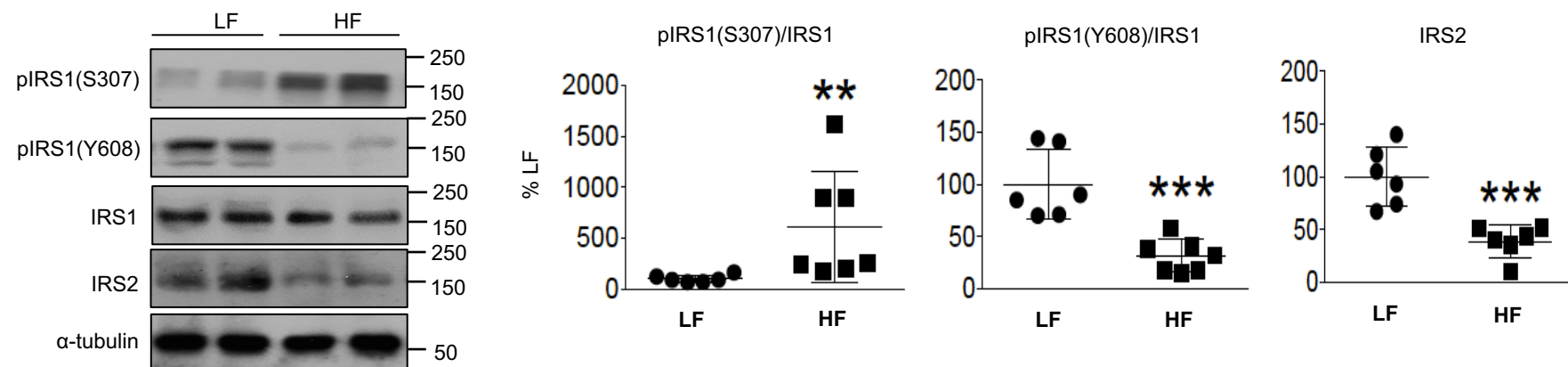
