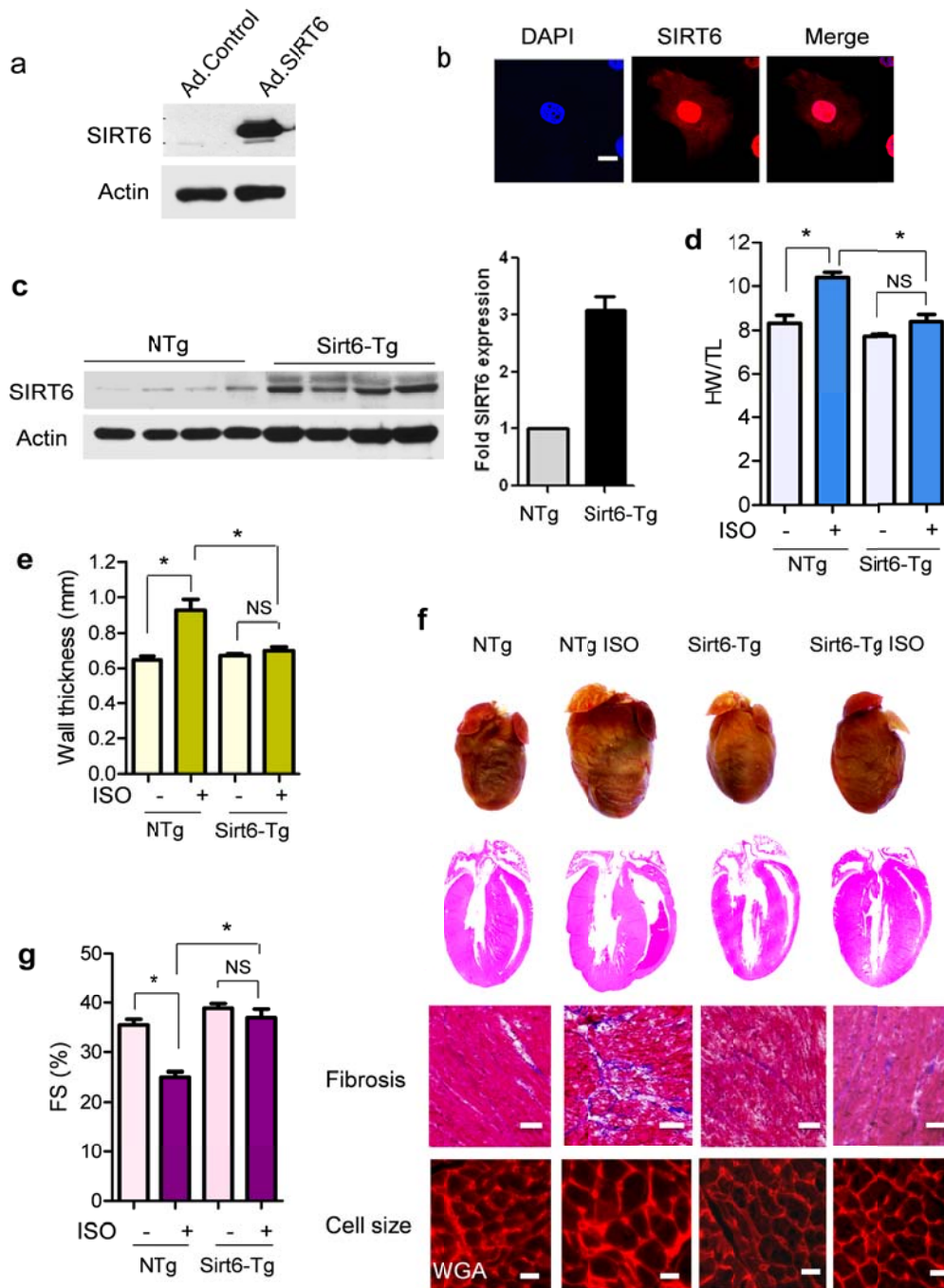
(a) Quantification of SIRT6 levels in non-failing and failing human hearts (for details see supplementary table1). **(b)** Hypertrophy of SIRT6 (+/-) hearts, compared to WT control at 6 months of age. **(c)** HW/BW and HW/TL ratios in WT and SIRT6 (+/-) mice. **(d)** Heart sections showing increased cardiomyocyte size (scale bar 10 μ M) and fibrosis (bar 40 μ M) in SIRT6 (+/-) hearts compared to WT controls.

Supplementary Figure 2



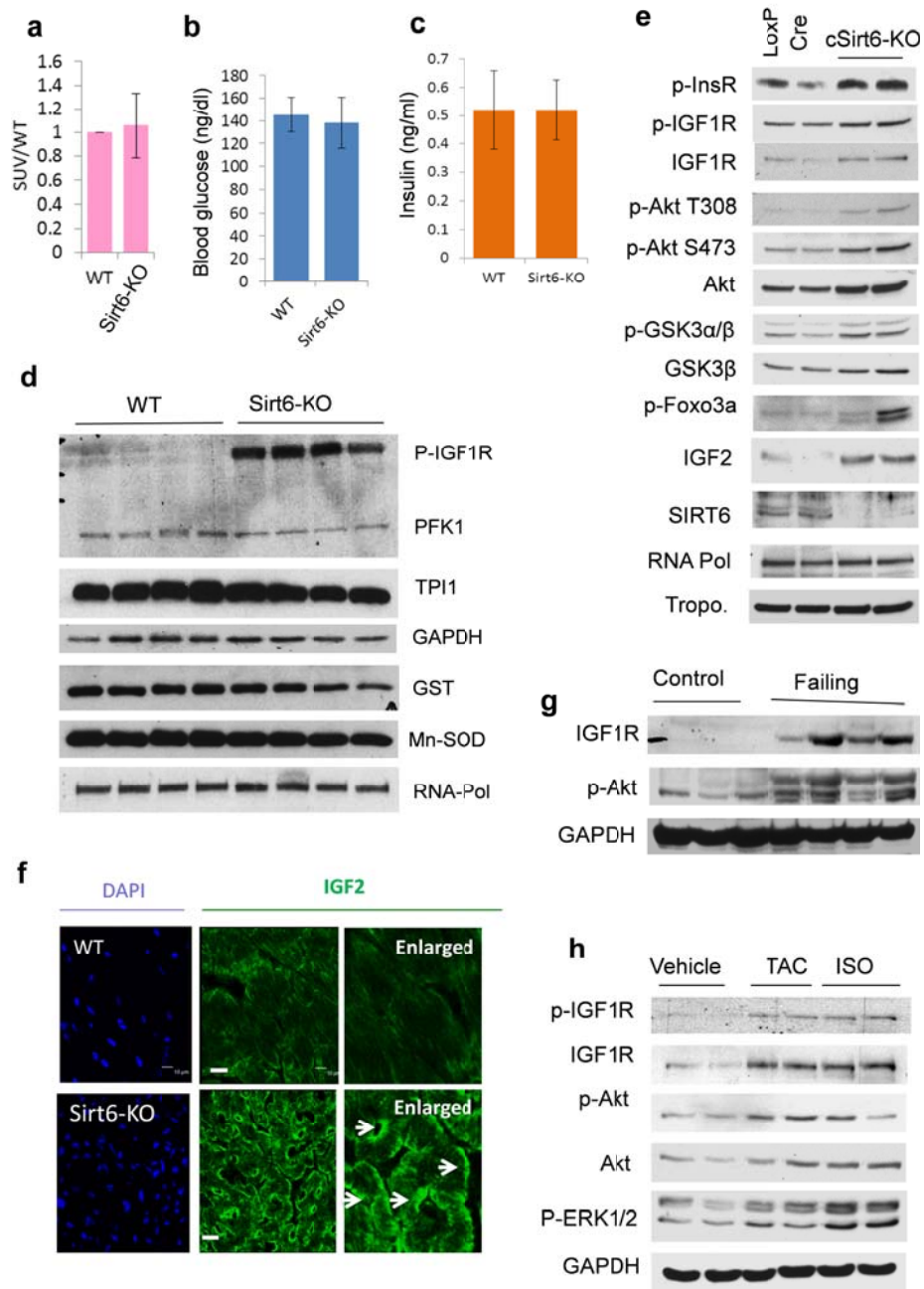
Supplementary figure 2: Generation of inducible cardiac specific SIRT6 knockout mice. Mice carrying both *loxp/loxp* and *α-MHC-Cre* alleles were injected with tamoxifen at 4 weeks of age for three consecutive days. SIRT6 deletion was determined by western blotting of different tissue lysates.

