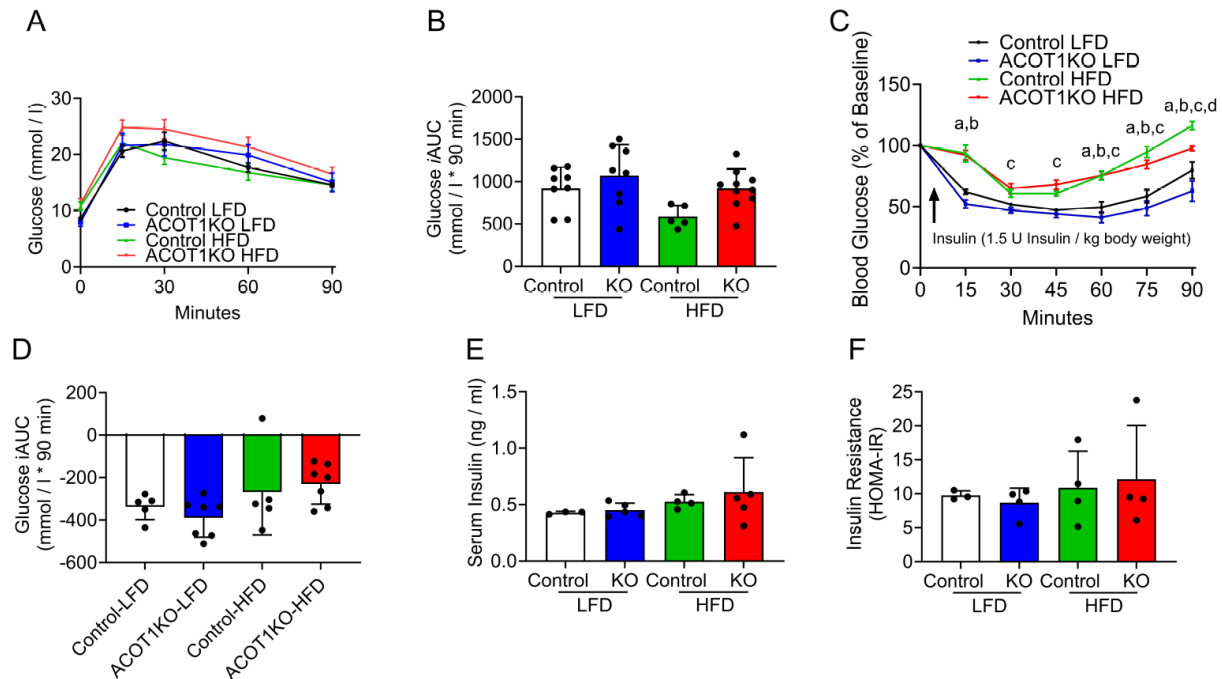
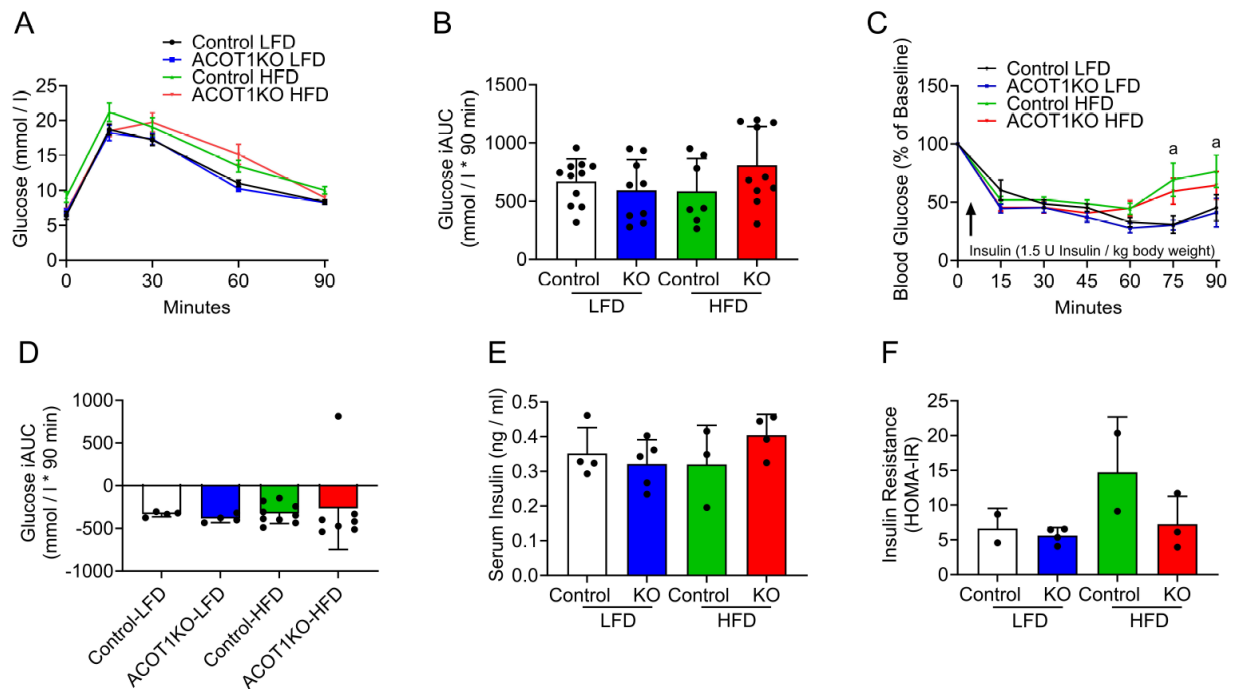


**Supplemental Figure 1. Protein densitometry and female ACOT1KO mice.** (A) Protein densitometry quantification from Figure 1D ( $n = 2$  per group). (B) Body weight changes in female mice ( $n = 12-16$  per group). (C) Body weight gain (calculated as the difference between final body weight and initial body weight in 20-week-old female mice given the LFD or HFD for 12 weeks ( $n = 11-16$  per group)). (D) Relative tissue weight of various female tissues ( $n = 1-12$  per group). All data are presented as means  $\pm$  S.E.M. <sup>a</sup> $P \leq 0.01$  control LFD vs. control