

## Supplementary Information

### Identification of *Creb3l4* as an essential negative regulator of adipogenesis

Tae-Hyun Kim<sup>1</sup>, Seong-Ho Jo<sup>1,2</sup>, Hyeonjin Choi<sup>1</sup>, Joo-Man Park<sup>1</sup>, Mi-Young Kim<sup>1</sup>, Hiroshi Nojima<sup>3</sup>,  
Jae-Woo Kim<sup>1,2</sup>, Yong-Ho Ahn<sup>1,2,\*</sup>

<sup>1</sup>Department of Biochemistry and Molecular Biology, Yonsei University College of Medicine, Seoul 120-752, Republic of Korea

<sup>2</sup>Brain Korea 21 PLUS Project for Medical Sciences, Yonsei University College of Medicine, Seoul 120-752, Republic of Korea

<sup>3</sup>Department of Molecular Genetics Research Institute for Microbial Diseases Osaka University, Osaka, Japan

**\*Corresponding author:** Yong-Ho Ahn, MD, PhD;

Dept. of Biochemistry and Molecular Biology,

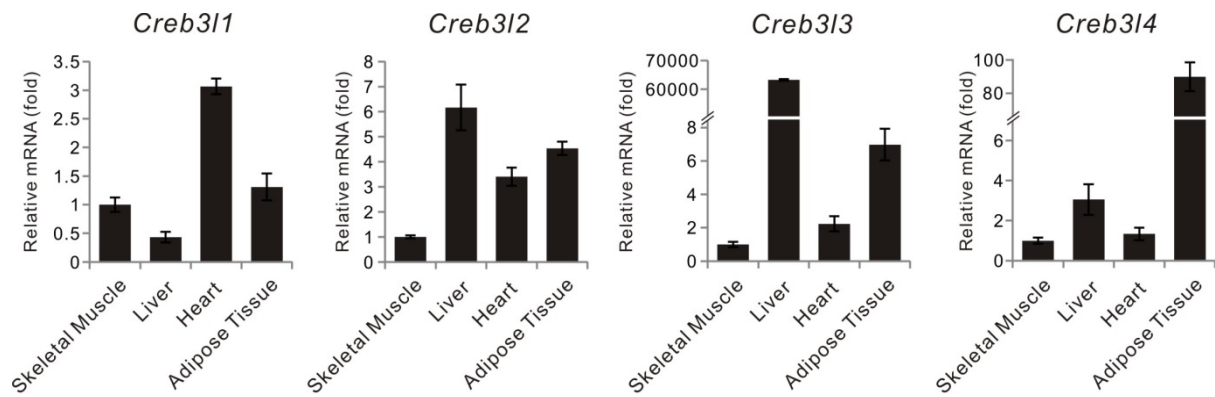
Yonsei University College of Medicine,

50-1 Yonsei-ro, Seodaemun-gu, Seoul 120-752, Republic of Korea.