Supplementary Figure 3



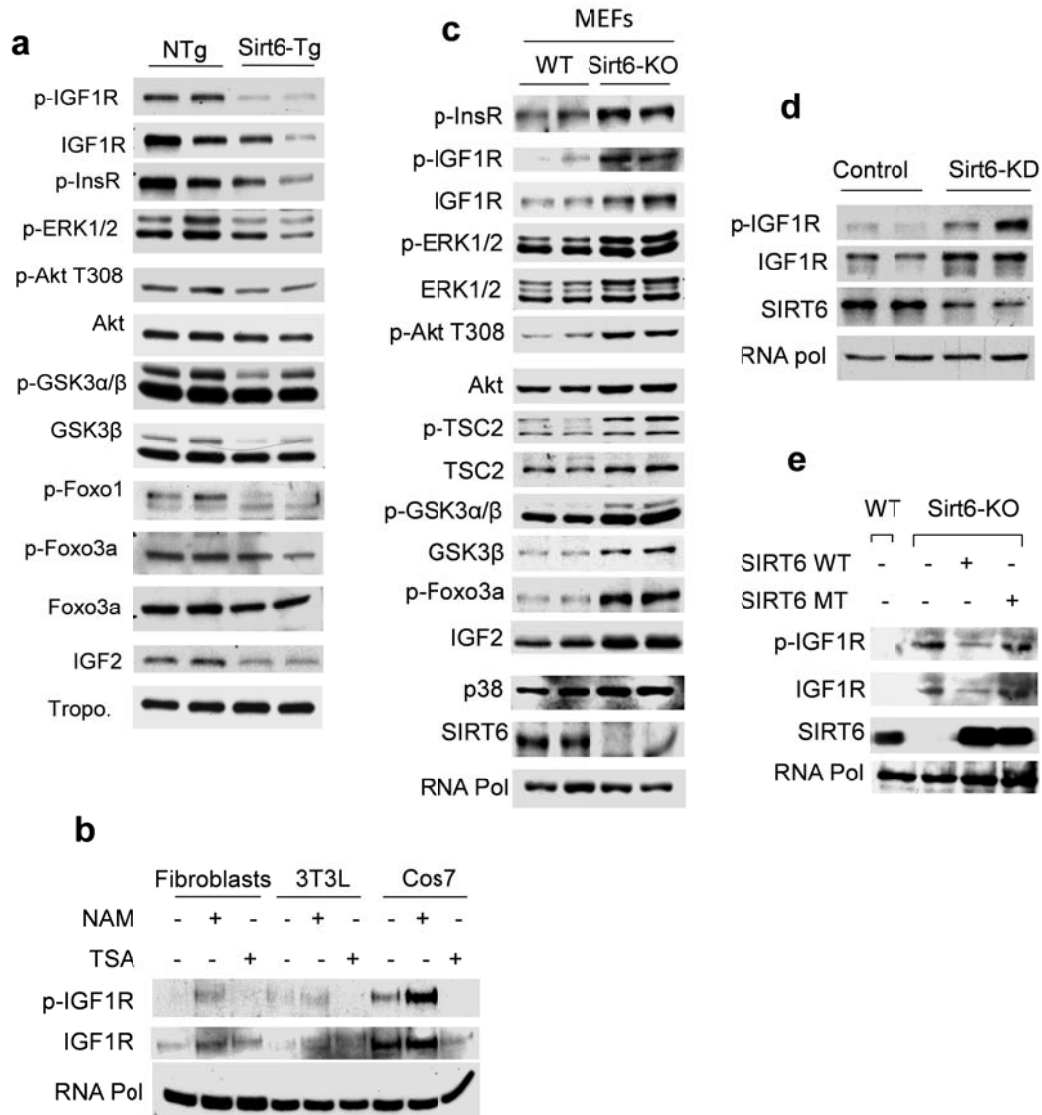
Supplementary figure 3: SIRT6 over expression blocks agonist-mediated cardiac hypertrophy. (a) Western blots showing expression of SIRT6 adenovirus in cardiomyocytes. (b) Nuclear localization of SIRT6 in cardiomyocytes (scale bar 10 μ M). (c) Expression levels of SIRT6 in non-transgenic (NTg) control and SIRT6.Tg hearts. Mean \pm SD, n=4. (d) N.Tg and SIRT6-Tg mice were treated with vehicle or isoproterenol (ISO, 8.7 mg/kg/d for 7 days), and their HW/TL ratio was determined; mean \pm SD, n =5-6. (e) Left ventricular wall thickness as determined by echocardiography. Mean \pm SD, n = 5-6. (f) Representative whole hearts and heart sections of N.Tg and SIRT6-Tg mice subjected to develop ISO-mediated hypertrophy. Scale bar to detect fibrosis 40 μ M, and for cell size 10 μ M. WGA indicates wheat germ agglutinin. (g) Fractional shortening of same mice as in panel e. Mean \pm SD, n = 5-6. *p<0.001.

Supplementary Figure 4



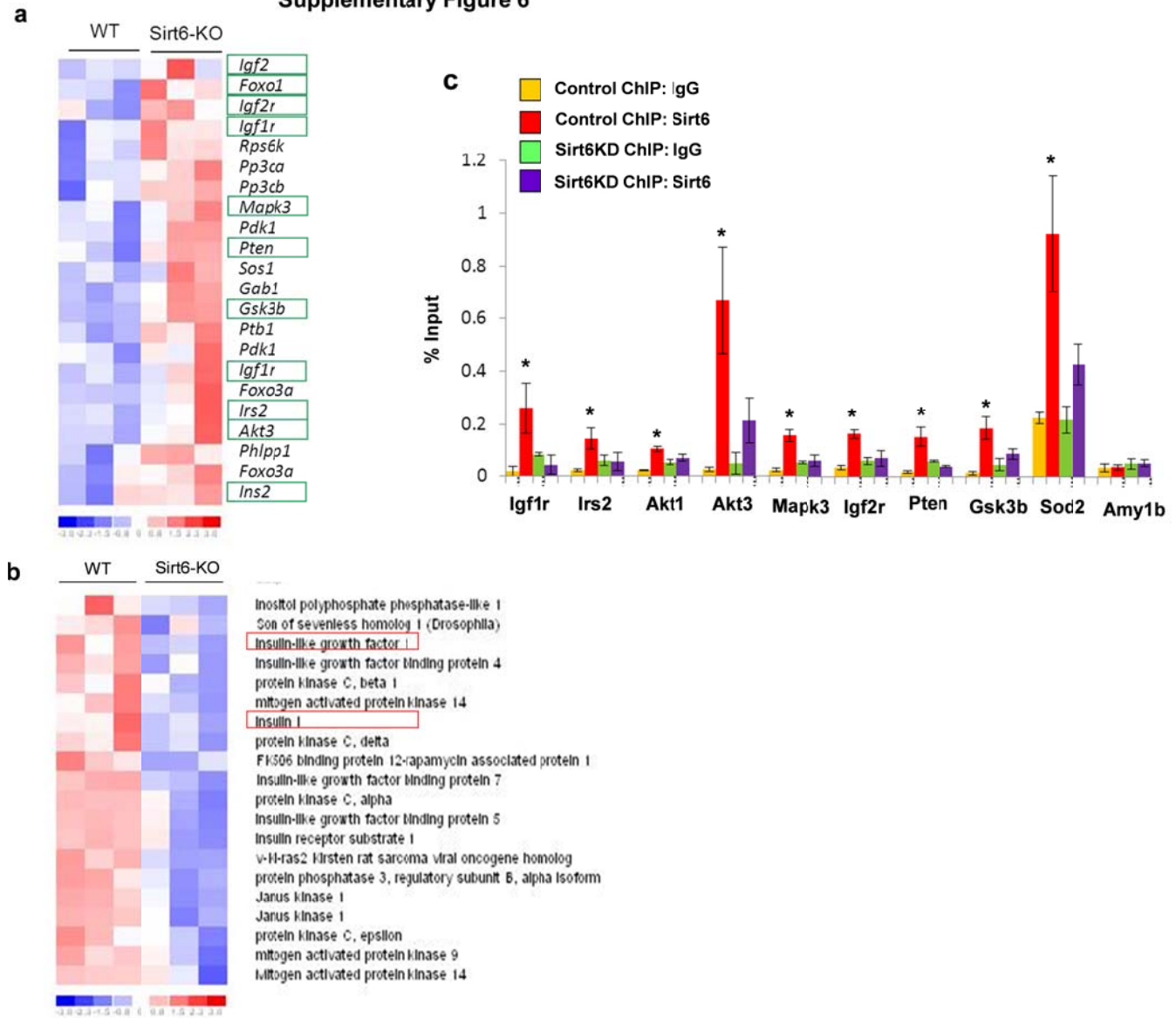
Supplementary figure 4: Biochemical characteristics of SIRT6-deficient hearts. (a) Standard uptake value (SUV) ratio of labeled ^{18}F FDG-Glucose incorporation in the heart of WT and SIRT6-KO mice, $n = 3$. (b) Blood glucose levels in WT and SIRT6KO mice at 3 months of age. (c) Serum insulin levels of the WT and SIRT6KO mice at 3 months of age, $n = 10-12$. (d) Western analysis of Hif-1 α target glycolytic genes and NF- κ B target antioxidant genes. Phospho-IGF1R is used as positive control. (e) Western analysis of IGF signaling related genes in controls and cardiac-specific SIRT6KO (cSIRT6KO) heart lysates. (f) Confocal images showing membrane localization of IGF2 (green) in WT and SIRT6KO hearts (scale bar 10 μ M). Arrows indicate membrane bound IGF2 in myocytes. (g) Western analysis of control and failing human heart lysates. (h) Western analysis of IGF/Akt signaling related targets in control and hypertrophied mouse heart lysates. Cardiac hypertrophy was induced by pressure overload (TAC) or isoproterenol (ISO) infusion in mice.