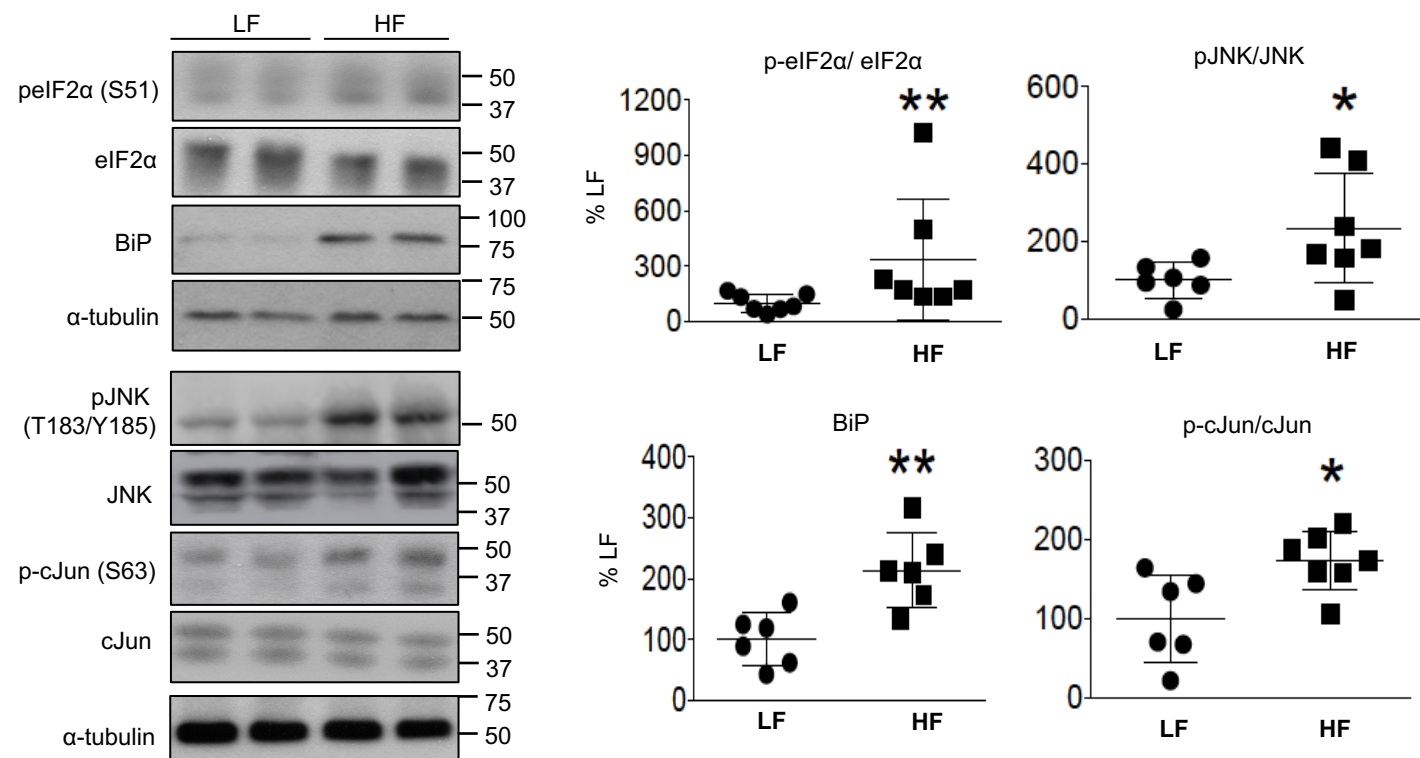
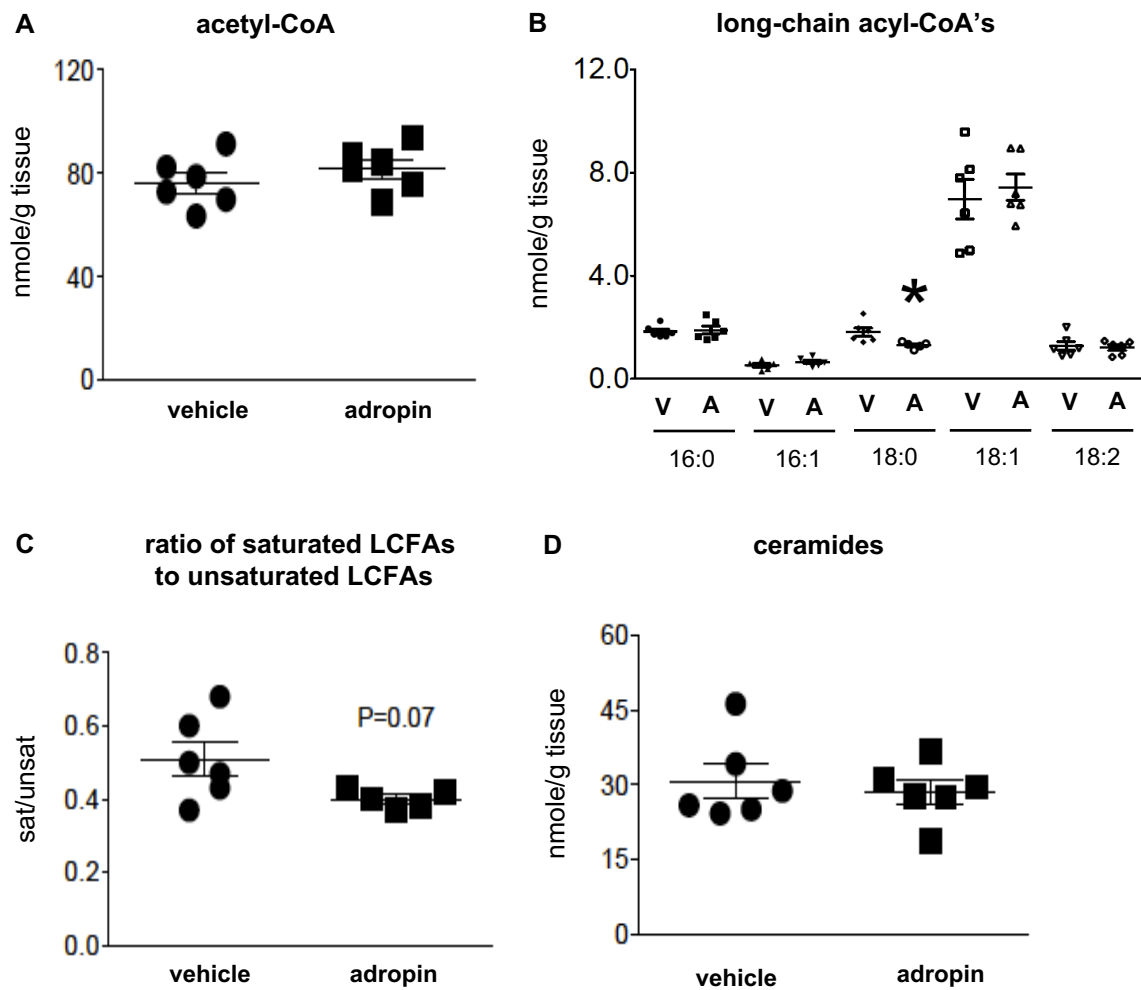