HFD, <sup>b</sup> $P = 0.02$  ACOT1KO LFD vs. ACOT1KO HFD, <sup>c</sup> $P \leq 0.01$  control HFD vs. ACOT1KO HFD. ANOVAs were used for statistical tests.

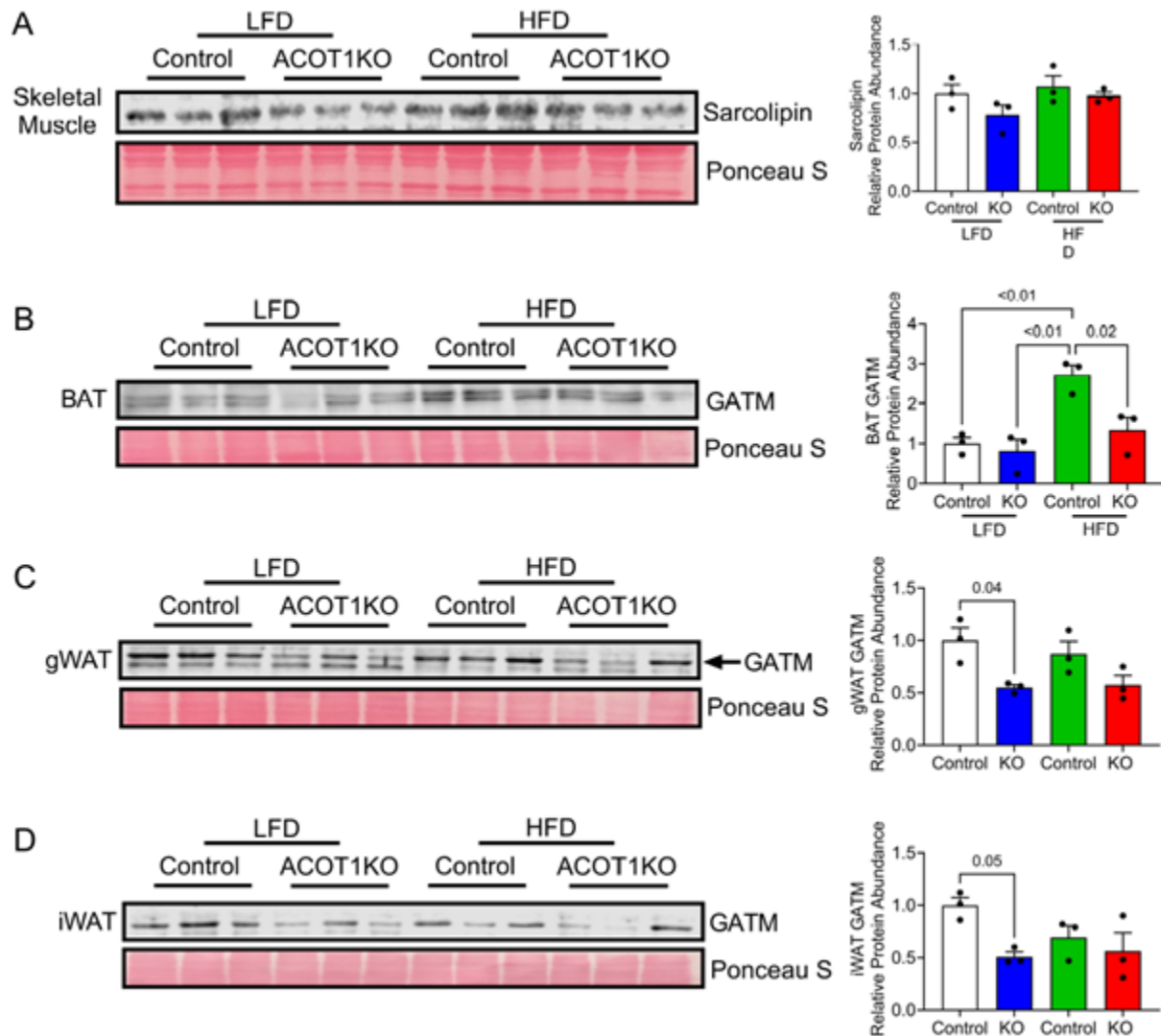


**Supplemental Figure 2. No impact of ACOT1 deficiency on glucose metabolism or insulin**

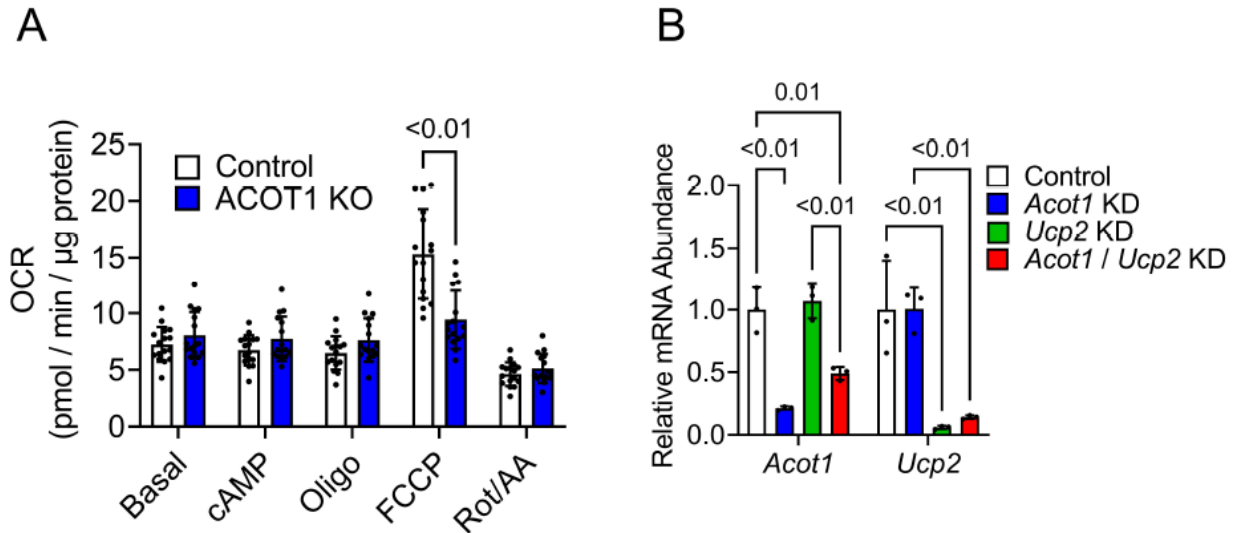
**resistance in male mice.** (A) Glucose time course during an oral glucose tolerance test in 20-week-old male mice given the LFD or HFD for 12 weeks. (n = 5-10 per group). (B) Glucose incremental area under the curve (n = 5-10 per group). (C) Glucose time course during an insulin tolerance test (n = 4-7 per group). (D) Glucose incremental area under