

An *Adcy3* coding mutation causes partial loss of enzymatic function, contributing to obesity in a rat model by reducing lipolysis

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Supplemental Figures

Figure S1. Body weight curves and tissue weights at sac in Adcy3^{mut/mut} vs WT rats. (a)

Adcy3^{mut/mut} males (M) and females (F) gain more weight than wild-type (WT) rats during the study. **(b)** Adcy3^{mut/mut} M and F have more retroperitoneal fat (RetroFat), gonadal fat, and omental fat (OmenFat) than WT rats at necropsy. **(c)** Adcy3^{mut/mut} M both have larger pancreas at necropsy than WT M with no significant differences in the necropsy weights of the other non-adipose tissues. Mean \pm SEM. T-test or rmANOVA, *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001

Figure S2. *Ex vivo* cAMP production in response to stimulation with a β 3-AR agonist in

RetroFat of Adcy3^{mut/mut} vs WT rats, separated by diet. (a) cAMP production in response to β 3-AR agonist in RetroFat is blunted in males (M) on high fat diet (HFD) for nine weeks relative to those on standard diet (SD) (p=0.042) **(b)** There is not a significant effect of diet in females (F). Mean \pm SEM. rmANOVA

Figure S3. Body weight curves, fat and lean mass, tissue weights at sac, fasting serum FFA

and body temperature during acute cold exposure in Adcy3^{+/-} vs WT rats. (a) Adcy3^{+/-} males (M) and females (F) gain more weight than wild-type (WT) rats during the study. **(b)** Adcy3^{+/-} M and F have increased fat mass with no differences in lean mass relative to WT rats. **(c)** Adcy3^{+/-} M and F have more retroperitoneal fat (RetroFat), gonadal fat, and omental fat (OmenFat) than WT rats at necropsy. **(d)** Adcy3^{+/-} M have larger pancreas at necropsy than wild-type (WT) M with no significant differences in the necropsy weights of any other non-adipose tissues. **(e)** During a 48-hour fast, Adcy3^{+/-} M and F have less serum FFA than WT rats as shown by a significant genotype by time interaction in the two-way ANOVA. **(f)** During an acute 5-hour cold

exposure at 4°C, there were no significant differences in body temperature between Adcy3^{+/} M and WT M. Adcy3^{+/} F maintain a significantly lower body temperature during acute cold exposure than WT F. Mean \pm SEM. T-test or rmANOVA, *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001

Figure S4. Gene expression changes in retroperitoneal fat (RetroFat), subcutaneous white adipose tissue (SubQ), and brown adipose tissue (BAT) in Adcy3^{+/} vs WT rats. **(a)** After a prolonged 48-hour cold exposure at 4°C, Adcy3^{+/} males (M) and females (F) generally have lower lipase gene expression (*Lipase E (Lipe)* and *Patatin-like phospholipase domain containing 2 (Pnpla2)*) in retroperitoneal adipose (RetroFat) and subcutaneous adipose (SubQ) than wild-type (WT) rats, as shown by both three-way ANOVA (results on legend) and stratified analyses (results in graph). There are no genotypic differences in general lipase gene expression in brown adipose tissue (BAT) after prolonged cold exposure, except that Adcy3^{+/} M express less *Pnpla2* than WT rats. **(b)** After a prolonged 48-hour cold exposure at 4°C, there were no significant differences in *Uncoupling protein 1 (Ucp1)* expression in Adcy3^{+/} M or F in BAT or SubQ. **(c)** After standard necropsy (no cold exposure), there were no significant differences in *Peroxisome proliferator-activated receptor gamma (Pparg)* expression in RetroFat between genotypes in males, while Adcy3^{+/} F express less *Pparg* than WT F. There were also no significant differences in *CCAAT/enhancer-binding protein alpha (Cebpa)* expression in RetroFat between genotypes in either sex. Mean \pm SEM. T-test, *p<0.05, ***p<0.001

Table S1. Primer sequences used in rt-qPCR.

Primer	Forward Sequence	Reverse Sequence
β -actin	TGAGGTAGTCTGTCAGGTCCCG	ACCACTGGCATTGTGATGGACT
<i>Ucp1</i>	GCAGAACACAGGTCTGGATG	ACGATCGTGCCTCTTCACTG
<i>Lipe</i>	AACTCACCCCTGACCGTGAAC	CTTGTTCGCCTGTGGGTAT
<i>Pnpla2</i>	GCACACAAACGTTCTCCTCA	GGGTGAACCTCCAGGTACCAA
<i>Pparg</i>	AGGGCTGATCAACTTCTCCA	GCAGGGGAATGTCTGTCT
<i>Cebpa</i>	ATTCTGGTGTGCGCTGAAAG	CCCCCTACTCGGTAGGAAAA