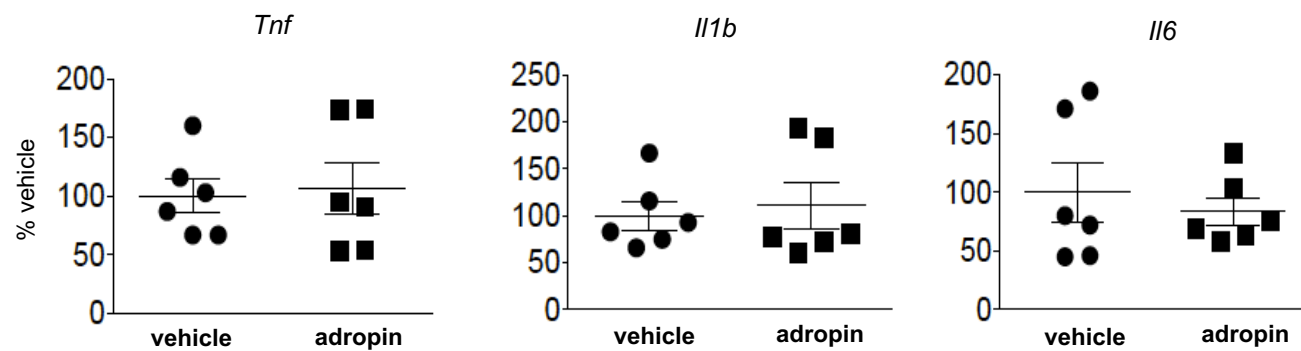