Tel: +82-2-2228-0835, Fax: +82-2-312-5041,

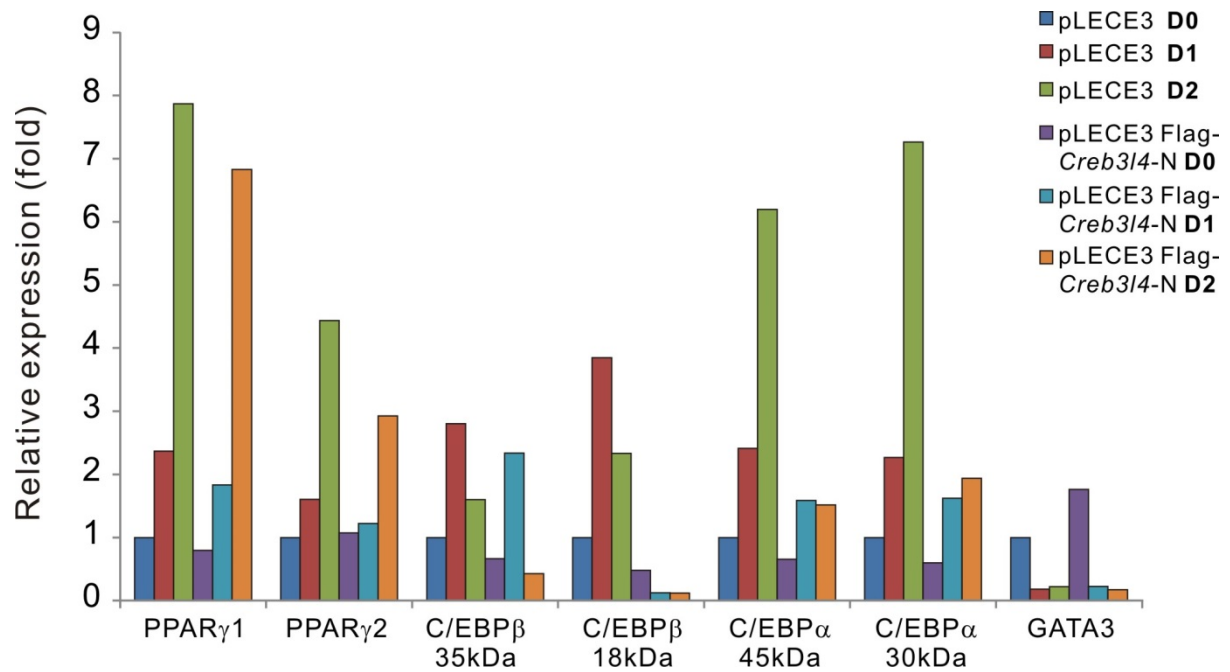
e-mail: [yha111@yuhs.ac](mailto:yha111@yuhs.ac)

## Supplementary Figure S1



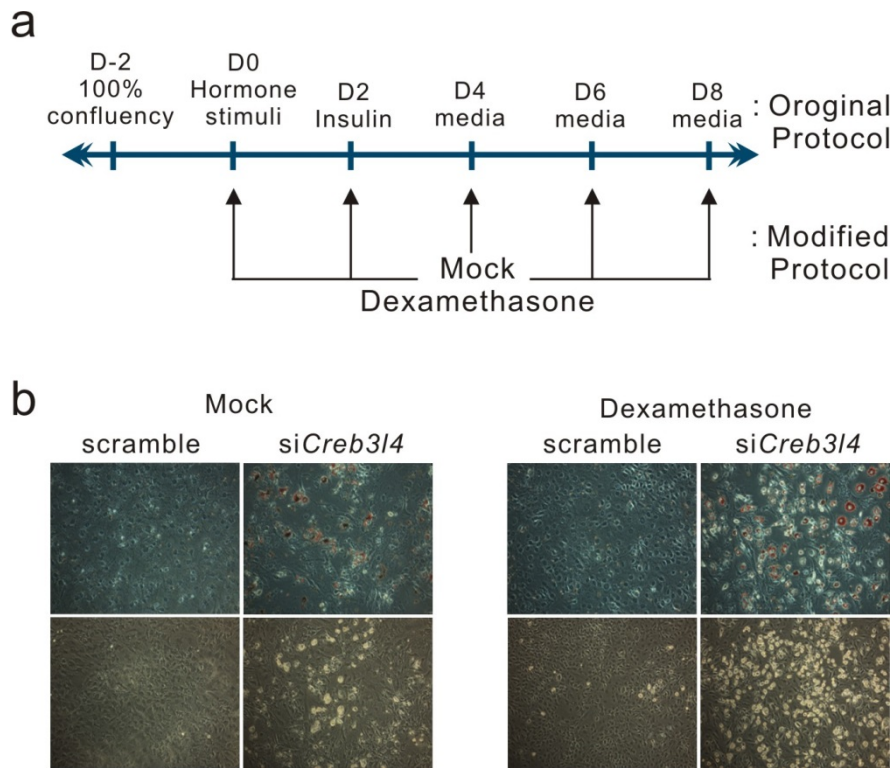
**Supplementary Figure S1.** Expression of CREB3L-family members in mice. The mRNA levels of *Creb3l* family members in the liver, heart, muscle, and white adipose tissue (WAT) was measured by real-time PCR; expression levels were normalized to that of the *Rplp0* (*36B4*) control. Error bars represent mean  $\pm$  SD.