the curve during insulin tolerance test (n = 5-7). (E) Fasting serum insulin levels (n = 3-5 per group). (F) Insulin resistance was calculated using the Homeostatic Model for Insulin Resistance (HOMA-IR) formula (n = 3-4 per group). All data are presented as means  $\pm$  S.E.M. <sup>a</sup>P  $\leq$  0.01 control LFD vs. control HFD, <sup>b</sup>P = 0.02 ACOT1KO LFD vs. ACOT1KO HFD, <sup>c</sup>P  $\leq$  0.01 ACOT1KO LFD vs. ACOT1KO HFD, <sup>d</sup>P = 0.02 control HFD vs. ACOT1KO HFD. ANOVAs were used for statistical tests.



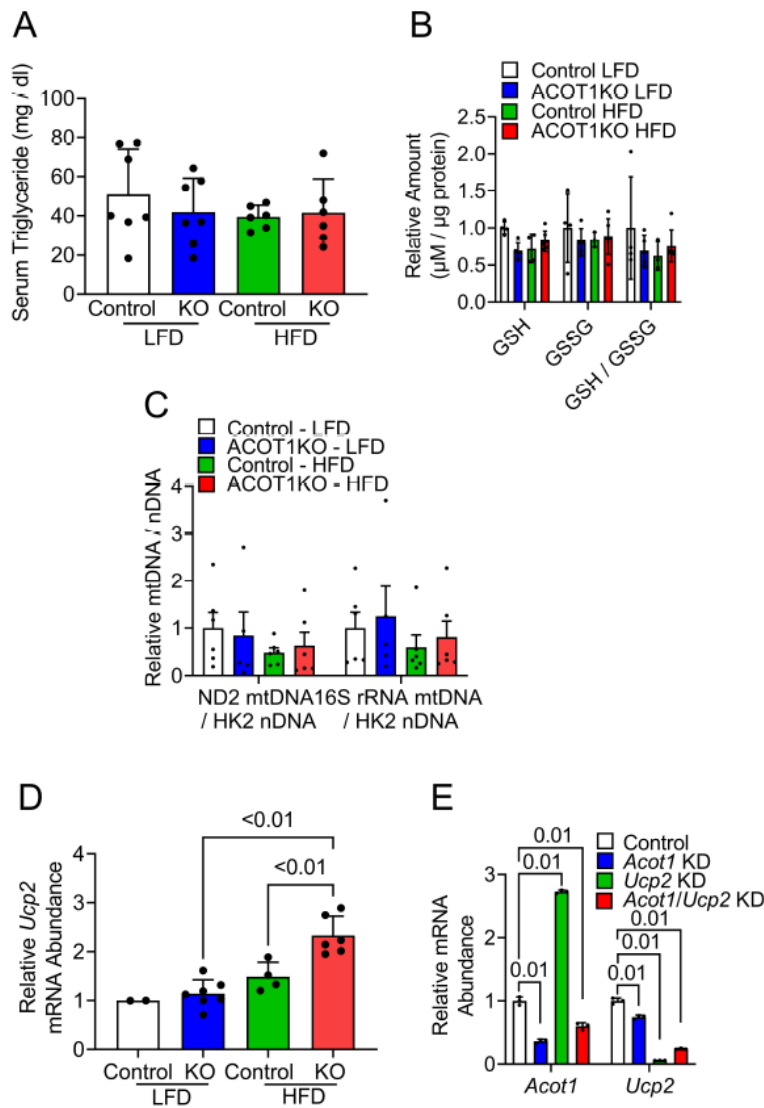
**Supplemental Figure 3. Glucose tolerance and insulin sensitivity in female mice.** (A) Glucose time course during an oral glucose tolerance test in 20-week-old female mice given the LFD or HFD for 12 weeks. (n = 7-11 per group). (B) Glucose incremental area under the curve (n = 7-11 per group). (C) Glucose time course during an insulin tolerance test (n = 4-9 per group). (D) Glucose incremental area under the curve during insulin tolerance test (n = 4-9). (E) Fasting serum insulin levels (n = 3-5 per group). (F) Insulin resistance was calculated using the Homeostatic Model for Insulin Resistance (HOMA-IR) formula (n = 2-4 per group). All data are presented as means  $\pm$  S.E.M. <sup>a</sup>P = 0.04 control LFD vs control HFD. ANOVAs were used for statistical tests.