Figure S1

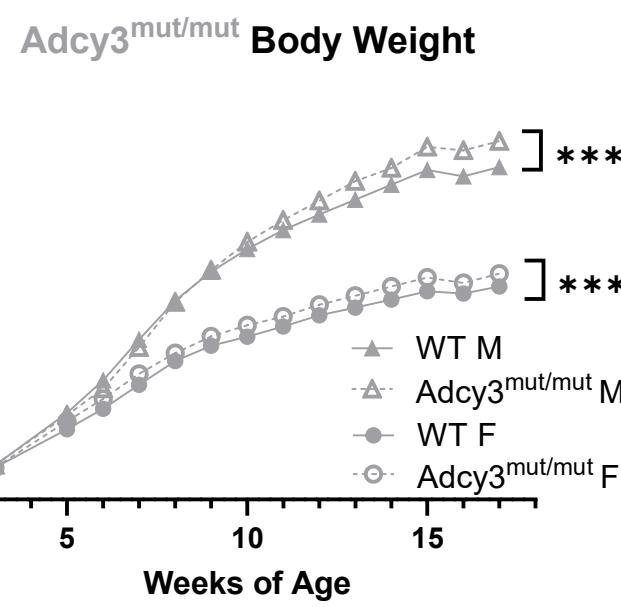
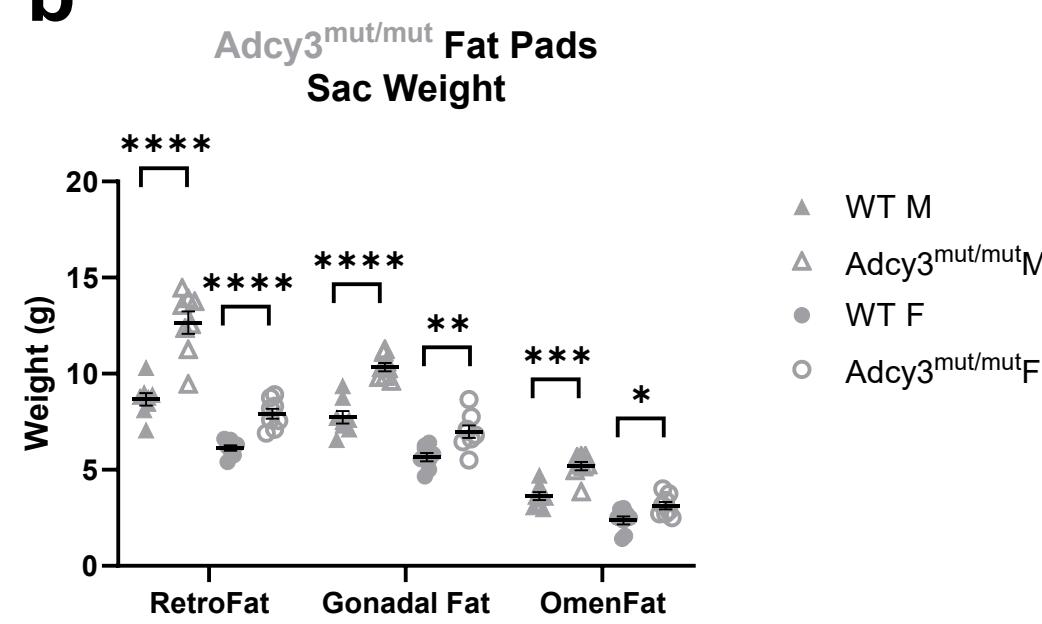