Supplementary Figure S2



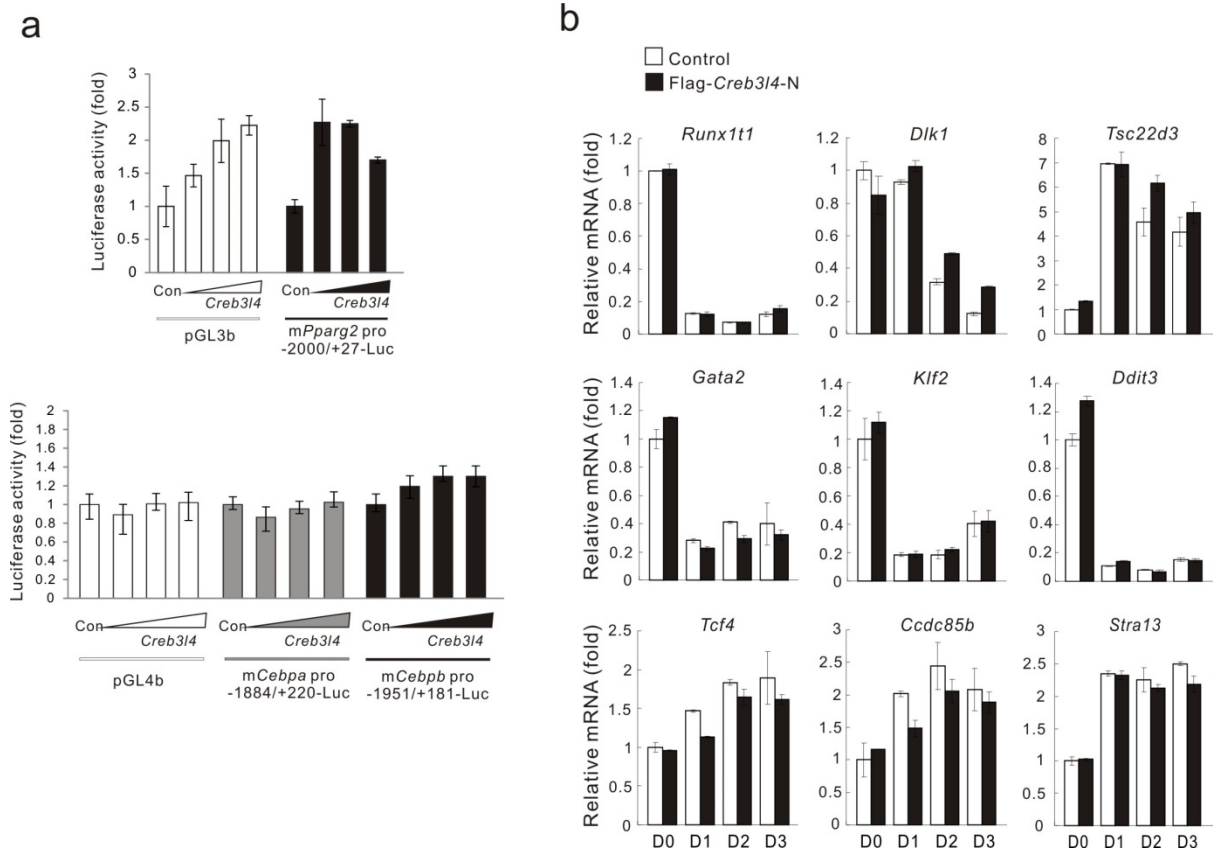
**Supplementary Figure S2.** Quantification analysis of western blot data from Figure 2b. Signal intensities were quantitated using Image J after normalization to GAPDH levels.

### Supplementary Figure S3



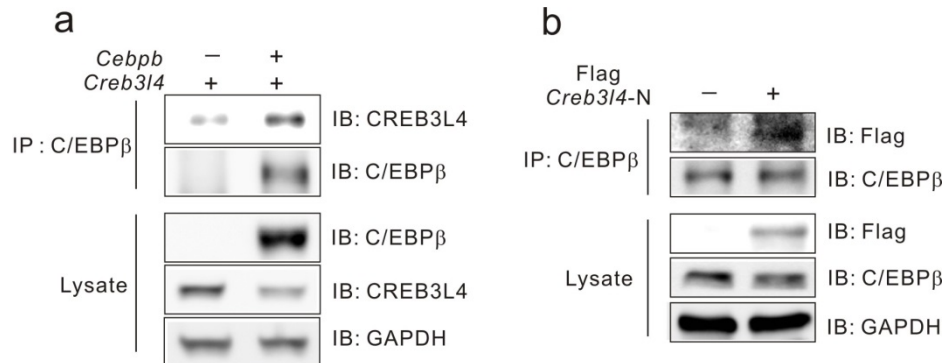
**Supplementary Figure S3.** Effect of *Creb314* ablation on adipocyte differentiation (a) A modified protocol for inducing adipocyte differentiation. Induction of adipogenesis in 3T3-L1 cells expressing si*Creb314*. (b) Cells were treated with either mock or dexamethasone (1  $\mu$ M) in 10% FBS–DMEM. The medium was changed every 2 days. After 8 days, the cells were stained with oil-red O. Microscopic views representative of three independent experiments are shown.