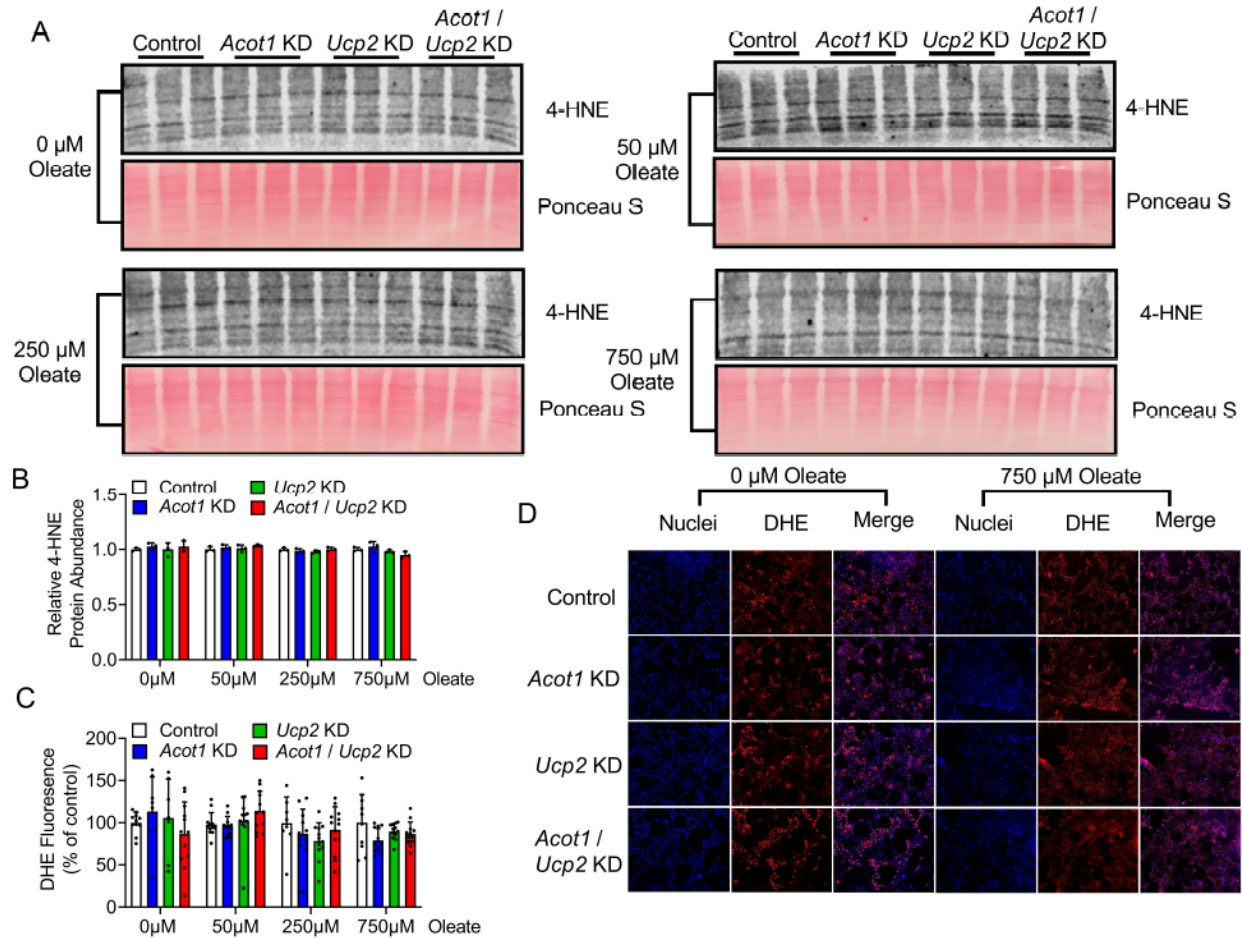
**Supplemental Figure 4. Abundance of proteins involved in UCP1 independent mechanisms of thermogenesis.** (A) Western blot image (left) and densitometry quantification (right) of sarcolipin protein abundance in skeletal muscle from 20-week-old control or ACOT1KO mice fed the LFD or HFD for 12 weeks ( $n = 3$  per group). (B-D) Western blot image (left) and densitometry quantification (right) of arginine:glycine amidinotransferase (GATM) protein abundance in BAT (B), gWAT (C), and iWAT (D) (all  $n = 3$  per group). All data are presented as means  $\pm$  S.E.M. ANOVAs were used for statistical tests.



**Supplemental Figure 5. Oxygen consumption rates and mRNA abundance in adipocytes.** (A) OCR in stromal vascular cells was measured with a Seahorse under basal conditions and after sequential injections of 8-bromo cyclic adenosine monophosphate (cAMP), oligomycin (oligo), carbonyl cyanide-p-trifluoromethoxyphenyl hydrazone (FCCP), and rotenone/antimycin A (Rot/AA) (n = 16 per group). (B) Relative mRNA abundance of *Acot1* or *Ucp2* in 3T3-L1 cells treated with control (scrambled), sh*Acot1*, sh*Ucp2*, or both sh*Acot1* / sh*Ucp2* lentivirus to induce gene knockdown (KD) (n = 3 per group). All data are presented as means  $\pm$  S.E.M. ANOVAs were used for statistical tests.



**Supplemental Figure 6. ACOT1 deficiency and hepatic metabolism.