

Comparison of insulin signaling pathway and gene expression in *Atf4-/-* liver and hepatocytes. (**A**) Quantification of Akt phosphorylation in *Atf4-/-* liver and muscle at basal and insulin stimulated condition. (**B**) Phosphorylation of GSK-3 β in *Atf4-/-* liver (upper panel) or cultured hepatocytes at basal and insulin stimulated condition. (**C**) Expression of *Pgc1* α , *Ppar* γ and *Mcad* in *Atf4-/-* liver or cultured hepatocytes. N.S.: not significant.



Quantification of *Atf4* transgene. (**A**) Atf4 protein expression in several tissues of $\alpha 1(I)$ Collagen-Atf4 transgenic mice. (**B**) Atf4 protein expression in bone of WT and $\alpha 1(I)$ Collagen-Atf4 transgenic mice. (**C**) Expression of *CMV*-Atf4 transgene in various tissues relative to pancreas in *CMV*-Atf4 transgene in various tissues relative to pancreas in *CMV*-Atf4 transgene in various tissues relative to bone in $\alpha 1(I)$ Collagen-Atf4 transgenic mice. (**D**) Expression of $\alpha 1(I)$ Collagen-Atf4 transgene in various tissues relative to bone in $\alpha 1(I)$ Collagen-Atf4 transgenic mice. N.D.: not detected.



Levels of Adipokines in mutant mice. Serum (A) leptin, (B) resistin and (C) adiponectin levels in $\alpha 1(I)$ Collagen-Atf4, $\alpha 1(I)$ Collagen-Cre; Atf4fl/fl and $\alpha 1(I)$ Collagen-Cre; Crebfl/fl mice.



Generation of *Atf4osb-/-* mice. (**A**) Targeting construct for conditional inactivation of *Atf4*. White box, exon; black triangles, LoxP sites. (**B**) Southern blot analysis of *Atf4-flox* clone. The 5' probe detected a 12 kb WT and 14 kb targeted band while 3' probe detected 12 kb WT and 14 kb targeted band. (**C**) PCR genotyping of *Atf4osb-/-* mice. WT and floxed allele yield 600 bp and 640 bp products, respectively. $\alpha 1(l)$ collagen-Cre transgenic mice harbor a transgene-specific band.



Level of osteocalcin in mutant mice. Serum total osteocalcin level in $\alpha 1(I)$ Collagen-Atf4 and $\alpha 1(I)$ Collagen-Cre; Atf4fl/fl mice.



Serum uncarboxylated osteocalcin levels in *Esp+/-* and *Esp+/-*; *Atf4+/-* mice. Error bars, mean + SEM. *: P<0.05 Table S1 Insulin sensitivity analysis of eight week-old Atf4_{osb} -/- mice analyzed by

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Genotype	Control (n=3)	<i>Atf4_{osb} -/-</i> (n=3)
Body weight (g)	25.0 ± 1.1	20.0 ± 0.3*
Basal glucose (mg/dl)	120.0 ± 12.7	94.7 ± 9.0
Clamp glucose (mg/dl)	101.7 ± 8.7	118.0 ±14.0
Glucose infusion rate (mg/kg/min)	47.2 ± 4.2	69.3 ± 2.5*
Basal hepatic glucose production (mg/kg/min)	10.4 ± 0.8	13.2 ± 0.9
Clamp hepatic glucose production (mg/kg/min)	2.4 ± 1.1	0.6 ± 1.6
Liver action (%)	77.5 ± 11.2	90.5 ± 9.0
Glucose turnover (mg/kg/min)	49.6 ± 4.2	69.9 ± 3.0*
Whole body glycolysis (mg/kg/min)	32.9 ± 7.6	28.2 ± 1.6
Glycogen synthesis (mg/kg/min)	16.7 ± 7.2	41.7 ± 4.3*

*: P < 0.05