Supplementary Figure 5



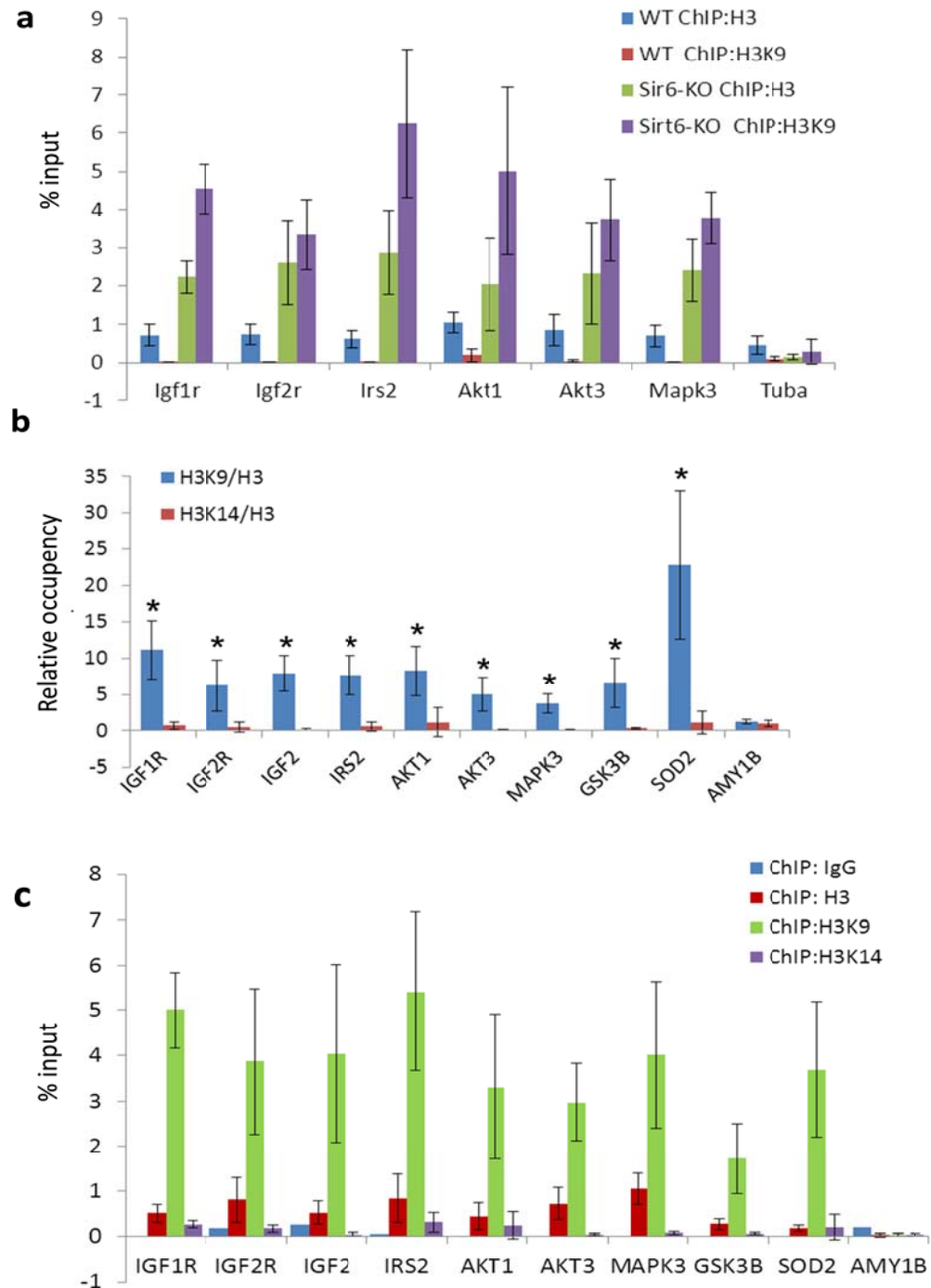
Supplementary figure 5: Western analysis of IGF signaling related genes. (a) Expression levels of IGF signaling related genes in NTg and SIRT6-Tg hearts. **(b)** Expression and phosphorylation of IGF1R in vehicle, nicotinamide (NAM) and trichostatin-A (TSA) treated cells. **(c)** Increased expression and phosphorylation of IGF signaling related genes in WT and SIRT6-KO MEFs. **(d)** IGF1R levels in control and SIRT6-KD 293T cells. **(e)** Over expression of WT, but not the mutant (MT) SIRT6 suppressed the IGF1R signaling in SIRT6-KO fibroblasts.

Supplementary Figure 6



Supplementary figure 6: SIRT6-dependent regulation of IGF signaling related genes. (a and b) Comparative microarray analysis of IGF signaling related genes in wild type and SIRT6-KO skeletal muscle samples. **(c)** The ChIP analysis data from Fig. 4e is shown without normalization as % input.

Supplementary Figure 7



Supplementary figure 7: The ChIP analysis with wild-type and SIRT6 knockout hearts and cells. (a) The ChIP analysis was performed in WT and SIRT6-KO heart samples with use of Ac-H3K9 (H3K9) and H3 antibodies. Acetylation of H3K9 at promoters is shown relative to total H3 levels and the data is shown as % input. (b and c) ChIP analysis was performed with use of Ac-H3K9, Ac-H3K14 and H3 antibodies in SIRT6-KD stable 293T cells. The data presented as either normalized with H3 levels (b) or shown without normalization as % input (c).

Supplementary Figure 8

Insulin-like growth factor I receptor (Igf1r) promoter

Mouse acggcagcggcgctgctcggctt**tgaccttcagcga**gcccggagcccccgcgcacggagtc
 Rat acggcagcggcgctgctcggctt**tgaccttcagcga**gcccggagcccccgcgcacggagtc
 Human acggcagcggcgctcgc-cctcggctt**tgaccttcagcga**gcccggagcccccgcgcacggagtc
 Opossum acaacacctc-----ttcggctt**tgaccttcagcga**aaccggatcccccctacgttcgga
 -85

Insulin-like growth factor II receptor (Igf2r) promoter

Mouse ggacg**tcacgtga**g-----caggaggcggggcggggcccgaactcagg**tcacgtga**cgcctcggggacgg
 Rat gaacg**tcacgtga**g-----aggaggcggggcaggggcccgaactcagg**tcacgtga**cgccttggggacgg
 Human gggag**tcacgtga**gcgggggcccggggggcggtgcggggcggctg**tcacgtga**cgcggttccggggg
 Orangutan gggag**tcacgtga**gcgggggcccggggggcggtgcggggcggctg**tcacgtga**cgcggttccggggg
 Opossum tgcg**tcacgtga**gcgggggaggggtgaaggcagggcagcagtagagccggaccagcggggcagagca
 -49 -8

Thymoma viral proto-oncogene 1 (Akt1) promoter

Mouse gcctccag**ggcgctaagtcagag**gctcagcag//taag-gactc**gggtgacagggga**ggacgggagca
 Rat gcctccag**ggcgctaagtcagag**gcccagtag//taag-gactc**gggtgacagggga**ggagggggca
 Chimpanzee acctccag**ggggccaagtcagag**gcccagtag//aggggactt**ggggagtgaggga**gaagaggggca
 Human acctccag**ggggccaagtcagag**gcccagtag//aggggactt**ggggagtgaggga**gaagaggggca
 Orangutan acctccag**ggggccaagtcagag**gcccagtag//aggggactt**ggggagtgaggga**gaagaggggca
 Rhesus acctccag**ggggccaagtcagag**gcccagtag//aggggactt**ggggagtgaggga**gaagaggggca
 -1000 -431

Thymoma viral proto-oncogene 3 (Akt3) promoter

Mouse ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcagca
 Rat ttttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcaaca
 Human ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcagagaat
 Orangutan ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcagagaat
 Dog ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcagagaat
 Horse ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcagagaat
 Opossum ctttcttcaataacttccttca**ggctgagtcatca**ctagagagtgggaaagggcagcagcagcaaaagag
 -115

Insulin receptor substrate 2 (Irs2) promoter

Mouse cgt**aacgcagagtcac**atggttgttttgct---ct**agttcagtcactc**gggtcgcgatgt**gttactcad**tgt
 Rat cgt**aacgcagagtcac**atggttgttttgct---ct**agttcagtcactc**gggtcgcgatgt**gttactcad**tgt
 Human cgt**aacgccagagtcac**atggttgttttgctcttct**agttcagtcactc**gggtcgcgatgt**gttactcad**tgt
 Opossum cgt**aacgcagagtcac**atggttgttttgctct--t**agttcagtcactc**aggtcgcgatgt**gttactcad**tgt
 -553 -524 -499

Phosphatase and tensin homolog (Pten) promoter

Mouse caagaggcggggcagggaaaccggagcccga**tgagggtga**ccacgcgggagacacaatagggg
 Rat caagaggcggggcagggaaaccggagcccga**tgagggtga**ccacgcgggagacacaatagggg
 Human cgaaaggtggggcgctgcaagggagccgg**tgagggtga**tacacgcctggcagacacaatagcag
 Orangutan cgaaaggtggggcgctgcaagggagccgg**tgagggtga**tacacgcctggcagacacaatagcag
 Horse caggaggtggggcggtgaaagggagcccga**tgagggtga**tacacactggagacacaatagggg
 -585

Mitogen-activated protein kinase 3 (Mapk3) promoter

Mouse ggcgggaaagccgggtgcgcgccagagaccat**gtgacgct**ccggggcc
 Rat ggcgggaaagccgggtgcgcgccagagaccat**gtgacatc**ccggggcc
 Human ggcaagaagggcgggcccgcgcggggcgggtc**gtgacagc**ccaggcc
 Chimpanzee ggcaagaagggcgggcccgcgcggggcgggtc**gtgacagc**ccaggcc
 Orangutan ggcgagacgggcccgcgcgcggggtc**gtgacggc**ccaggcc
 -135

Glycogen synthase kinase 3 beta (Gsk3b) promoter

Mouse ttctcttaggattcgaggtac-tgt**cctgaccacat**ttccccctcacctgttttcggggctgct
 Rat ttctcttaggattcgaggtat-cgg**cctgaccacat**ttccccctcacctgttttcggggctgct
 Human ttctctggaggcttcaggatatccgg**cctgacggcat**ttccccctcacctgttttcggggcccg
 Orangutan ttctctggaggcttcaggatatccgg**cctgacggcat**ttccccctcacctgttttcggggcccg
 Dog ttctctggaggcttcaggatatccgg**cccagcggcgt**ttccccctcacctgttttcggggctcg
 Horse ttctctggaggcttcaggatatccgg**cctgacggcat**ttccccctcacctgttttcggg-cccgt
 Opossum tttcaccagggcctaaggatatccgg**tatgactccgt**ttccccctcacctgttttcgggaccact
 -792