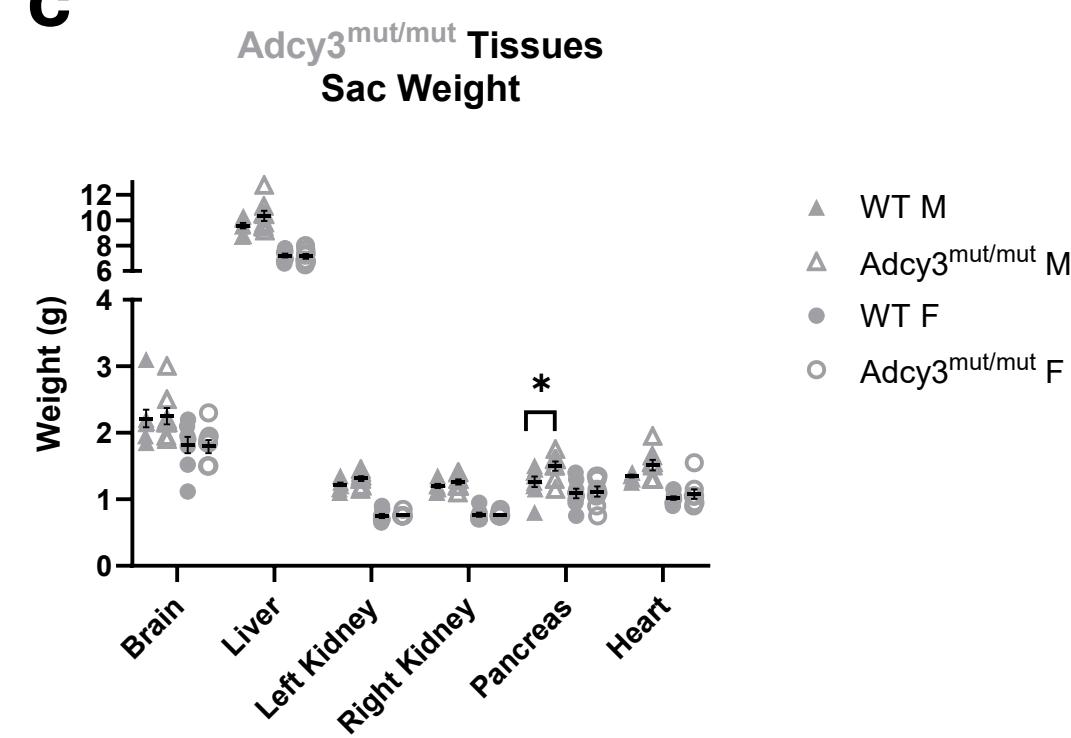
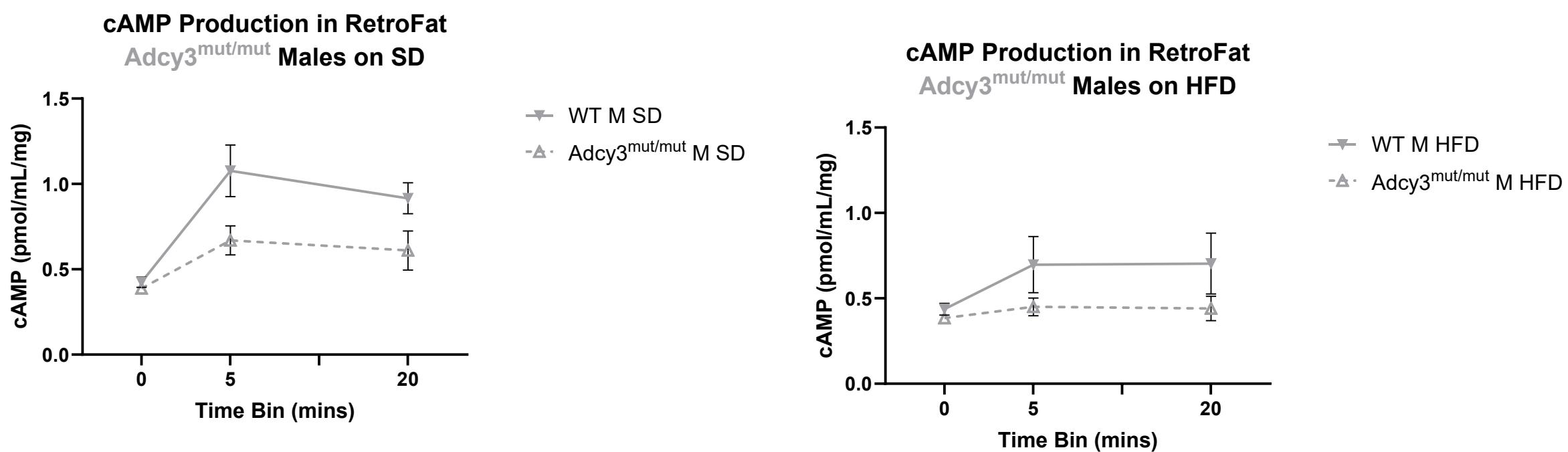
a**b****c**

