

Supplementary Section to:

Hyperlipidemia and hepatitis in liver-specific CREB3L3 knockout mice generated using a one-step CRISPR/Cas9 system

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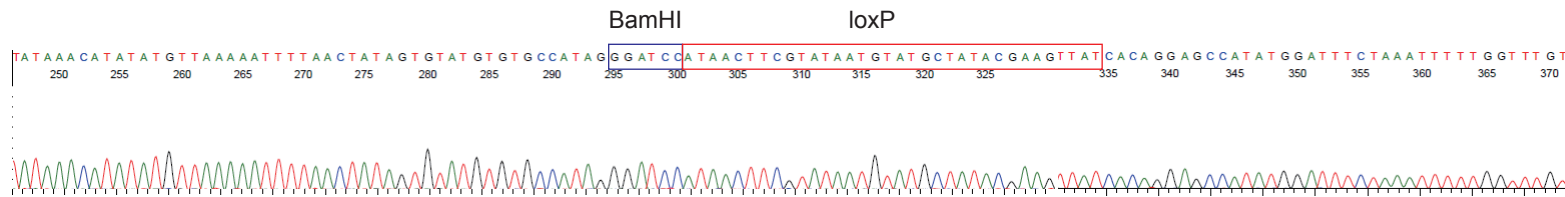
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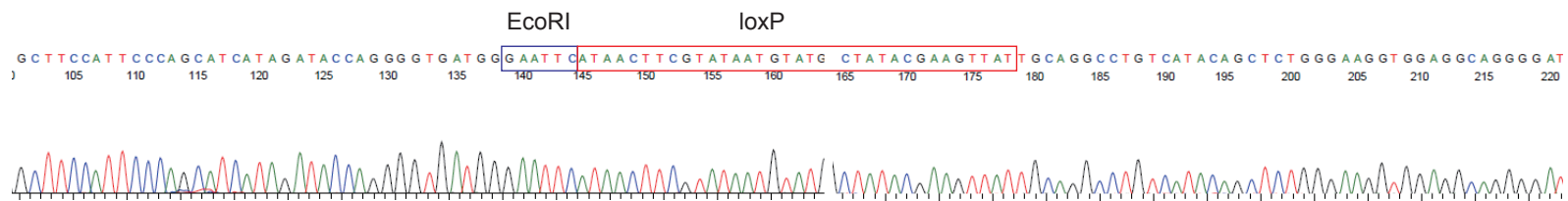
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Creb3l3 intron 3



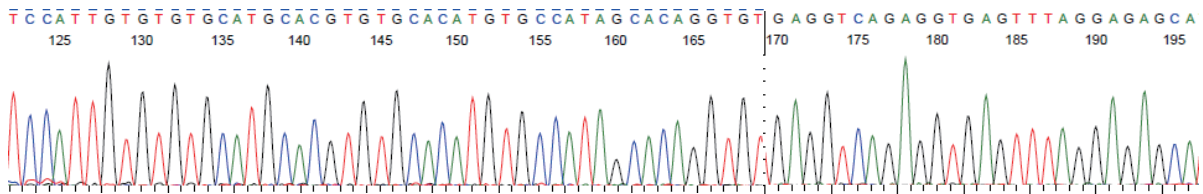
Creb3l3 intron 11



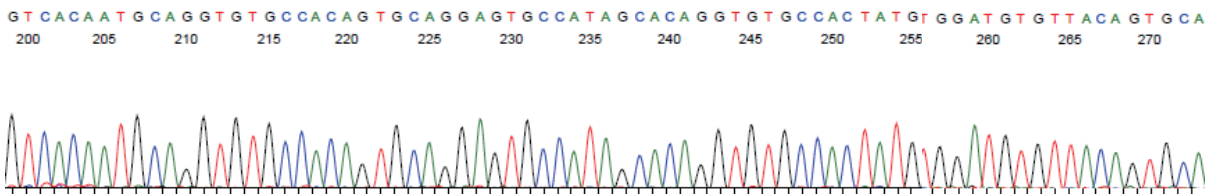
Supplementary Figure 2 The integration of loxP sites in the intron 3 and 11 of the *Creb3l3* genome. The sequences across the targeting regions confirmed correct fusion of loxP sites to the intron 3 and 11 of the *Creb3l3* genome.

a

Seed sequence PAM
GTGTATGT **GTGCCATAGCACAGG** i3 targeting site
TCCATTGTGTGTGCATGCACGTGTGCACATGTGCCATAGCACAGGTGTGAGGTCAGAGGTGAGTTTAGGAGAGCA
chr1:134783344-134783358
▲
134783344