Mechanistic target of rapamycin (serine/threonine kinase) (Mtor) promoter

Mouse	acttgtagtagcgacg----taccggaagtgttcggtg---gaagtgaccga- agtcaactcac
Rat	actagtagtagcgacg----taccggatgtgtgtgtgtgggaagtgaccga- agtcaactcac
Human	aatcctagcagcgccg----taccggatgtgtgagtg---gaagtgactga- ggtgaactcac
Chimp	aatcctagcagcgccg----taccggatgtgtgagtg---gaagtgactga- ggtgaactcac
Orangutan	aatcctagcagcgccg----caccggatgtatgagtg---gaagtgactga- ggtcaactgac
Rhesus	aatcctagcagcgccg----caccggatgtatgagtg---gaagtgactga- ggtcaactcac
Opossum	gctcattcaacgcagcactcactggaagccgggtcg---gaagtggctga- agtcaactcgc

-221

Aldolase C, fructose-bisphosphate (ALDOC) promoter

Mouse	cccc---agggagt ca ggtgactctgcggaagtgtgccaccttatttactccagcttggaccgagcta
Rat	cccc---agggagt ca ggtgactctgcgatctgctgctgcttatttactccagcttggactgagcta
Human	ccc---cggaggagt ca ggtgactctgacatccgcagcctcatttaccagaggagccagggtgcagc
Orangutan	ccct-cggaggagt ca ggtgactctgacatccgcagcctcatttaccagaggagccagggtgcagc
Dog	ccct-cggaggagt ca ggtgactctgcgcatccgcagcctcatttaca---ggaggctgagctagagc
Horse	cccccgggggagt ca ggtgactctgcgcatccgcagcctcatttaca---ggaggctgagctgagc
Opossum	ctcc-gctggagt ca ggtgactctgcgcatccgcagcctcttttacc---ggagcccagcccgatc

-1102 AP-1 -1098 HIF-1 α

Lactate dehydrogenase A (LDHA) promoter

Mouse	agcctacacgtgggttcccc gcacgt ccgct-gggt--cc-cact tgacgtca c-gcggagcttcc
Rat	agcctacacgtgggttcccc gcacgt ccgct-gggt--tc-cact tgacgtca c-acggagcttcc
Human	gactcacacgtgggttcccc gcacgt ccgcc-ggccc--cc-ccc tgacgtca c-atagctgttcc
Orangutan	gactcacacgtgggttcccc gcacgt ccgcc-ggcccccc-ccc tgacgtca c-atagctgttcc
Rhesus	gaccacacgtgggttcccc gcacgt ccgcc-ggccc--cc-ccc tgacgtca c-atagctgttcc
Dog	gaccacacgtgggttcccc gcacgt ccgcc-cggt--gc-ccc tgacgtca c-ctcgcttcc
Horse	gaccatacgtgggttcccc gcacgt ctcc-cagtt---c-ccc tgacgtca c-cgcgcgttcc
Opossum	atcccacacgtgggttcc gcacgt ccgct-cccag--cc-ccc gaacgtca gctttttggtggt

-34 HIF-1 α -11 c-Jun

Lactate dehydrogenase B (LDHB) promoter

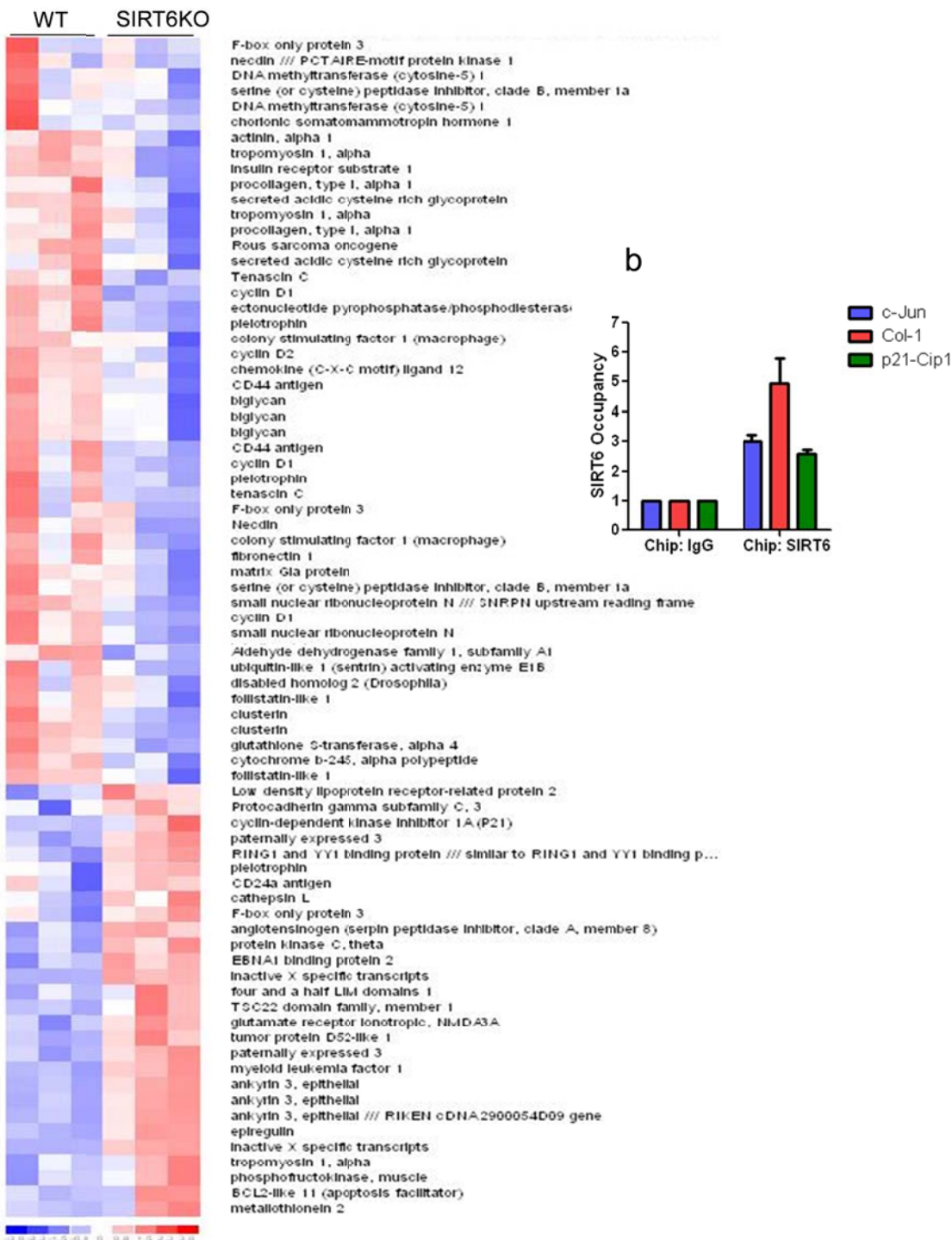
Mouse	ggaagt gac ctcttacttttagagagcggggagag-- cgcg tg ca ctccagc-cttgtccttgaagg
Rat	agaagt gac cttccacttttagagagcggggagag ca g cgcg tg ca ctccagc-cttgtccttgaagg
Human	ggaagt gagac ctcctatctggaggtgggggaggg-- agtgtg ca c acttgacccttgctccttgaagg
Orangutan	ggaagt gac ctccttttgggagatgggggaggg-- gtgtg ca c acttgacccttgtccttgaagg
Rhesus	ggaagt gac cccctatctggaggtgggggaggg-- agtgtg ca c acttgacccttgtccttgaagg
Dog	ggaagt gac ctcct-----caggggagggagag-- cgcg tg ca cttgacccttgtccttgaagg
Horse	agaagt gac ctcctatt--tggggagggcggg-- cg tg ca cttgacccttgtccttgaagg
Opossum	-----ctctccccagcataggggaagga-- cg act gca acttgacccttgtccttgaagg

-173 AP-1 -141 HIF-1

Supplementary figure 8: In-silico analysis of evolutionarily conserved c-Jun binding sites in the promoters of different genes. The transcription factor binding site analysis was done using TRANSFAC software (Biobase GmbH); evolutionary conservation analysis was carried out using UCSC Genome Browser. Nucleotide sequences in bold denote AP-1/c-Jun consensus binding sites, and number below denotes distance (bp) from the transcription start site of the respective mouse gene. Upon extensive analysis, we confirmed that InsR does not have any conserved binding site for AP1/c-Jun, though it showed increased expression in SIRT6KO hearts. Interestingly, our in silico analysis does not find any c-Jun binding sites in HIF-1 α target glycolytic genes, PFK1, TP11, PDK1 and GAPDH. However, few glycolytic genes especially ALDOC, LDHA, and LDHB have both c-Jun and HIF-1 α binding sites very close to their proximal promoter regions.