Figure S2

a



b

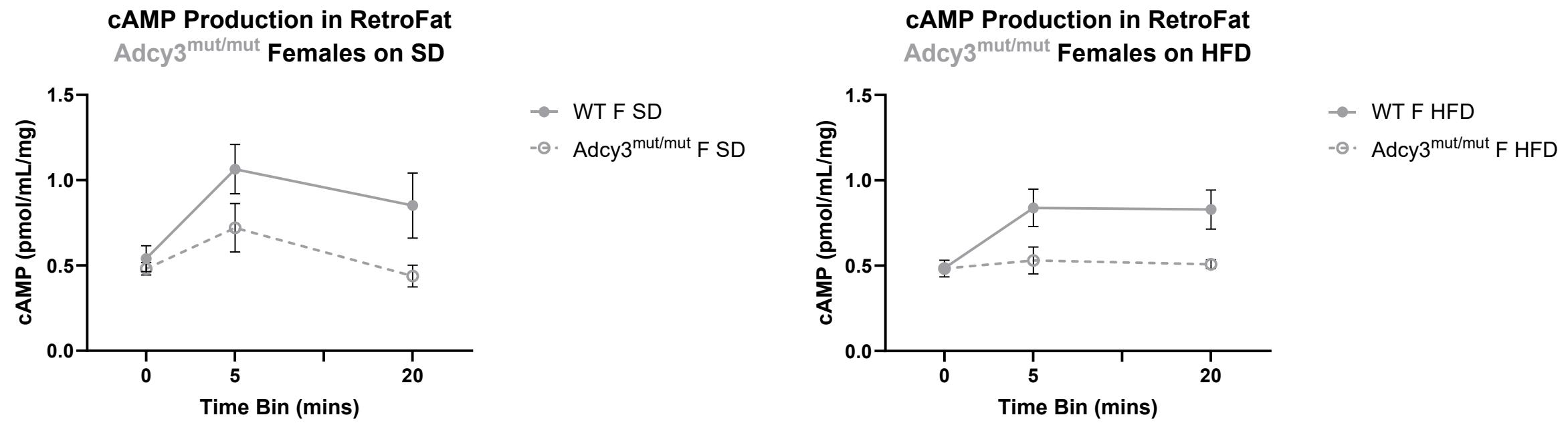


Figure S3

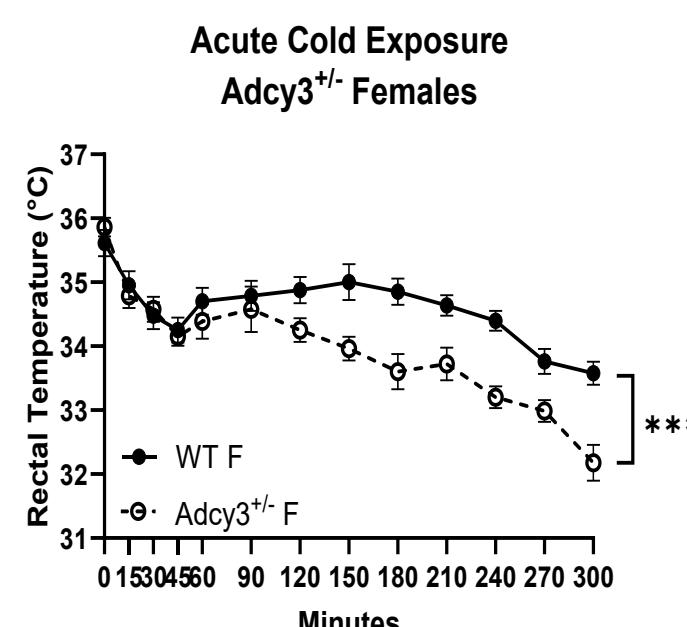