Supp. Fig. 1

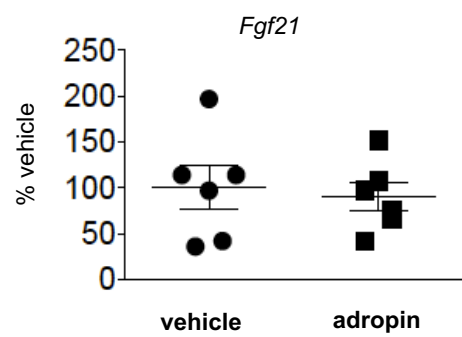
A**B**



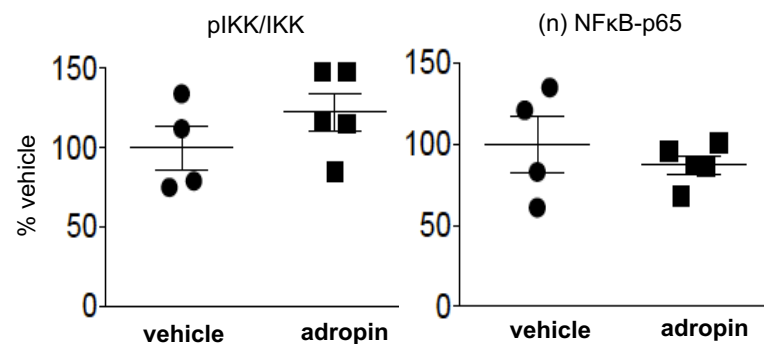
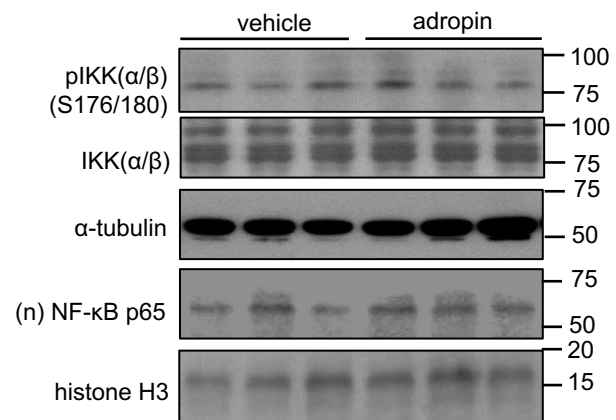
Supp. Fig. 3



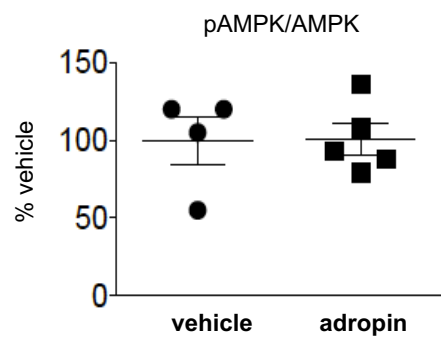
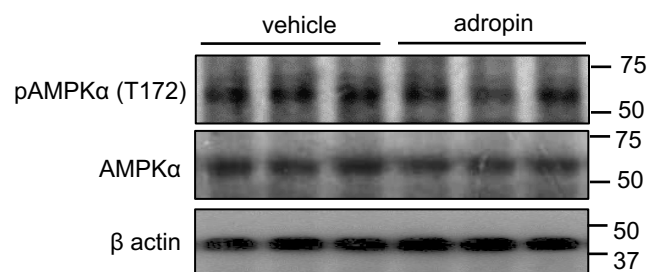
Supp. Fig. 4



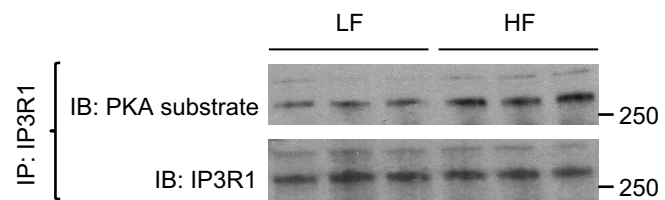
Supp. Fig. 5



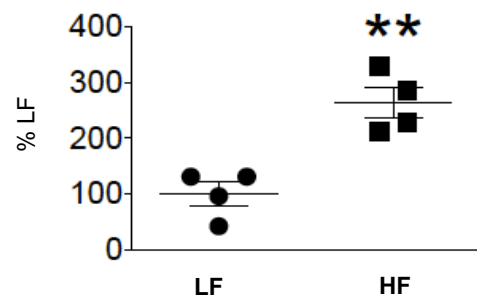
Supp. Fig. 6



Supp. Fig. 7



PKA phosphorylation of IP3R1



Supp. Fig. 8

Supplementary figure legends

S1 Adropin³⁴⁻⁷⁶ treatment reduced blood glucose level in 6h-fasted DIO mice.