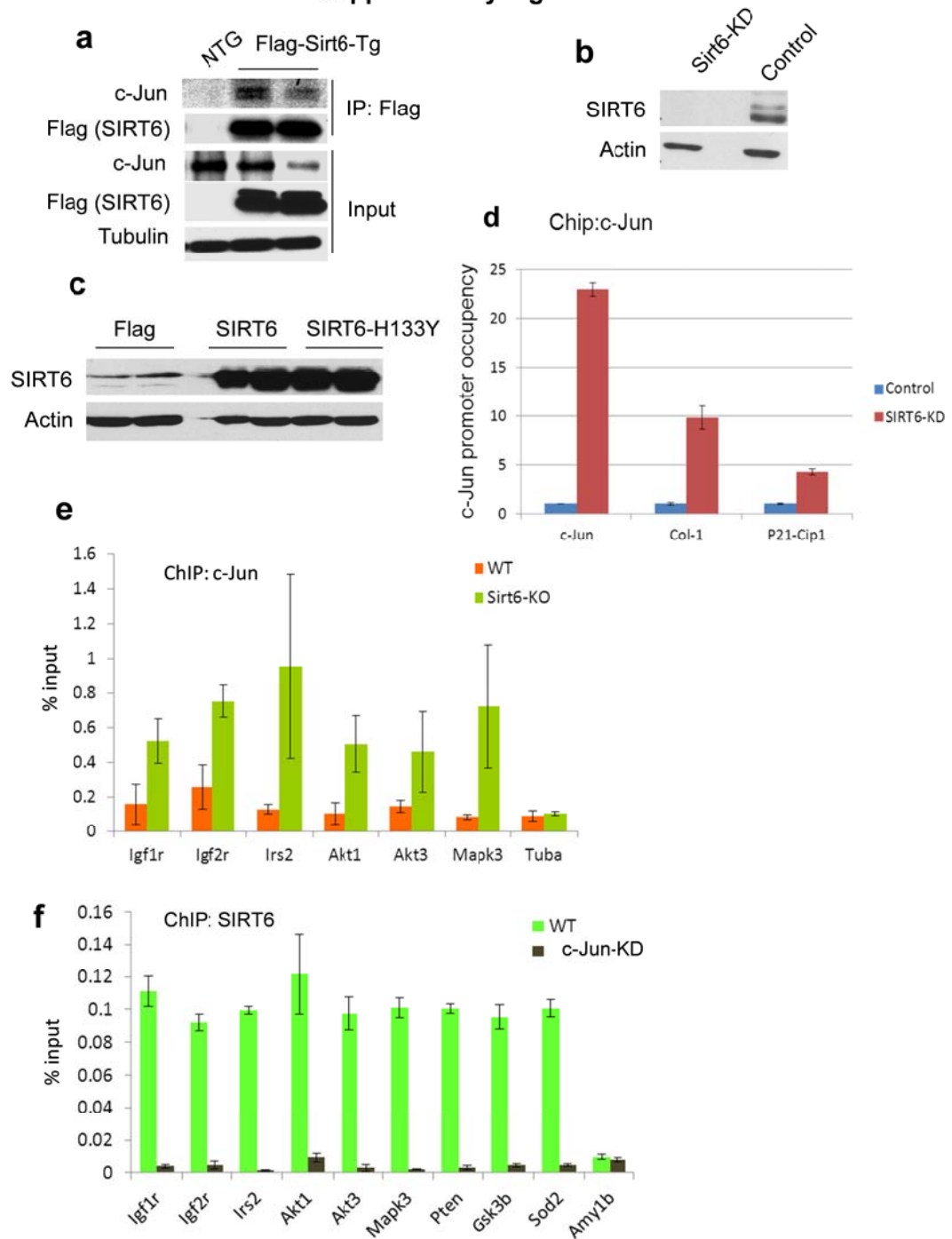
a

Supplementary Figure 9



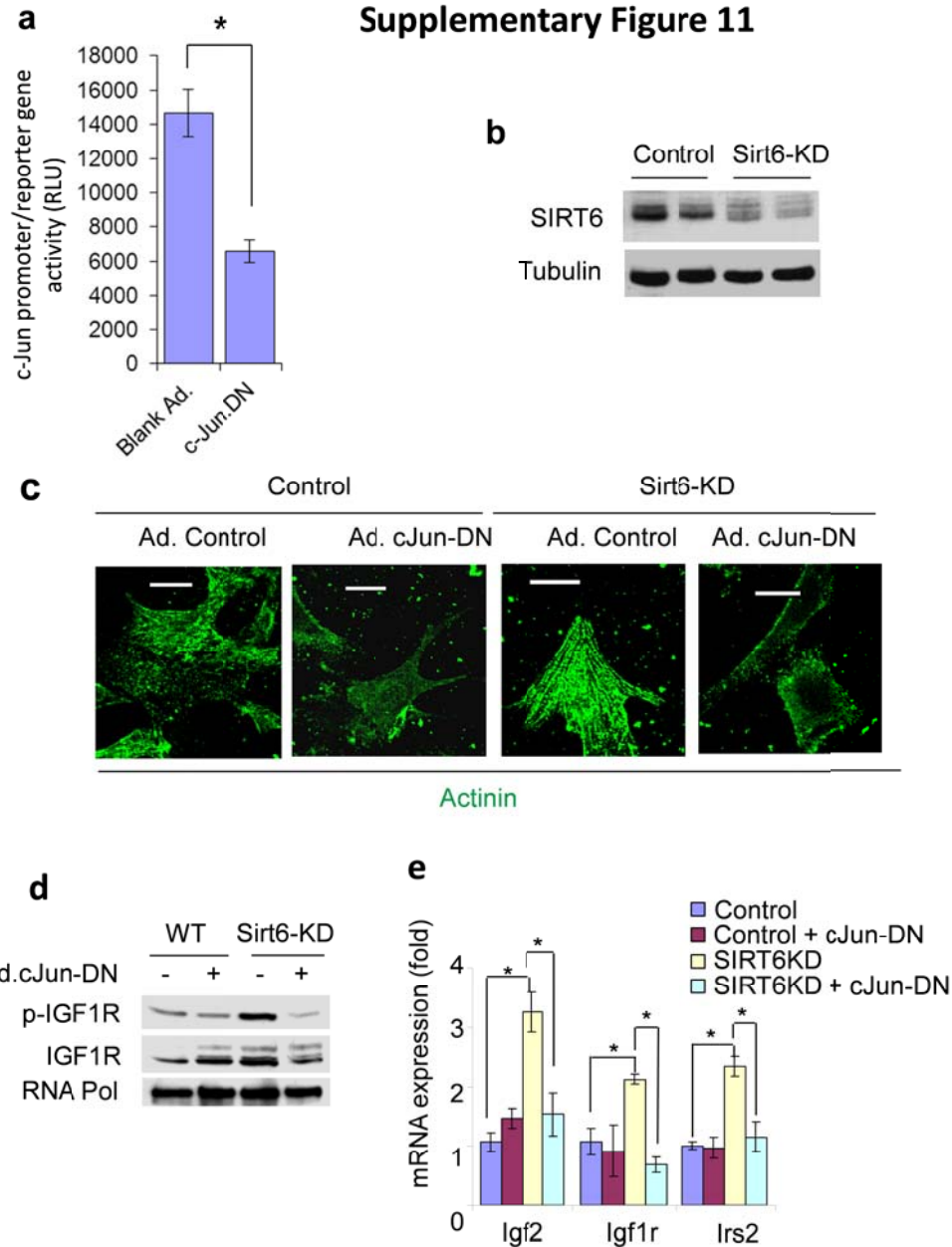
Supplementary figure 9: Analysis of AP1 target genes. (a) Comparative microarray analysis of AP-1 target genes in wild type and SIRT6-KO skeletal muscle samples. **(b)** ChIP analysis showing SIRT6 binding to AP1 target gene promoters. Mean \pm SD, n = 3.

Supplementary Figure 10



Supplementary figure 10: SIRT6 associates with c-Jun on IGF signaling related genes. (a) SIRT6 was immunoprecipitated from the SIRT6 transgenic mouse hearts using Flag-tag specific antibody. The interaction of SIRT6 with c-Jun was determined by western blotting. **(b and c)** SIRT6 levels in SIRT6KD (b) and overexpressing (c) cells determined by western blotting. **(d)** The ChIP analysis of c-Jun occupancy in the promoters of IGF signaling related genes of control and SIRT6-KD stable 293T cells. **(e)** The ChIP analysis data from Fig. 5e is presented without normalization as % input. **(f)** The ChIP analysis data from Fig. 5f is shown without normalization as % input.