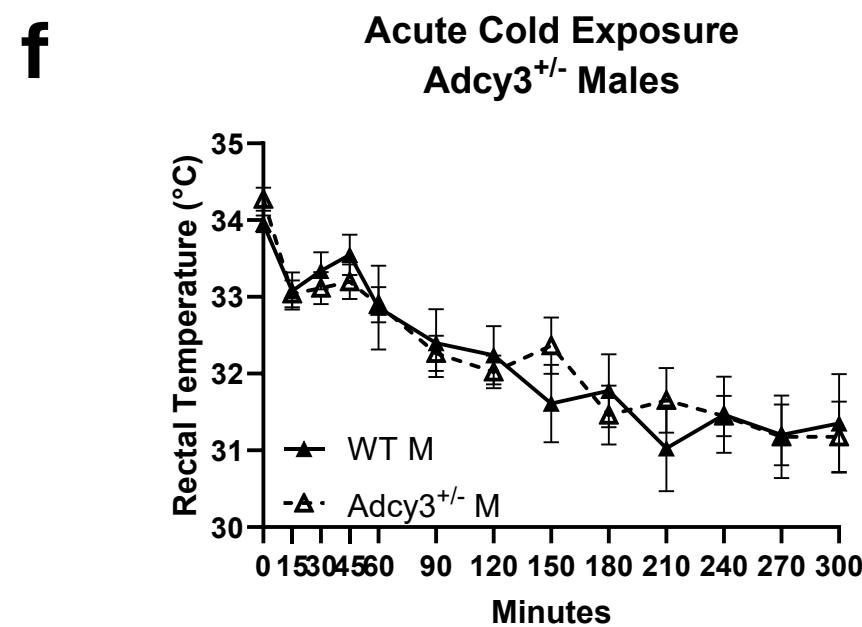
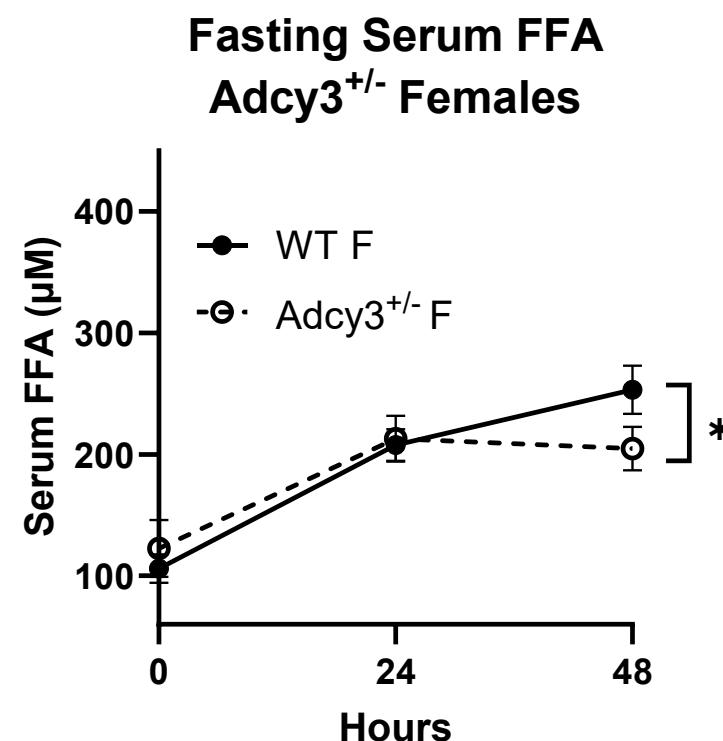
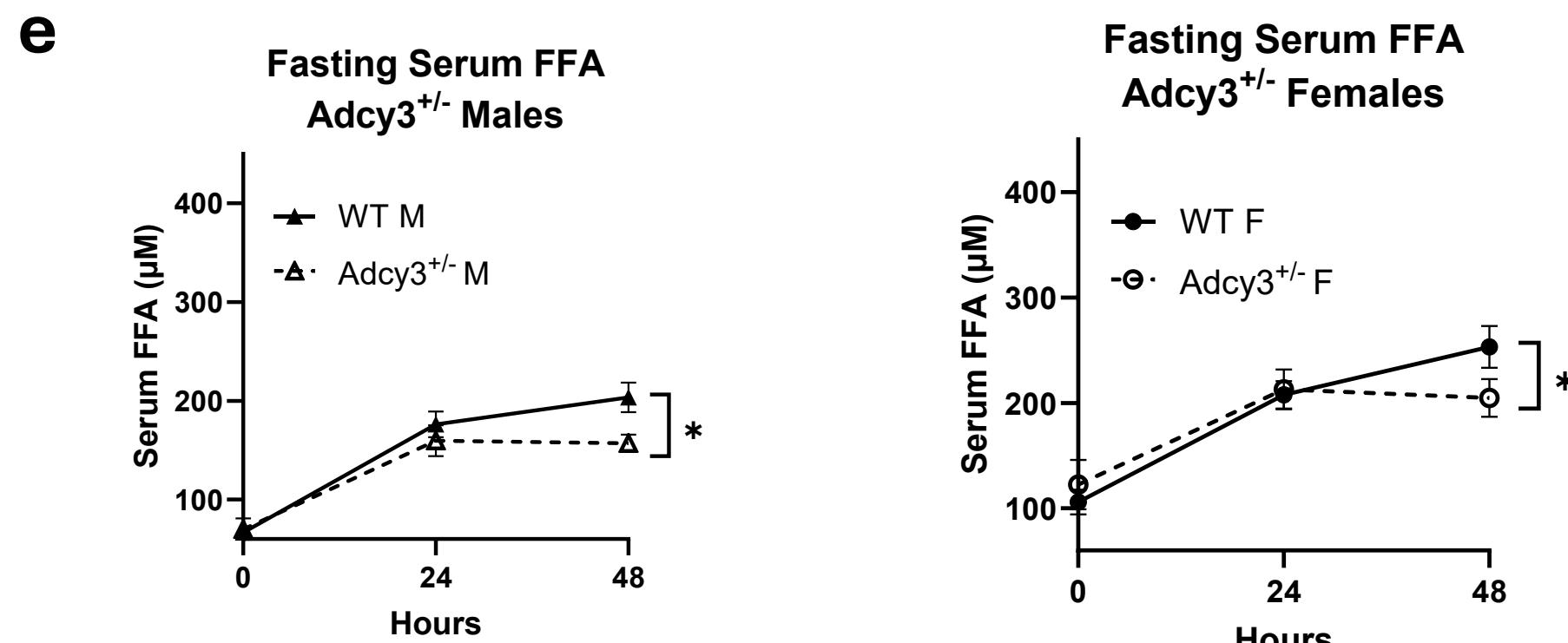