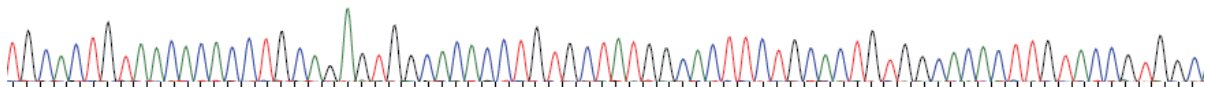
GTGTATGT **GTGCCATAGCACAGG** i3 targeting site
GTCACAATGCAGGTGTGCCACAGTGCAGGAGTGCCATAGCACAGGTGTGCCACTATGTGGATGTGTTACAGTGCA
chr1:181415553-181415567
▲
181415553



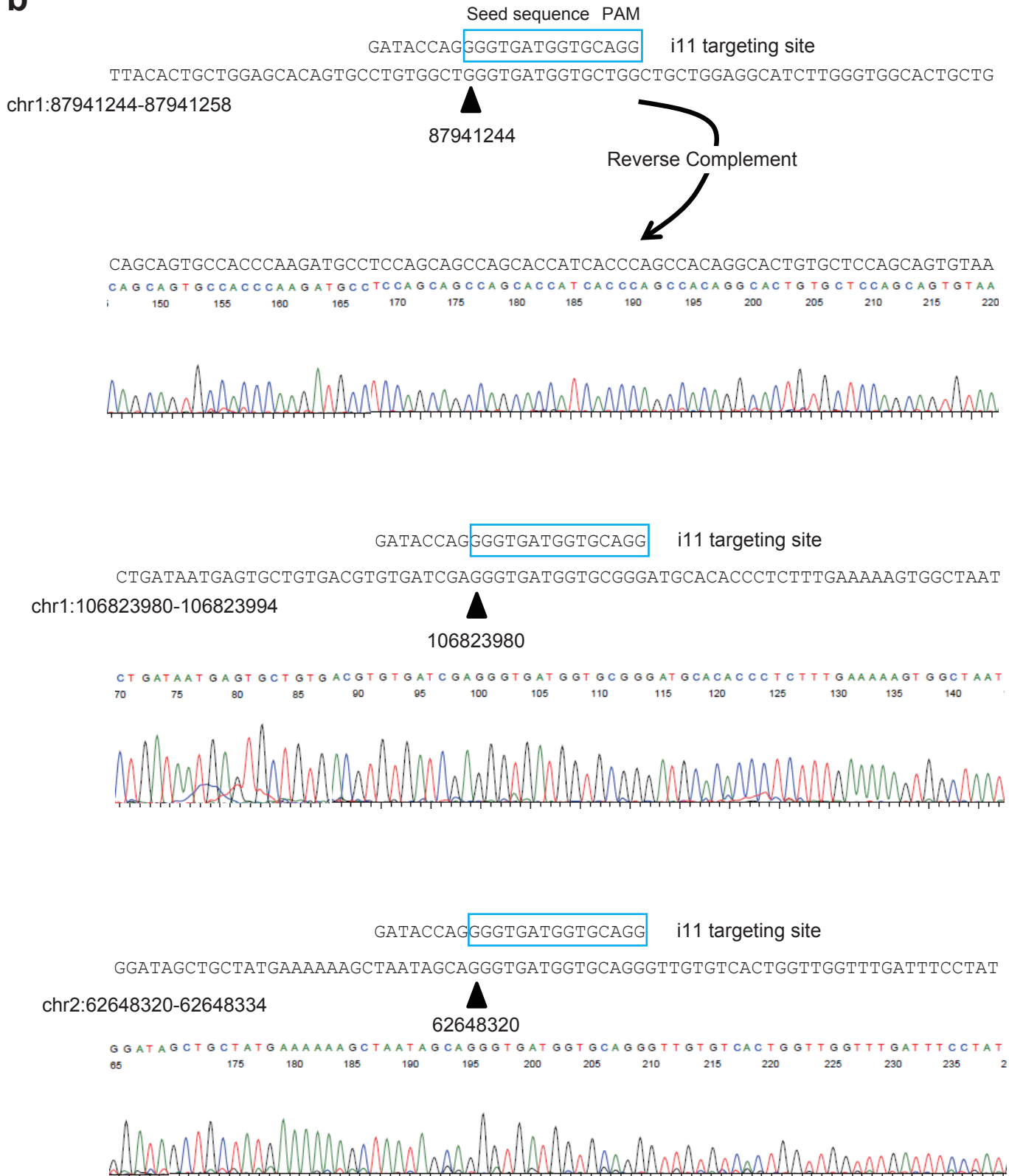
GTGTATGT **GTGCCATAGCACAGG** i3 targeting site
GCCACGGTACAAGTGTGCCACAGTGCAGAAGTGCCATAGCACAGGTGTGCCACTCTGCAGGTGTGTTACAGTGCA
chr1:181415729-181415743
▲
181415729