** (A) Serum triglyceride levels in 20-week-old mice fed the LFD or HFD for 12 weeks (n = 6-7 per group). (B) Relative hepatic glutathione (GSH), oxidized glutathione (GSSG), and the GSH / GSSG ratio (n = 3-6 per group). (C) Relative hepatic mtDNA content (n = 5-6 per group). (D) Relative mRNA abundance of hepatic *Ucp2* (n = 2-7 per group). (E) Relative mRNA abundance of *Acot1* or *Ucp2* in AML12 cells treated with control (scrambled), sh*Acot1*, sh*Ucp2*, or both sh*Acot1* / sh*Ucp2* lentivirus to induce gene knockdown (KD) (n = 3 per group). All data are presented as means ± S.E.M. ANOVAs were used for statistical tests.



**Supplemental Figure 7. Oxidative stress in AML12 cells treated with various doses of oleate for 24 hours.** (A-B) Western blot images (A) and densitometry quantification (B) of 4-HNE protein abundance in AML12 cells treated with control (scrambled), sh*Acot1*, sh*Ucp2*, or both sh*Acot1* / sh*Ucp2* lentivirus to induce gene knockdown (KD) and after a 24 h incubation with the indicated amount of oleate (n = 3 per group, per condition). (C-D) Dihydroethidium (DHE) fluorescence (n = 7-12 per group) (C) and representative images (D) of select conditions of AML12 cells treated with the indicated lentivirus and pre-treated with oleate (10X objective). A total of 3-4 images per condition were taken. All data are presented as means  $\pm$  S.E.M. ANOVAs were used for statistical tests.