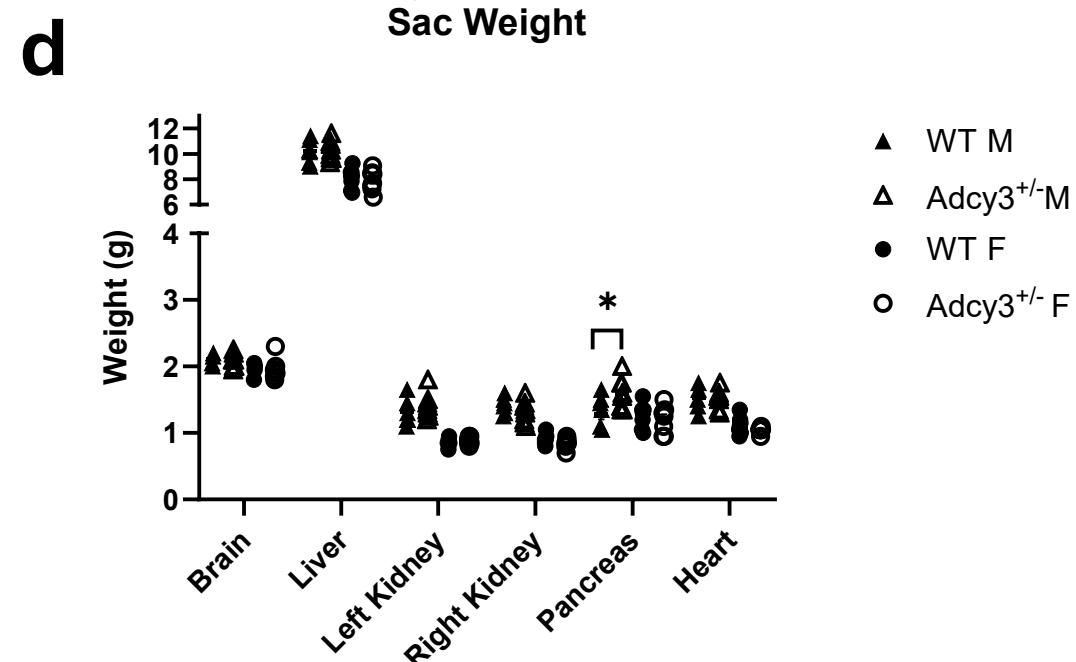
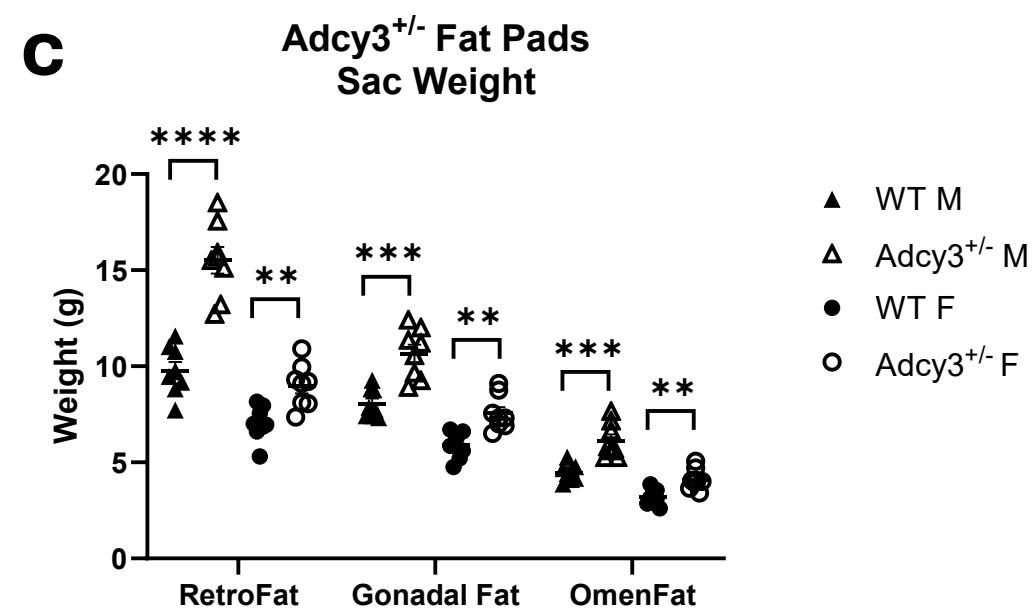
