Protein target	Company (catalog number)	Specie	Dilution
FFAR1 β -Actin UCP1 UCP3 PGC1 α PPAR γ DIO2 α -Tubulin p-IKK α/β (Ser- 180) p- SAPK/JNK (Thr183/Tyr185) GPR78 p-IRE1 (Ser724) p-PERK (Thr980) p-eiF2a (Ser 52) ATF-6 α (F-7) CHOP p-STAT3 (Y705) SOCS3 p-Altr (Ser 472)	Sigma-Aldrich (SAB2108661) Sigma-Aldrich (A5316) Abcam (ab10983) Santa Cruz (sc-33182) Santa Cruz (sc-17380) Santa Cruz (sc-17380) Santa Cruz (sc-7273) Abcam (ab ab77779) Sigma-Aldrich (T5168) Santa Cruz (sc-23470) Cell Signaling (#4671) Cell Signaling (#4671) Cell Signaling (#4671) Cell Signaling (#3183) Abcam (ab48187) Cell Signaling (#3179) Santa Cruz (sc-101670) Santa Cruz (sc-101670) Santa Cruz (sc-166659) Cell Signaling (#5554) Abcam (ab76315) Cell Signaling (#23064)	rabbit mouse rabbit mouse mouse rabbit mouse rabbit rabbit rabbit rabbit rabbit rabbit rabbit rabbit rabbit	1:5000 1:5000 1:5000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000 1:1000
AKT	Cell Signaling (#9272)	rabbit	1:1000

Supplementary Table 1. Antibodies used for western blot

Primers	ThermoFisher TaqManGene Expression Assays ID
Ucp1	Mm01244861_m1
Pparg	Mm01184322_m1
Ppargc1a	Mm01208835_m1
Prdm16	Mm01266512_m1
Hprt	Mm01545399_m1

Supplementary Table 2. Primers used for RT-PCR



Supplementary Figure 1. Effect of central TUG-905 administration on energy balance and lipid metabolism

(A) Absolute body weight change during the 7 days experimental period (n=20-24 mice/group), (B) representative microphotographs of Oil-Red staining (10X, scale bar: 200 μ m), (C) hepatic lipid content in Oil Red O-stained sections (n=8-9 mice/group), circulating (D) triglycerides, (E) NEFA and (F) cholesterol (n=6-7 mice/group) of vehicle and TUG-905 ICV-treated DIO mice. Data are expressed as MEAN±SEM. Statistical significance was determined by two-way ANOVA (A) or Student's t-test (C, D, E, and F); **P<0.01 vs. vehicle.



Supplementary Figure 2. Effect of POMC-specific Ffar1 knockdown on energy balance

(A) Negative control probes for GFP and FFAR1, (B) number of POMC⁺/GFP⁺, POMC⁺/GFP⁻, POMC⁺/GFP⁻ cells (n= 3-4 mice/group) and (C) absolute body weight change (n= 11 mice/group) of POMC-Cre mice treated with AAV encoding shScramble or shFfar1. Data are expressed as MEAN±SEM. Statistical significance was determined by two-way ANOVA (C) or Student's t-test (B). *P < 0.05 vs. shScramble.