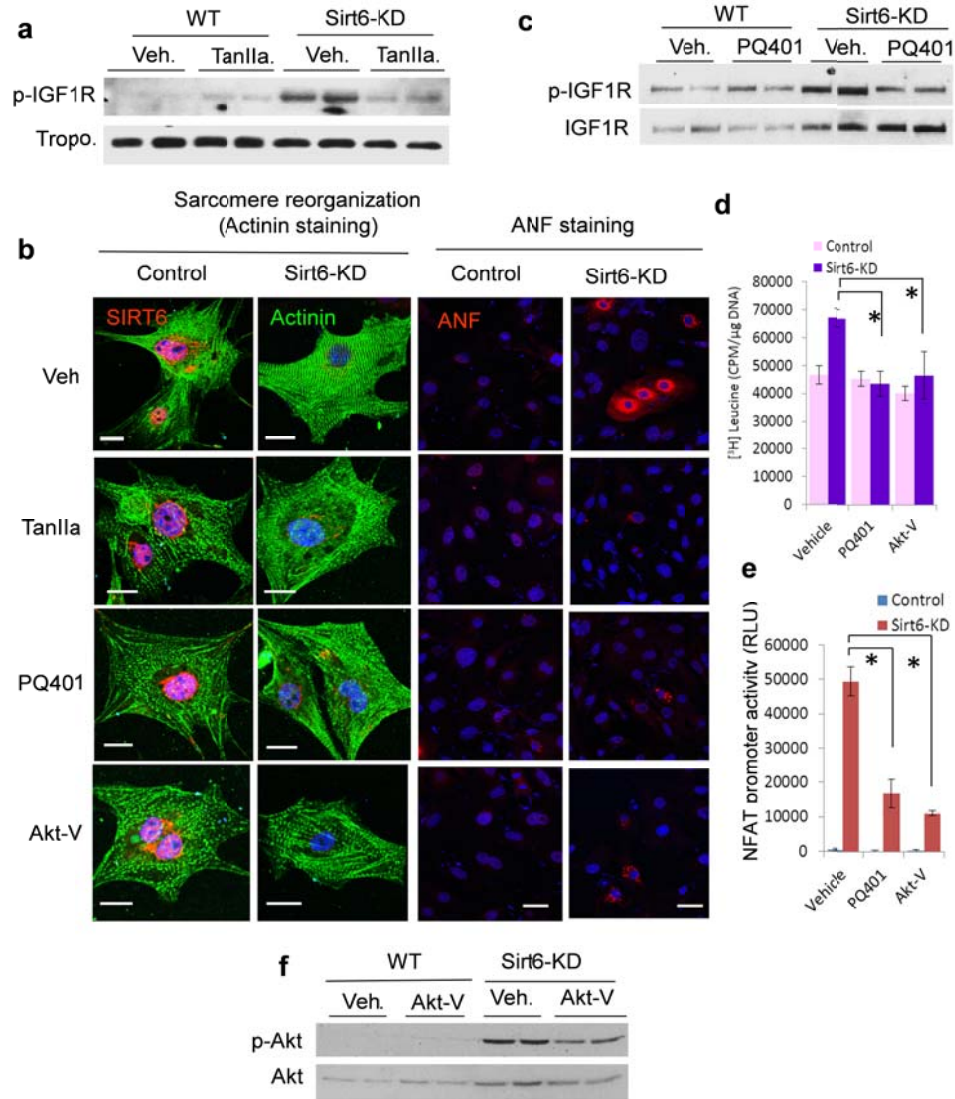
Supplementary Figure 11



Supplementary figure 11: c-Jun inhibition blocks hypertrophy of SIRT6-KD cardiomyocytes.

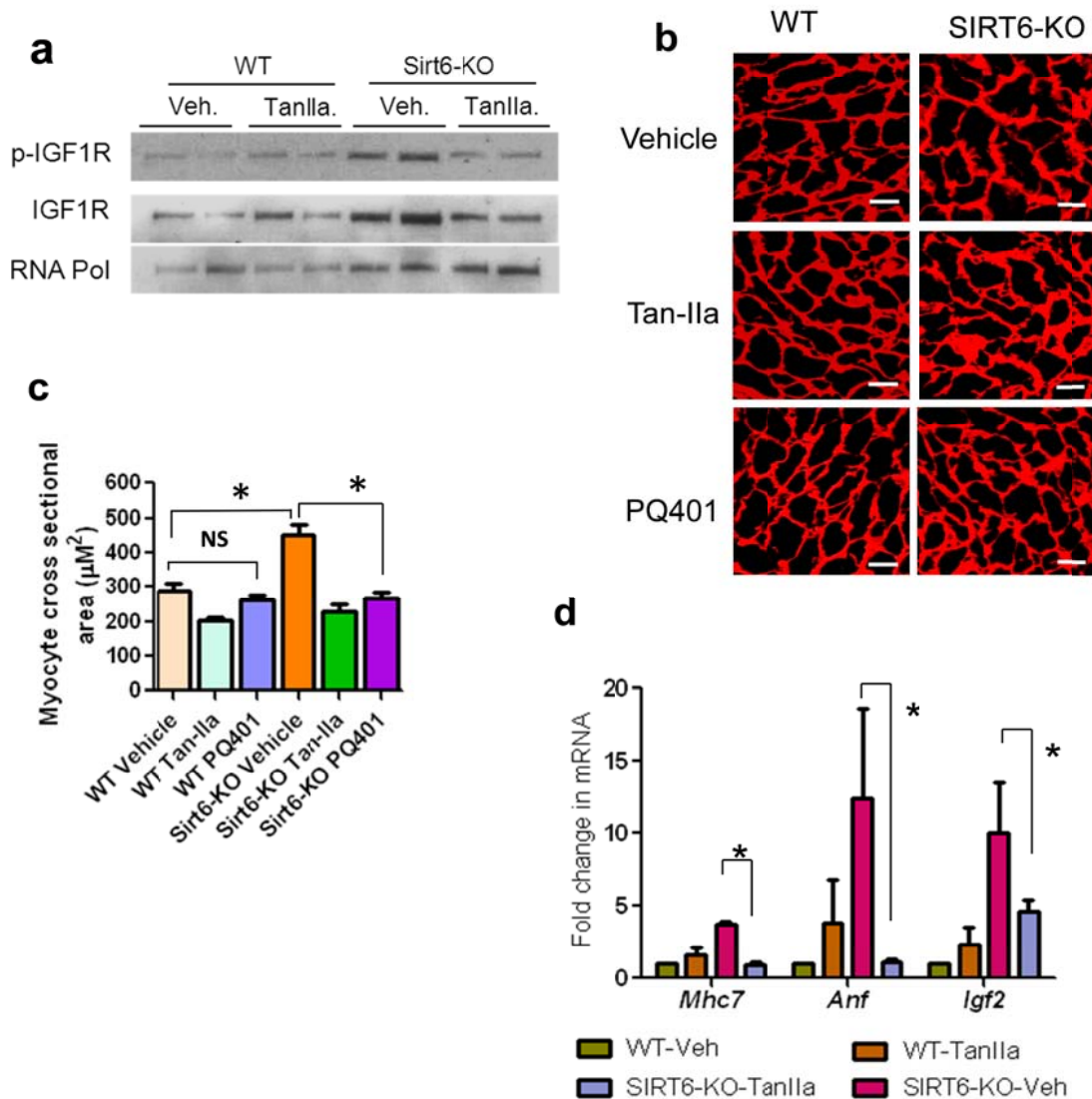
(a) A promoter/ reporter assay showing that adenovirus vector expressing dominant negative (DN) form of c-Jun inhibits activity of endogenous c-Jun. **(b)** Western blots showing SIRT6 knockdown in neonatal rat cardiomyocytes by use of lentiviral vectors (Open Biosystem). **(c)** Confocal imaging of control and SIRT6KD cardiomyocytes over expressed with dominant-negative (DN) form of c-Jun and stained for actinin to visualize sarcomere organization (scale bar 20 μ M). **(d)** Western blot analysis of IGF1R expression and phosphorylation in WT and SIRT6-KD cardiomyocytes infected with control or c-Jun-DN adenovirus. **(e)** Real-time PCR analysis of representative IGF2 signaling related genes in WT and SIRT6-KD cardiomyocytes infected with control or c-Jun-DN adenovirus. Mean \pm SD, n=3-5, *p<0.001.

Supplementary Figure 12



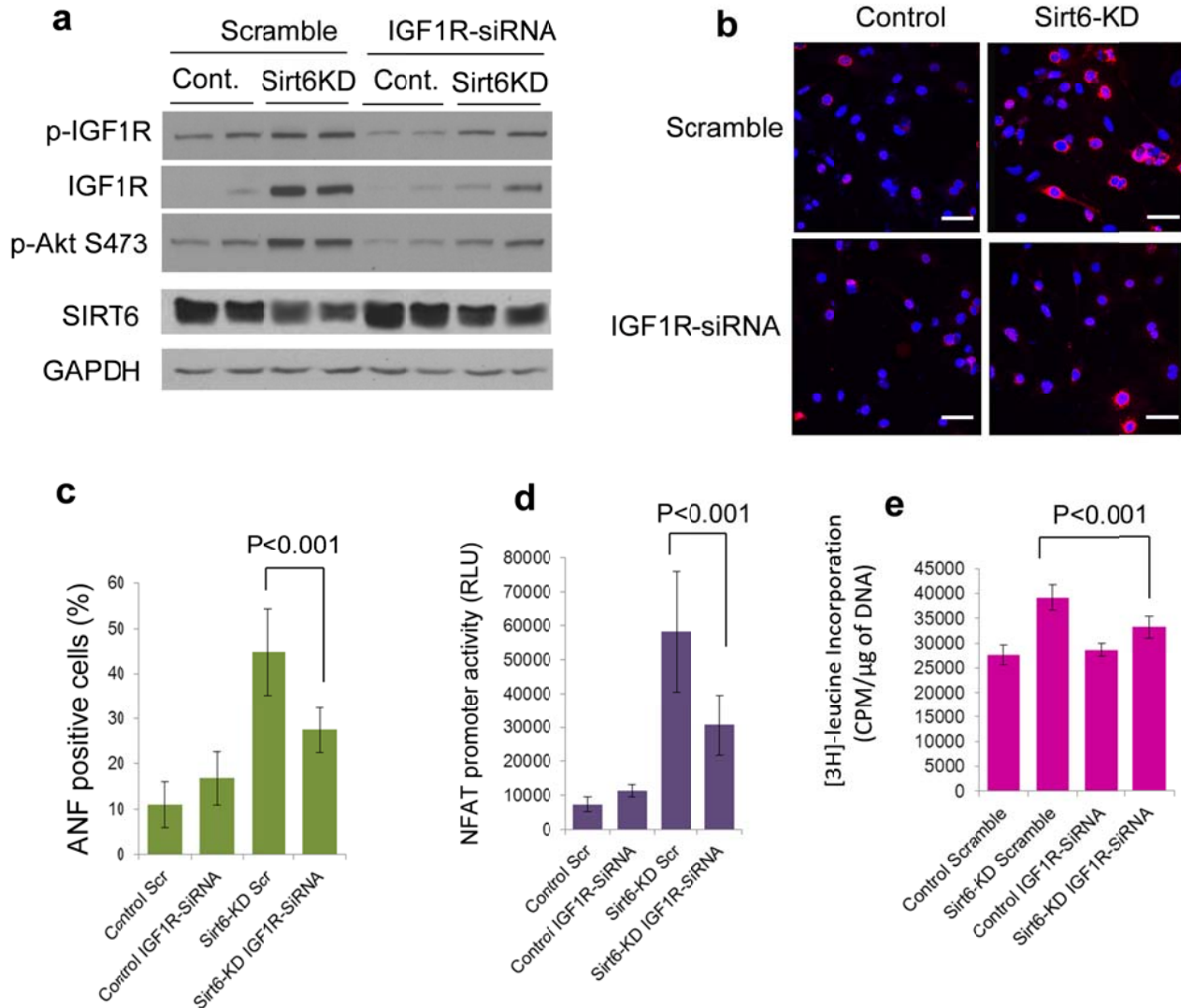
Supplementary figure 12: AP1 and IGF-Akt signaling inhibitors block hypertrophy of SIRT6-deficient cardiomyocytes in vitro. (a) Western analysis of IGF1R phosphorylation in WT and SIRT6KD neonatal rat cardiomyocytes treated with vehicle or Tan-IIa. (b) Sarcomere organization and ANF expression in cardiomyocytes treated with different inhibitors. Sarcomeric organization was determined by staining cells for sarcomeric α -actinin (green) and SIRT6 (red) (Scale bar 20 μ M). ANF release (red) was determined by staining cells with an anti-ANF antibody. DAPI stain was used to mark position of nuclei (Scale bar 50 μ M). (c) Western analysis of IGF1R phosphorylation in cardiomyocytes treated with vehicle or the inhibitor, PQ401 (d) Control and SIRT6KD cardiomyocytes were treated with vehicle, PQ401 (10 μ M) or Akt-V (0.5 μ M) for 48 hours. Incorporation of [³H]-leucine into cellular protein was determined and normalized to DNA content of cells. (e) Cardiomyocytes were infected with control and NFAT-luciferase promoter/reporter expressing adenovirus vector and treated with vehicle, PQ401 (10 μ M) or Akt-v (0.5 μ M). Cell extracts were analyzed for luciferase activity after 24 hours. Mean \pm SD, n = 5-8, *P < 0.001. (f) Western analysis of Akt expression and phosphorylation in WT and SIRT6KD cardiomyocytes treated with vehicle or Akt-v inhibitor.