Reverse Complement

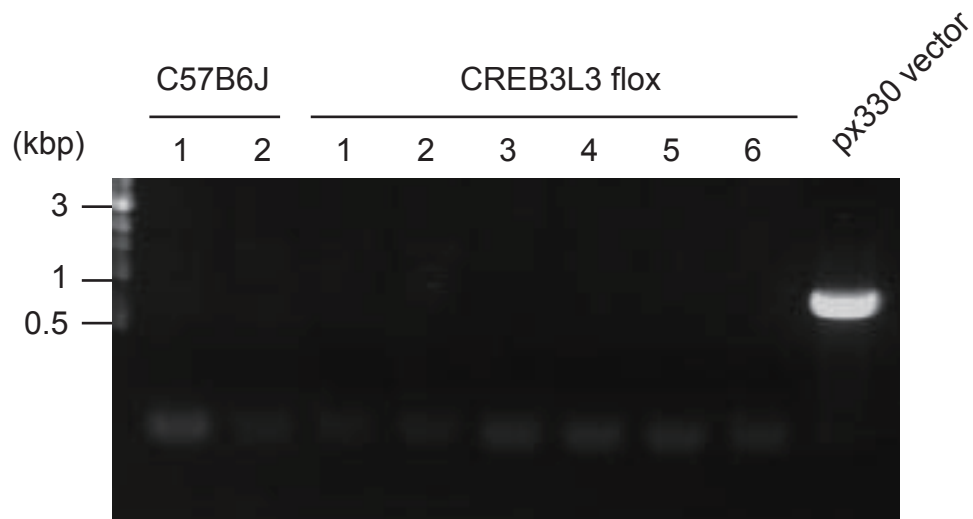
TGCACTGTAACACACCTGCAGAGTGGCACACCTGTGCTATGGCACTTCTGCACACTGTGGCACACTTGTACCGTGGC
T **GC** A C T G T A A C A C A C C T G C A G A G T G G C A C A C C T G T G C T A T G G C A C T T C T G C A C T G T G G C A C A C T T G T A C C G T G G C
260 265 270 275 280 285 290 295 300 305 310 315 320 325 330



b



Supplementary Figure 3 Analysis of the off-target effect. The off-target sites that exactly match 12 bases at the 3' end and the NGG (blue box) were searched with free software CRISPR direct. The correct sequences around the off-target candidates against intron 3 (a) and 11 (b) of the *Creb3l3* genome were confirmed by sequencing.



Supplementary Figure 4 The analysis of integration of pX330 vector into mouse genome. The Cas9 cDNA were not observed in genome from CREB3L3 floxed mice by PCR.

	Fasted			Fed		
	flox	LKO	IKO	flox	LKO	IKO
BW (g)	22.83 ± 0.71	23.72 ± 0.39	21.95 ± 0.96	26.03 ± 0.92	26.69 ± 0.46	24.55 ± 1.16
Lean wight (g)	19 ± 0.69	19.62 ± 0.33	18.15 ± 0.87	21.9 ± 0.87	22.19 ± 0.43	20.2 ± 1.01
Fat weight (g)	3.83 ± 0.09	4.1 ± 0.16	3.73 ± 0.09	4.13 ± 0.12	4.49 ± 0.19	4.35 ± 0.15
Fat weight (%)	16.81 ± 0.56	17.29 ± 0.55	17.05 ± 0.36	15.94 ± 0.55	16.84 ± 0.66	17.675 ± 0.23*

Supplementary Table 1 Body composition of CREBH LKO and IKO mice

Body composition of 8-week-old male mice fasted for 24 h and fed was determined by DEXA. n = 7 ~ 10 / group, * p < 0.05.

	Fasted		Fed	
	flox	IKO	flox	IKO
Glu (mg/dl)	74.0 ± 3.8	72.6 ± 4.6	161.8 ± 9.3	136.5 ± 5.6*
TG (mg/dl)	232.4 ± 23.2	207.2 ± 15.6	101.5 ± 5.7	105.1 ± 2.9
Cholesterol (mg/dl)	53.8 ± 3.4	64.4 ± 13.6	52.0 ± 3.4	46.2 ± 4.4
FFA (mEq/l)	2.65 ± 0.15	2.39 ± 0.19	1.95 ± 0.13	2.09 ± 0.17
FGF21 (pg/ml)	774.7 ± 154.9	787.0 ± 258.2	88.3 ± 38.6	76.0 ± 27.0

Supplementary Table 2 Plasma parameters in CREBH IKO mice

Samples were collected from 8-week-old male mice fasted for 24 h and fed. n = 7 ~ 10 / group, * p < 0.05.

