

Figure S1

Protein sequence

ORIGIN

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1 mafmkkyllp ilglfmayyy ysaneefrpe mlqgkkvi tv gaskgigrem ayhlakmgah
61 vvv tarsket lqkvvshcle lgaasahyia gtmedmtfae qfvaqagklm ggldmlilnh
121 itntslnlfh ddihhvrksm evnflsyvvl tvaalpmlkq sngsivvvss lagkvaypmv
181 aaysaskfal dgffssirke ysvsrvnvsi tlcvlglidt etamkavsgi vhmqaapkee
241 caleiikgga lrqeevyyds slwttllirn pcrkilefly stsynmdr fi nk
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HSD11B1 DNA sequence

ORIGIN

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1 acaattcaga ggctgctgcc tgcttaggag gttgtagaaa gctctgtagg ttctctctgt
61 gtgtcctaca ggagtcttca ggccagctcc ctgtcggatg gcttttatga aaaaatatct
121 cctccccatt ctggggctct tcatggccta ctactactat tctgcaaacg aggaattcag
181 accagagatg ctccaaggaa agaaagtgat tgtcacaggg gccagcaaa ggcgcggaag
241 agagatggct tatcatctgg cgaagatggg agcccatgtg gtggtgacag cgaggtcaaa
301 agaaactcta cagaagggtg tatcccactg cctggagcct ggagcagcct cagcacacta
361 cattgctggc accatggaag acatgacctt cgcagagcaa tttgttgccc aagcaggaaa
421 gctcatggga ggactagaca tgctcattct caaccacatc accaactctt ctttgaatct
481 ttttcatgat gatattcacc atgtgcgcaa aagcatggaa gtcaacttcc tcagttacgt
541 ggtcctgact gtagctgcct tgcccattgt gaagcagagc aatggaagca ttgttgctgt
601 ctctctctct gctgggaaa gctggttatcc aatggttgct gcctattctg caagcaagtt
661 tgctttggat ggggttcttct cctccatcag aaaggaatat tcagtgtcca ggggtcaatgt
721 atcaatcact ctctgtgttc ttggcctcat agacacagaa acagccatga aggcagtttc
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1201 taaacatagg tataattacc agatagttat attaaattta tatcttatat ataataatat
1261 gtgatgatta atacaatatt aattataata aaggtcacat aaactttata aattcataac
1321 tggtagctat aacttgagct tattcaggat ggtttcttta aaaccataaa ctgtacaaat
1381 gaaa
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Fusion Protein design

NheI Kozak

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5' GCTAGCGCCACCATGGCGAGC ATGGATAGCACTGAGAACGTCATCAAGCCCTTCATGCGCTTCAAGGT
GCACATGGAGGGCTCCGTGAACGGCCACGAGTTCGAGATCGAGGGCGTGGGCGAGGGCAAGCCCTACGAG
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CAGAGCAATGGAAGCATTGTTGTCGTCTCCTCTCTGGCTGGGAAAGTGGCTTATCCAATGGTTGCTGCCT
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CAATGTATCAATCACTCTCTGTGTTCTTGGCCTCATAGACACAGAAACAGCCATGAAGGCAGTTTCTGGG
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AAGAAGAAGTGTATTATGACAGCTCACTCTGGACCACTCTTCTGATCAGAAATCCATGCAGGAAGATCCT
GGAATTCTCTACTCAACGAGCTATAATATGGACAGATTCATAACAAG**TAG**CGGGCCGC**3'**

Not1

Table S1. Antibodies used in Western blot studies.

Antibody	Host	Source/Catalog	WB dilution
Acetyl H4K12	Sheep	R&D Systems/AF5215	1:1000
H4	Rabbit	Cell Signaling/#2592	1:1000
Acetyl H3K9	Rabbit	Millipore/06-599	1:1000
H3	Rabbit	Cell Signaling/#9715	1:1000
H3K9me2	Rabbit	Cell Signaling/#4658	1:1000
NF-κB	Rabbit	Cell Signaling/DE14E12	1:1000
NRF2	Rabbit	Santa Cruz/sc-722	1:500
Beclin 1	Rabbit	Abcam/ab62557	1:1000
p-TORC1 S151	Rabbit	Cell Signaling/#3359	1:1000
TORC1	Rabbit	Cell Signaling/#2501	1:1000
LC3B	Rabbit	Cell Signaling/#2775	1:1000
p-CREB S133	Rabbit	Cell Signaling/#9198	1:1000
CREB	Rabbit	Cell Signaling/#4820	1:1000
BDNF	Rabbit	Santa Cruz/N-20	1:500
PSD95	Rabbit	Abcam/ab12093	1:2000
Synaptophysin	Mouse	Millipore/MAB5258	1:1000
SNAP25	Mouse	Santa Cruz/ SP-12	1:500
GAPDH	Mouse	Millipore/MAB374	1:2000
β-Tubulin	Mouse	Millipore/Clone AA2	1:2000
TBP	Mouse	Abcam/ab818	1:2000
Goat-anti-mouse HRP conjugated		Biorad/170-5047	1:2000
Goat-anti-rabbit HRP conjugated		Biorad/170-6515	1:2000
Donkey-anti-goat HRP conjugated		Santa Cruz/sc-2020	1:2000

Table S2. Primers and probes used in qPCR studies.

SYBR-Green primers

Target	Product size (bp)	Forward primer (5'-3')	Reverse primer (5'-3')
<i>Tet2</i>	113	CCATCATGTTGTGGGACGGA	ATTCTGAGAACAGCGACGGT
<i>Hdac2</i>	280	CTATCCCGCTCTGTGCCCT	GAGGCTTCATGGGATGACCC
<i>Il-18</i>	179	ACAGAATATCAACCAACAAGTTG ATATTCTC	GATTCTTTCCTTTGAGGCCCA
<i>Cxcl2</i>	100	AGCCACACTTCAGCCTAGCG	TGTAGCCTGGTGGTTGGTGG
<i>Tnf-α</i>	157	TCGGGGTGATCGGTCCCCAA	TGGTTTGCTACGACGTGGGCT
<i>Gfap</i>	125	CCTTCTGACACGGATTTGGT	ACATCGAGATCGCCACCTAC
<i>Aox1</i>	286	CATAGGCGGCCAGGAACATT	TCCTCGTTCAGAAATGCAGC
<i>iNOS</i>	101	GGCAGCCTGTGAGACCTTTG	GAAGCGTTTCGGGATCTGAA
<i>Adam10</i>	125	CCTTCTGACACGGATTTGGT	CTGTACAGCAGGGTCCTTGAC
<i>Bace1</i>	67	ACAAGCCTTTCGCCTCC	TCAGGCCACCATAATCCAGC
<i>Aβ-precursor</i>	99	TCGGGGTGATCGGTCCCCAA	GTCACGTTCCACCTCCCCAG
<i>Neprilisin12</i>	196	TTGGGAGACCTGGCGGAAAC	CATTCTTGACCCTCACCCC
<i>Creb</i>	86	GGCTGCTGCTGCCTGT	ACACACCGCGTCAAACCTACA
<i>β-actin</i>	218	CTGTCCCTGTATGCCTCTG	ATGTCACGCACGATTTCC

Taqman probes

Target	Product size (bp)	Reference
<i>Dnmt1</i>	58	Mm01151063_m1
<i>Gapdh</i>	107	Mm99999915_g1