Supplementary Figure 13



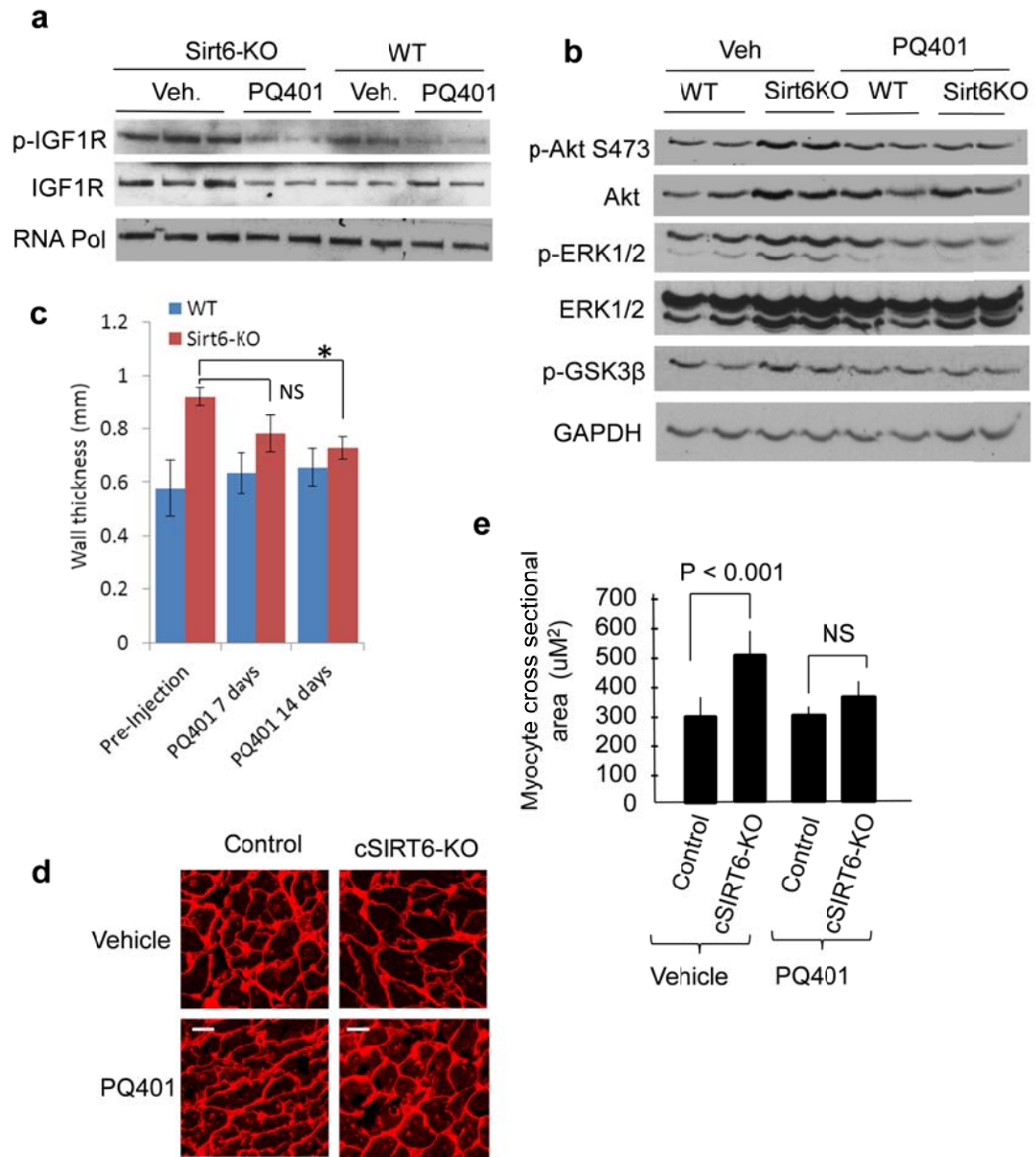
Supplementary figure 13: AP1 and IGFR inhibitors block hypertrophy of SIRT6KO hearts. (a) Western analysis of IGF1R expression and phosphorylation in heart lysates of WT and SIRT6-KO mice treated with vehicle or Tan-IIa. (b and c) Cardiomyocyte cross sectional area in WT and SIRT6KO mice injected with vehicle or AP1 inhibitor, Tan-IIa (1 mg/kg/day) or IGF1R inhibitor, PQ401 (1 mg/kg/day) for two weeks. Mean \pm SD, n= 6 (scale bar 10 μ M). (d) Real-time PCR analysis of cardiac fetal genes (*Mhc7* and *Anf*) and *Igf2* gene expression in WT and SIRT6-KO mice injected with vehicle or Tan-IIa. (mean \pm SD, n=4-5 mice, *p<0.001).

Supplementary Figure 14



Supplementary figure 14: IGF1R knockdown blocks hypertrophy of SIRT6-KD cardiomyocytes. (a) Western analysis showing IGF1R knockdown in control and SIRT6-deficient (SIRT6KD) cardiomyocytes by use of siRNA. (b to e) IGF1R knockdown blocks hypertrophic response of SIRT6-deficient cardiomyocytes, as measured by ANF release from nuclei (b and c), NFAT promoter/reporter activity (d), and ³H-Leucine incorporation into cellular proteins (e). Values are mean ±SD, n = 4-8. Scr indicates scrambled siRNA. Panel b scale bar 50μM.

Supplementary Figure 15



Supplementary figure 15: IGF1R inhibitor, PQ401 blocks cardiac hypertrophy of SIRT6KO mice. (a and b) Western analysis of the heart lysate of wild-type and SIRT6-KO mice treated with vehicle or PQ401 (1 mg/kg/day) for two weeks. **(c)** Analysis of left ventricular wall thickness in WT and SIRT6KO mice injected with vehicle or IGF1R inhibitor, PQ401 for indicated time periods. Mean \pm SD, $n=4-5$, $*p<0.001$. **(d and e)** Cardiomyocyte cross sectional area in the WT and cardiac-specific SIRT6-KO (cSIRT6KO) mice injected with vehicle or PQ401 for two weeks. Mean \pm SD, $n= 6-8$. Panel d scale bar 10 μ M.

Supplementary table 1: Clinical characteristics of patients and their corresponding cardiac SIRT6 levels.