## Supplementary Figure S4



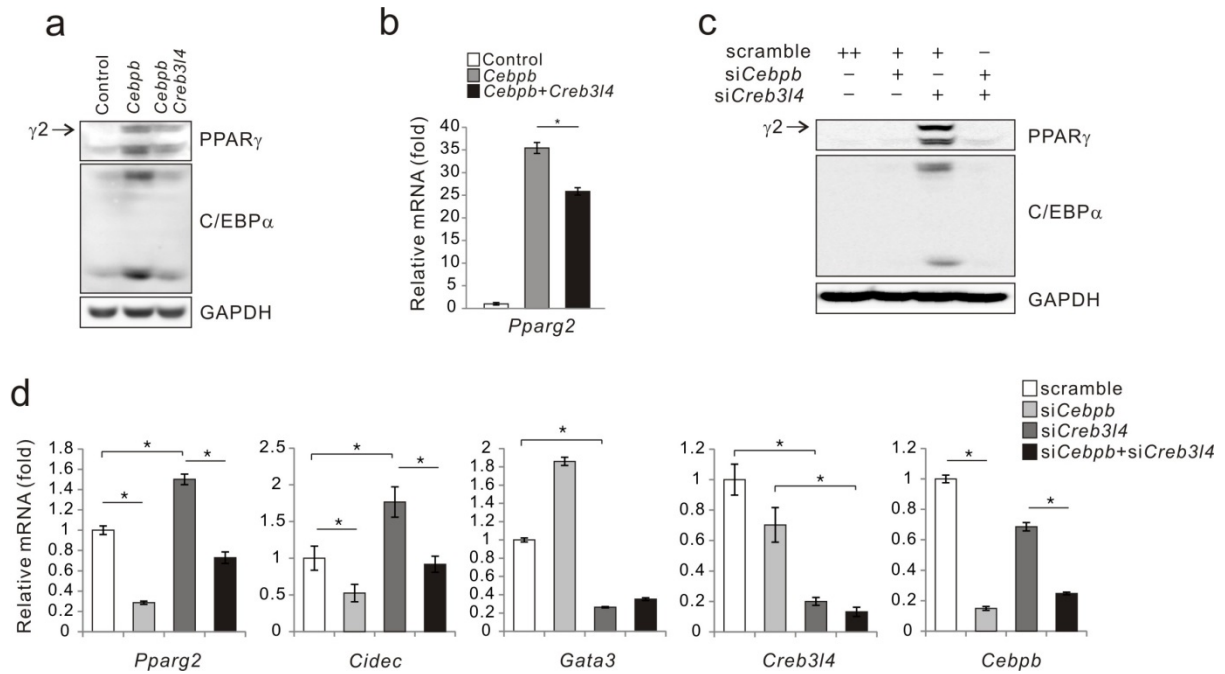
**Supplementary Figure S4.** Effect of CREB3L4 on the promoter activities of adipogenic factors and gene expression of negative transcriptional regulators. (a) Effect of CREB3L4 on the promoter activities of *Pparg2*, *Cebpa*, and *Cebpb*. NIH-3T3 cells were cotransfected with pcFlag-*Creb314* (0, 50, 100, or 200 ng), luciferase reporters (500 ng; mouse *Pparg2* promoter covering -2000/+27, mouse *Cebpa* promoter covering -1884/+220, mouse *Cebpb* promoter covering -1951/+181), and *Renilla* luciferase plasmid. The total amount of transfected plasmid was adjusted to 200 ng by adding empty vector. Luciferase activities were normalized to *Renilla* luciferase activities to adjust for transfection efficiency. Normalized activities are shown as the mean  $\pm$  SD of three independent experiments performed in triplicate, and are expressed as fold-increase relative to the basal activity. (b) The mRNA levels of negative regulators of adipogenesis, as determined by real-time PCR. 3T3-L1 preadipocytes cells were transfected with Flag-*Creb314* (N) by electroporation, and differentiation was induced with medium containing DMI (0.5 mM 3-isobutyl-1-methylxanthine [IBMX; M], 1  $\mu$ M dexamethasone [D], and 1  $\mu$ g/ml of insulin [I]). The cells were harvested at the indicated times. Error bars represent mean  $\pm$  SD.