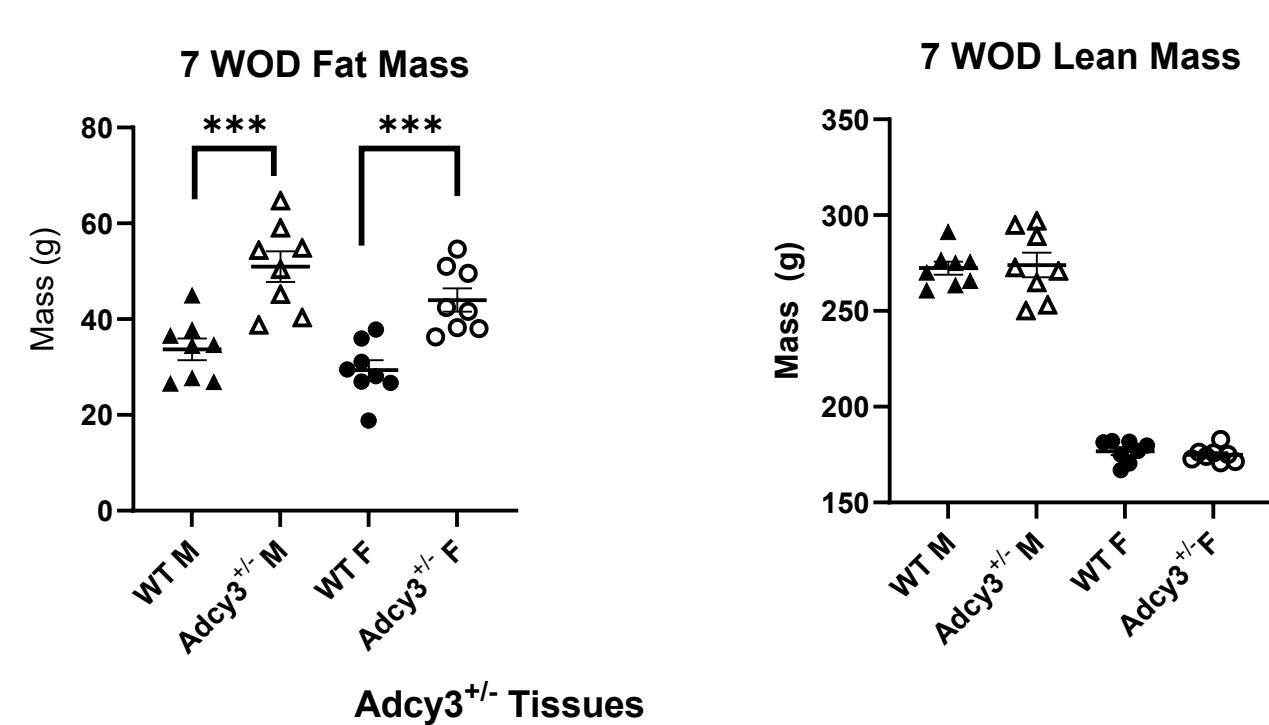
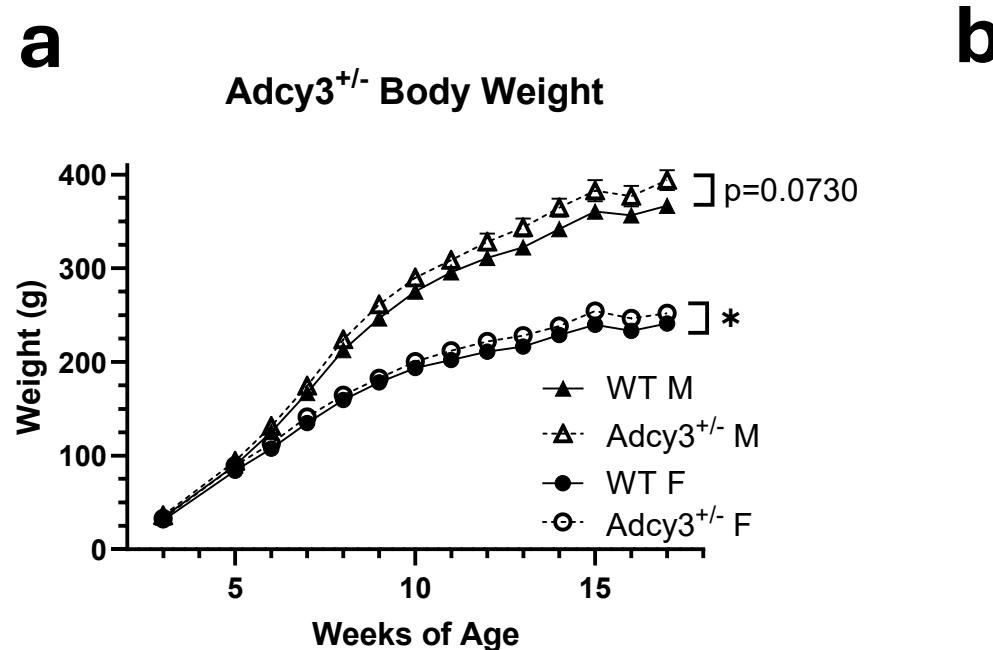
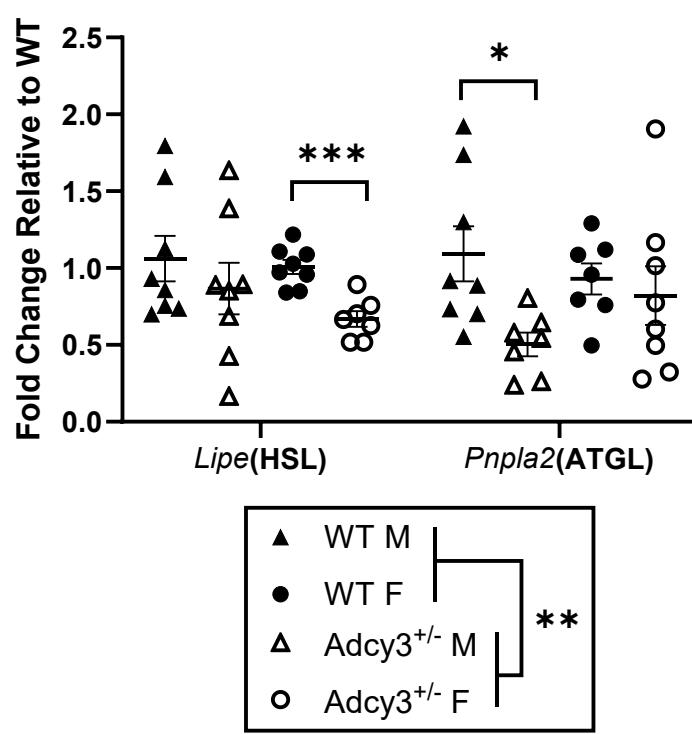