Patients	Age	Gender	Diagnosis	Dgn	Dbt	Medical Treatments				SIRT6
						β -Blk	ACEI	Diuretics	Ca-Blx	
Non-failing hearts										
NF1	48	M	MVD	--	--	+	--	+	--	1
NF2	39	F	MVD	--	--	+	--	+	--	1.85
NF3	65	M	MVD	--	--	--	--	+	--	1.5
NF4	50	M	Donor dysfunction	--	--	--	--	--	--	2.4
NF5	76	M	MVD	--	--	--	+	+	--	1.6
NF6	7	F	VSD	--	--	--	--	--	--	0.7
NF7	45	M	Donor dysfunction	--	--	--	--	--	--	0.7
NF8	38	M	Donor dysfunction	--	--	--	--	--	--	0.8
Failing hearts										
F-1	57	M	IsCMP	NA	NA	NA	NA	NA	NA	0.1
F-2	56	M	IdCMP	--	+	+	--	+	--	0.15
F-3	65	F	DCM	--	--	+	+	--	--	0.25
F-4	55	M	IsCMP	--	--	+	--	+	--	0.1
F-5	59	M	IsCMP	--	--	+	--	--	--	0.2
F-6	53	F	IdCMP	--	--	+	+	--	--	0.15
F-7	53	M	IsCMP	--	--	+	+	--	--	0.3
F-8	66	M	DCM	--	--	+	+	+	--	0.35
F-9	43	F	DCM	--	--	+	+	+	--	0.4
F-10	82	M	DCM	+	--	+	--	+	--	0.3
F-11	13	M	IdCMP	NA	NA	NA	NA	NA	NA	0.8
F-12	26	M	DCM	--	+	--	--	--	--	0.15
F-13	61	F	IsCMP	--	--	--	--	--	--	0.5
F-14	68	M	IdCMP	--	--	+	--	+	--	0.1
F-15	57	F	IsCMP	--	--	+	--	+	--	0.4
F-16	70	M	IdCMP	--	--	+	+	+	--	0.5
F-17	66	M	IsCMP	--	--	--	+	+	--	0.35
F-18	60	F	DCM	--	--	--	--	--	--	0.1
F-19	32	F	IdCMP	NA	NA	NA	NA	NA	NA	0.1
F-20	56	F	IsCMP	NA	NA	NA	NA	NA	NA	0.05
F-21	53	M	DCM	--	--	+	+	+	--	0.1
F-22	57	M	IsCMP	--	--	--	--	--	--	0.35
F-23	52	M	IdCMP	--	--	+	+	--	--	0.1
F-24	61	M	IdCMP	--	--	--	--	+	--	0.1

MVD, mitral valve defect; VSD, ventricular septal defect; IsCMP, ischemic cardiomyopathy; DCM, dilated cardiomyopathy, IdCMP, idiopathic dilated cardiomyopathy; Dgn, digoxin, Dbt, dobutamine; β -Blk, β -adrenergic blocker, ACEI, angiotensin converting enzyme inhibitor, Ca-Blx, calcium channel blocker, NA, not available. SIRT6 levels determined by western blotting were normalized against expression of loading control (actin or tubulin) and presented as fold change over NF1.

Table 2: Primer sequences.

Primer Name	Forward Sequence (5'→ 3')	Reverse Sequence (5'→ 3')
Primer sequences used for the ChIP assay with mouse tissues or cell lines		
mAkt1ChIP1	ATTTCATCCTGGGCGATAGC	CCAGAAGCCCCGACTTGATAGTAAC
mAkt1ChIP2	GTA CTGGGTGGATGAGCCCTCAATAG	CAGCGTGGGAAGTGAATCAGTTTGAC
mAkt3ChIP1	GGAGCCATCATGAGCGATGTTAC	TTGATTCAACAGCGCCAGAGG
mAkt3ChIP2	TGCCACAAGACCAGAGCAGTGATTTC	CAATTGGCCTGACCGCACATAAAG
mMapk3ChIP1	CCGGGTGGGTTCCCTTAGCATTACTG	ACCCGGCTTTCCCGCCTAGTTAC
mMapk3ChIP2	GATGGCTCAGGCCATAAG	AGTGGCCCTCAGTAATGC
mGsk3bChIP1	TGCCAGTGCCACTCTAAC	TGTAGTCCAGCGTCCATTG
mGsk3bChIP2	GAACAATGGACGCTGGACTACATGTG	ATGTGGTCAGGACAGTACCTCGAATC
mlgf1rChIP1	CGTGCTCGGCTTTGACCTTC	CGCGAGCTCCTTCCCAAATCCAG
mlgf1rChIP2	TGTAGCCGCTTGGAGTGTGC	CCGCTCAGCGGAGTTAATG
mlgf2rChIP1	GTGGTGGTACACGCTTCTAAC	GGTGACTGGTGGACTAATATGC
mlgf2rChIP2	GTTGTCAGGCCCTCGAGTAG	ACGTGACGTCCTCGTTTACG
mlrs2ChIP1	TCACGCTCATTGGTCCGTCTCG	CCGCCGCACAGTGAGTAACACATC
mlrs2ChIP2	CCCTTTCCCGGCACTATGGAAACC	GGGCGTCATCAGAGCCATTCACTTG
mTorChIP1	CTCAGTGAACCGATTTC	ACGGTTTGGTACCCTAAG
mTorChIP2	TCCCAGACGATCCCTAAC	CCACGAACACTTCCGGTACG
mPtenChIP1	TGTGAGGTGCACTCTATTCACGGAGAC	GGGTTCAATTTGCCGAAAGATGAACG
mPtenChIP2	ACGCGGGAGACACAATAGG	GTCGGCGACAGTCTTTAC
mTsc2ChIP1	GGGCAAGGCATAGCCTAATCG	TGGAGACCTGCCAGGAGTTC
mTsc2ChIP2	CTTGGCTCTGTTGCCAAAG	ATCCTGCAGACCGATGATG
Primers used for the ChIP assay with human cell lines		
hAKT1ChIP-1	TGCTGGCCTGGTGATACG	CAGAGGGCTGGACTCAAAGAC
hAKT1ChIP-2	AGTTGTCCGAGGAACCTTCTG	CCGGGTATGGAATGAGTAAGTG
hAKT3ChIP-1	TATTTGGGTAGGCGTGACTG	AGCACTTCCCTAGTCTTG
hAKT3ChIP-2	CTGGCGACAGAGTGAGATTCC	AAACCAGTCACGCCTACCC
hMAPK3ChIP-1	TGCCTTTCTTCACTCCCTACCTTC	AGCTGAGATTGCACCACTTCAC
hMAPK3ChIP-2	TCGTAGTCCCAGCTCTTTG	TTCCATGCCTCTCAGAGTC
hGSK3bChIP-1	CACCAATCACCGAAGGAGGTACG	AATCAGAGTCGCCGGCCCTTACG
hGSK3bChIP-2	GCAGCTAGGTCTTGCGATTG	GGAGGGCGTCTATAAAGCG
hIGF1rChIP-1	GGAGCGAAGACTGAGTTTG	GGAGCCAGACTTCATTCC
hIGF1rChIP-2	GTTCCGGCTTTCCAGTACG	GGCTGGGAGAGGTTCAATTG
hIGF2rChIP-1	CCCACGCTTGGGAAATAAC	AGTGCCAGGGAGGTAATG
hIGF2rChIP-2	TTGACTCCACGGGCGAGTTAAAG	CAAAGTGCCAGGGAGGTAATG
hIRS2ChIP-1	CTCGGTGCGGATGTGTTACTC	TGCTGCTGCTGCTGGTGTG
hIRS2ChIP-2	GCGCTATGGAAACCGCACTTCTCCG	GCGCCATTCACTTGTCAGCTTGTCG
hmTORChIP-1	TCCATAAAGAGCGCTAGCC	CTCCCGGTGTAATTCTGAGAG
hMTORChIP-2	AAGGGAATCCTAGCAGCGCCGTACC	AAGTTCAGGACCCGGCTTCTCCAG
hPTENChIP-1	TGCAAGGGAGCCGGATGAGGTGATAC	CACCGCTGTCGGATCACAATCGTTCCG

hPTENChIP-2	CTGCTCAACGCACCCATCTCAGCTTTC	AGCAACGCGAGGCGAGGATAACG
hTsc2ChIP-1	CGAGCCTGAGATGCTTGAC	CTGGGCAAGCTCATCTACC
hTsc2ChIP-2	GCCGAGGCAGGAGAATAATTTG	TAGTGATGGGCCAGATAC
Primers used for the real-time PCR analysis		
Akt1	TGGACAAGGACGGGCACATCAAG	TACTCCGGCGTTCCGCAGAATG
Akt2	TGTGGGCGACTTCATCCTTTC	TTCGGCAAGGTCATTCTGGTTCG
Akt3	CATAGGCTATAAGGAGAAACC	TTGGATAGCTTCCGTCCAC
Insulin1	GAAGTGGAGGACCCACAAGTG	CTGAAGGTCCCCGGGGCT
Insulin2	TGCTGATGCCCTGGCCTGCTCT	CTGGTCCCACATATGCACATGCA
Gsk3 β	TCCATTCCCTTTGGAATCTGC	CAATTCAGCCAACACACACAGC
Irs2	CGCCGCTACAGCGAAGTACT	CGGACGCCAAGCACAAGT
FoxO1	TCGTACGCCGACCTCATCA	CTGTCGCCCTTATCCTTGAAGT
Pten	AATCCCAGTCAGAGGCGCTATGT	GATTGCAAGTTCGCCACTGAACA
mTor	GAGAACCAGCCATAAGA	ACCAGCCAATGTAGCACT
Igf1	TCATGTGCTCTTACACCTCTTCT	CCACACACGAACTGAAGAGCAT
Igf2	ACAACCTCGATTTGAACCACATTC	GAGAGCTCAAACCATGCAAACCT
Igf1r	GTGGGGGCTCGTGTCTTCTC	GATCACCGTGCAATTTTCCA
Igf2r	GGGAAGCTGTTGACTCCAAAA	GCAGCCCATAGTGGTGTGAA
Insr	ATGGGCTTCGGGAGAGGAT	GGATGTCCATACCAGGGCAC
Mapk3	AGGGCTACACCAAATCCATC	GGGAACCCAAGATACCTAGAA
Igfbp3	GAGTGTGGAAAGCCAGGTTGTC	GCATGGAGTGGATGGAACCTG
At1	GTTCTGCTCACGTGTCTCA	CATCAGCCAGATGATGATGC
At4	GATGAAGGCACACCATCCAT	AAACCAGCCTTTCTCGTTTCT
GAPDH	ATGGTGAAGAAGGTCCGGTGTGA	AATCTCCACTTTGCCACTGC
RPL32	ACAACAGGGTGCGGAGAAGATT	GTGACTCTGATGGCCAGCTGT
18S	GGACAGGATTGACAGATTGATAG	CTCGTTTCGTTTATCGGAATTAAC