## Supplementary Figure S5



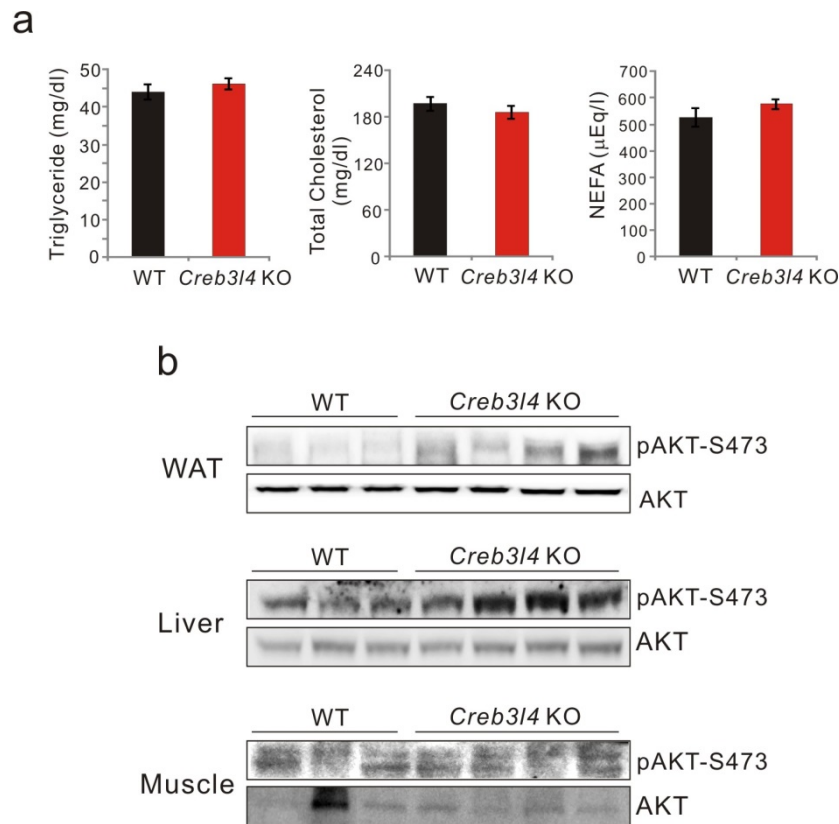
**Supplementary Figure S5.** Interaction between C/EBPβ and CREB3L4. (a) NIH-3T3 cells were co-transfected with *Creb3l4* and *Cebpβ* expression vectors and then immunoprecipitated using an anti-C/EBPβ antibody. (b) Interaction of endogenous C/EBPβ and CREB3L4 in 3T3-L1 cells. Cells were transfected with Flag-tagged *Creb3l4-N* expression vector and precipitated with anti-C/EBPβ antibody.

## Supplementary Figure S6



**Supplementary Figure S6.** Effect of C/EBP $\beta$  on siCreb3l4-induced PPAR $\gamma 2$  expression. (a) 3T3-L1 cells were transfected with *Cebpb* or *Cebpb/Creb3l4* and stimulated to differentiate with 10% FBS–DMEM containing insulin (1  $\mu\text{g}/\text{ml}$ ) every 2 days. Immunoblot of transcription factors involved in adipogenesis in 3T3-L1 cells at day 8. (b) 3T3-L1 cells were transfected with constructs containing *Cebpb* or *Cebpb/Creb3l4*. Expression of *Pparg2* in 3T3-L1 cells expressing *Cebpb* or *Cebpb/Creb3l4* at day 0, as analyzed by real-time PCR. (c) Immunoblot of C/EBP $\alpha$  and PPAR $\gamma 2$  at day 8. siCebpb and siCreb3l4 were transfected into 3T3-L1 cells. The cells were stimulated to differentiate with 10% FBS–DMEM containing insulin (1  $\mu\text{g}/\text{ml}$ ) every 2 days. (d) The mRNA levels of *Pparg2*, *Cidec*, *Gata3*, *Creb3l4*, and *Cebpb* in 3T3-L1 cells transfected with siCebpb and siCreb3l4 at day 0. Values are expressed as mean  $\pm$  SEM,  $n = 3$  per group, \* $P < 0.05$ .