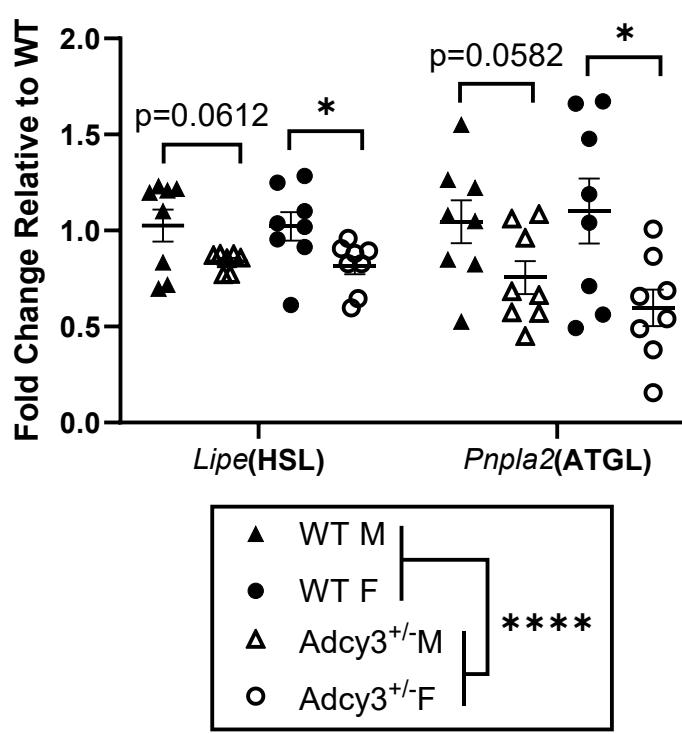
Figure S4

a

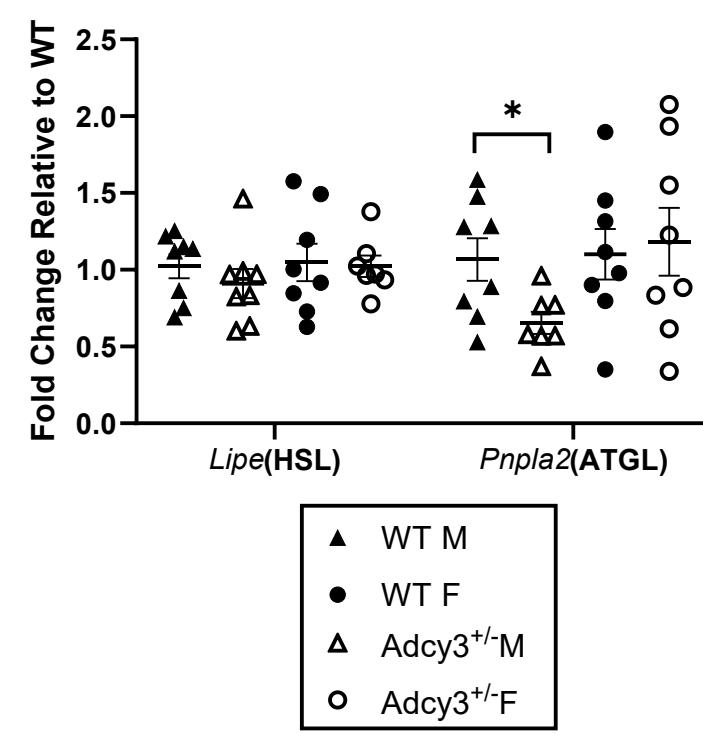
Lipase Gene Expression: Prolonged Cold
Adcy3^{+/−}, RetroFat



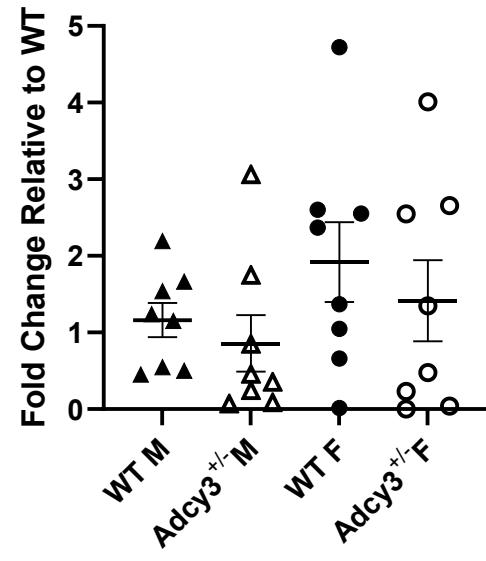
Lipase Gene Expression: Prolonged Cold
Adcy3^{+/−}, SubQ



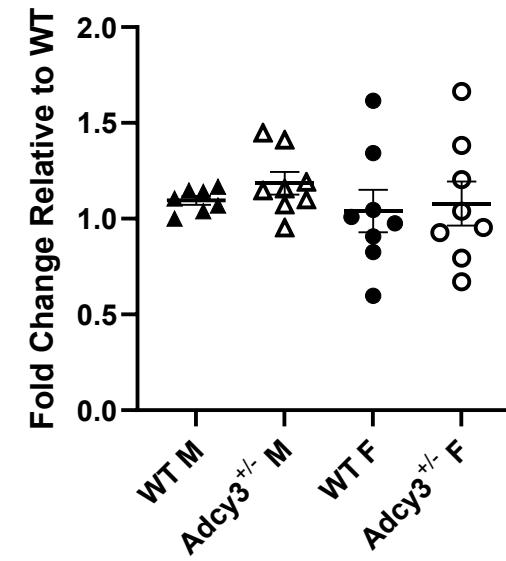
Lipase Gene Expression: Prolonged Cold
Adcy3^{+/−}, BAT

**b**

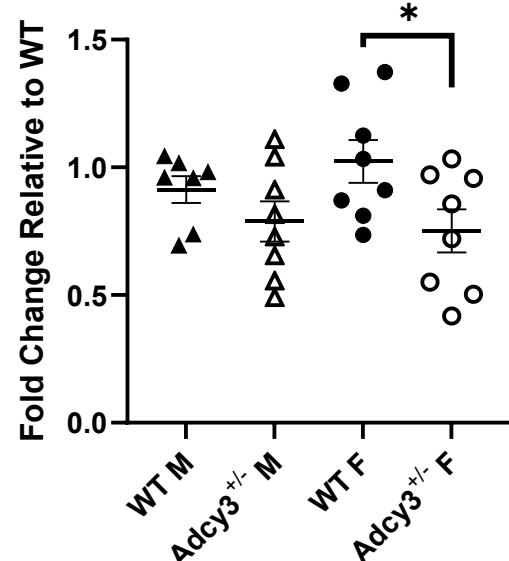
Ucp1 Gene Expression: Prolonged Cold
Adcy3^{+/−}, SubQ



Ucp1 Gene Expression: Prolonged Cold
Adcy3^{+/−}, BAT

**c**

Pparg Gene Expression: Room Temperature
Adcy3^{+/−}, RetroFat



Cebpa Expression: Room Temperature
Adcy3^{+/−}, RetroFat