Diet-induced obese (DIO) mice maintained on high fat diet (HF) and lean mice (control) maintained on low fat diet (LF) received 5 injections of adropin³⁴⁻⁷⁶ or vehicle. HF/veh: n=6, HF/adr: n=6, LF/veh: n=8. Six-hour fasting period encompassed 5h before and 1h after the last injection. One hour after the last injection, tail blood glucose levels were determined.

S2 Diet-induced obese mice displayed altered IRS signaling, increased JNK activity and ER stress responses in the liver.

DIO mice were maintained on high fat diet (HF), and lean mice maintained on low fat diet (LF) were used as the control. (A) The phosphorylation levels of Ser-307 in IRS1 (n=6-7) and Tyr 608 in IRS1 (n=6-7) as well as total IRS2 protein levels (n=6) were determined by Western blotting. HF vs LF: **, P<0.01; ***, P<0.0005. (B) The phosphorylation levels of Ser51 in eIF2 α (n=7), BiP protein levels (n=6), the phosphorylation levels of Thr-183/Tyr-185 in JNK (n=6-7) and the phosphorylation levels of Ser63 in c-Jun (n=6-7) were determined by Western blotting. α -tubulin was used as the loading control.

S3 Adropin³⁴⁻⁷⁶ treatment did not exert major effects on the levels of fatty acid metabolites in the liver.

Lipidomics was performed to quantify the levels of acetyl-CoA (A, n=6), individual long-chain acyl-CoA's (B, n=5-6), the ratio of saturated fatty acyl-CoA levels (16:0 plus 18:0)

to unsaturated fatty acyl-CoA levels (16:1 plus 18:1) (C, n=5-6) and ceramides (D, n=6).
adropin vs vehicle: *, $P \leq 0.05$.

S4 Adropin³⁴⁻⁷⁶ treatment did not alter the expressions of inflammatory cytokines in the liver.

The message levels of inflammatory cytokines including tumor necrosis factor (*Tnf*), interleukin-1b (*Il1b*) and interleukin-6 (*Il6*) were determined by real-time RT-PCR (n=6).

S5 Adropin³⁴⁻⁷⁶ treatment did not alter the expression of FGF21 in the liver.

The message levels of fibroblast growth factor 21(*Fgf21*) was determined by real-time RT-PCR (n=6).

S6 Adropin³⁴⁻⁷⁶ treatment did not affect IKK pathway in the liver.

The phosphorylation levels of Ser-176/180 in inhibitor- κ B kinase α/β (IKK α/β) in whole tissue lysates (n=4-5) and the levels of nuclear factor κ B p65 subunit (N- κ B p65) in the nuclear extract (n=4-5) were determined by Western blotting. α -tubulin and histone H3 were used as the loading control in whole tissue lysates and nuclear lysates, respectively. The same α -tubulin band was used as the loading control for the blot of total IRS1 (Fig. 1A) and the blots of pAkt (S473) and total Akt (Fig. 2A). The same histone H3 band was used as the loading control for the blots of (n)FoxO1 (Fig. 2D), (n)SREBP1c (Fig. 6A) and (n)CRTC2 (Fig. 8B).

S7 Adropin³⁴⁻⁷⁶ treatment did not affect the phosphorylation level of AMPK in the liver.

The phosphorylation levels of Thr-172 in AMP-activated protein kinase (AMPK) were determined by Western blotting (n=4-5). β -actin was used as the loading control.

S8 DIO mice displayed increased phosphorylation level of PKA substrate sites in IP3R in the liver.

The phosphorylation levels of PKA substrate sites in IP3R1 in DIO mice maintained on HF and lean mice on LF were measured by Western blotting following immunoprecipitation of IP3R1 (n=4). HF vs LF: **, $P < 0.01$.