## Supplementary Figure S7



**Supplementary Figure S7.** The phenotype of *Creb3/4* KO mice (a) Serum lipid profiles, triglyceride, total cholesterol, and non-esterified fatty acid (NEFA) levels were measured by enzymatic methods using an autoanalyzer (n = 7–15). Values are expressed as mean  $\pm$  SEM. (b) Mice fed HFD were fasted for 6hr and then received insulin (5U/kg) by intraperitoneal injection. After 5 min, the mice were sacrificed. One gram of each tissue was homogenized in RIPA buffer and subjected to western blot using anti-pAKT-S473, anti-AKT-total antibodies.



**Table S1.** List of primer sequence for gene-specific real-time PCR.

Gene	Forward (5' → 3')	Reverse (5' → 3')
<i>Creb3l1</i>	TCTTTGATGACCCTGTGCTGGA	TCCTCCATCTTGACAGGCACAA
<i>Creb3l2</i>	CAAGCTGAGCGAGCTGTCAGAG	CCACCTCCATTGACTCGCTCTT
<i>Creb3l3</i>	GACAGCATGGAGGTCCTTGA	TCCCAGGATGCAATTTAGGA
<i>Creb3l4</i>	ATATCTTCTCGACGGGATCCTT	TCCCTACCAGGAGATGTTTC
<i>Pparg2</i>	CCAGAGCATGGTGCCTTCGCT	CAGCAACCATTGGGTCAGCTC
<i>Cebpa</i> □	GAACAGCAACGAGTACCGGGTA	GCCATGGCCTTGACCAAGGAG
<i>Cebpb</i>	CAAGCTGAGCGACGAGTACA	CAGCTGCTCCACCTTCTTCT
<i>Gata3</i>	AACAGACCCCTGACTATGAAGAA	GGTTGAAGGAGCTGCTCTTGGG
<i>Gata2</i>	GGCACCTGTTGTGCAAATTGTCA	CCTTCTTCATGGTCAGTGGC
<i>Runx1t1</i>	GGCCAGCGGTACAGTCCAAATA	GATAGGAGTCCCTGTAGTGGTG
<i>Dlk1</i>	GACCCACCCTGTGACCCC	CAGGCAGCTCGTGCACCCC
<i>Tsc22d3</i>	GCTGCACAATTTCTCCACCT	GAGGTCCATGGCCTGCTCAAT
<i>Klf2</i>	GGCCCCAGCTTCGGCGGC	TGAGGAGACCGCGCGTGGCA
<i>Ddit3</i>	TGAGTCCCTGCCTTTCACCTTG	GGTCCTCTGTCAGCCAAGCTA
<i>Tcf4</i>	AGAGATGAGAGCGAAGGTGGT	CTGTTCGTTCCCTCCGGGCC
<i>Ccdc85b</i>	CTGATGCAGGAGGTGAATCG	CGAGTCCAGAAAGCAGCAGA
<i>Stra13</i>	GGAAGGAACTGGTGAGCAGAC	GCAGCCTCTAGTACGAAGATCC
<i>Cidec</i>	TCCAGGACATCTTGAAACTT	GGCTTGCAAGTATTCTTCTGT
<i>Lxra</i>	GAGAAGCTGGTGGCTGCCCA	AGCTGTAGGAAGCCAGGGAG
<i>Adipoq</i>	ATGGCAGAGATGGCACTCCT	CCTTCAGCTCCTGTCATTCC
<i>Pepck</i>	ACACACACACATGCTCACAC	ATCACCGCATAGTCTCTGAA
<i>Rplp0</i>	TGGCCAATAAGGTGCCAGCTGCTG	CTTGTCTCCAGTCTTTATCAGCTGCAC