Gene	Fwd primer	Rv primer
<i>Acox1</i>	CGA TCC AGA CTT CCA ACA TGA G	CCA TGG TGG CAC TCT TCT TAA CA
<i>Acta2</i>	CCC AGA CAT CAG GGA GTA ATG G	TCT ATC GGA TAC TTC AGC GTC A
<i>Apoa4</i>	TTA CCC AGC TAA GCA ACA ATG C	GAG GGT ACT GAG CTG CTG AGT GA
<i>Apoa5</i>	GCG AGT TCT GCC GTA GGA C	CCC AAC CCC ATC AAA TGT GA
<i>Apoc2</i>	CCA AGG AGC TTG CCA AAG AC	TGC CTG CGT AAG TGC TCA TG
<i>Apoc3</i>	TAG AGG GCT ACA TGG AAC AAG C	CAG GGA TCT GAA GTG ATT GTC C
<i>Ccl2</i>	CCC ACT CAC CTG CTG CTA CT	ATT TGG TTC CGA TCC AGG TT
<i>Cd68</i>	CCT CCA CCC TCG CCT AGT C	TTG GGT ATA GGA TTC GGA TTT GA
<i>Col1a1</i>	GCT CCT CTT AGG GGC CAC T	ATT GGG GAC CCT TAG GCC AT
<i>Cpt1a</i>	TCT TTC ACT GAG TTC CGA TGG G	ACG CCA GAG ATG CCT TTT CC
<i>Creb3l3</i>	CCA GAC CCT TTA CCC ATA CAT	ATG GTT GGA AGT TAG GGT TCA G
<i>Cyp7a1</i>	GCT GAG AGC TTG AAG CAC AAG A	TTG AGA TGC CCA GAG GAT CAC
<i>Cyclophilin</i>	TGG CTC ACA GTT CTT CAT AAC CA	ATG ACA TCC TTC AGT GGC TTG TC
<i>Elovl2</i>	ACG CTG GTC ATC CTG TTC TT	GCC ACA ATT AAG TGG GCT TT
<i>Elovl5</i>	ATG GAA CAT TTC GAT GCG TCA	GTC CCA GCC ATA CAA TGA GTA AG
<i>Elovl6</i>	ACA ATG GAC CTG TCA GCA AA	GTA CCA GTG CAG GAA GAT CAG T
<i>F4/80</i>	CTT TGG CTA TGG GCT TCC AGT C	GCA AGG AGG ACA GAG TTT ATC GTG
<i>Fasn</i>	ATC CTG GAA CGA CGA GAA CAC GAT CT	AGA GAC GTG TCA TCC TGG ACT T
<i>Fgf21</i>	AGA TCA GGG AGG ATG GAA CA	TCA AAG TGA GGC GAT CCA TA
<i>Fxr</i>	CTC TGC TCA CAG CGA TCG TC	CAC CGC CTC TCT GTC CTT GA
<i>Hmgcr</i>	GAG AAG AAG CCT GCT GCA TA	CGT CAA CCA TAG CTT CCG TAG TT
<i>Hmgcs1</i>	GTG GCA CCG GAT GTC TTT G	CTC TGA CCA GAT ACC ACG TTC
<i>Il1b</i>	GCA ACT GTT CCT GAA CTC AAC T	ATC TTT TGG GGT CCG TCA ACT
<i>Il6</i>	TAG TCC TTC CTA CCC CAA TTT CC	TTG GTC CTT AGC CAC TCC TTC
<i>Ldlr</i>	TGG AGG ATG AGA ACC GGC T	GCA CTG AAA ATG GCT TCG TTT A
<i>Ppara</i>	CCT CAG GGT ACC ACT ACG GAG T	GCC GAA TAG TTC GCC GAA
<i>Scd1</i>	AGA TCT CCA GTT CTT ACA CGA CCA C	CTT TCA TTT CAG GAC GGA TGT CT
<i>Shp</i>	CAA GGA GTA TGC GTA CCT GAA G	CCT GGC ACA TCT GGG TTG AAG
<i>Srebf1a</i>	AGG CGG CTC TGG AAC AGA	TCA AAA CCG CTG TGT CCA GTT
<i>Srebf1c</i>	CGG CGC GGA AGC TGT	TGC AAT CCA TGG CTC CGT
<i>Srebf2</i>	CTG CAG CCT CAA GTG CAA AG	CAG TGT GCC ATT GGC TGT CT
<i>Tgfb1</i>	CCA CCT GCA AGA CCA TCG AC	CTG GCG AGC CTT AGT TTG GAC
<i>Tnfa</i>	TCG TAG CAA ACC ACC AAG TG	AGA TAG CAA ATC GGC TGA CG

Supplementary Table 3 QPCR primer sequences