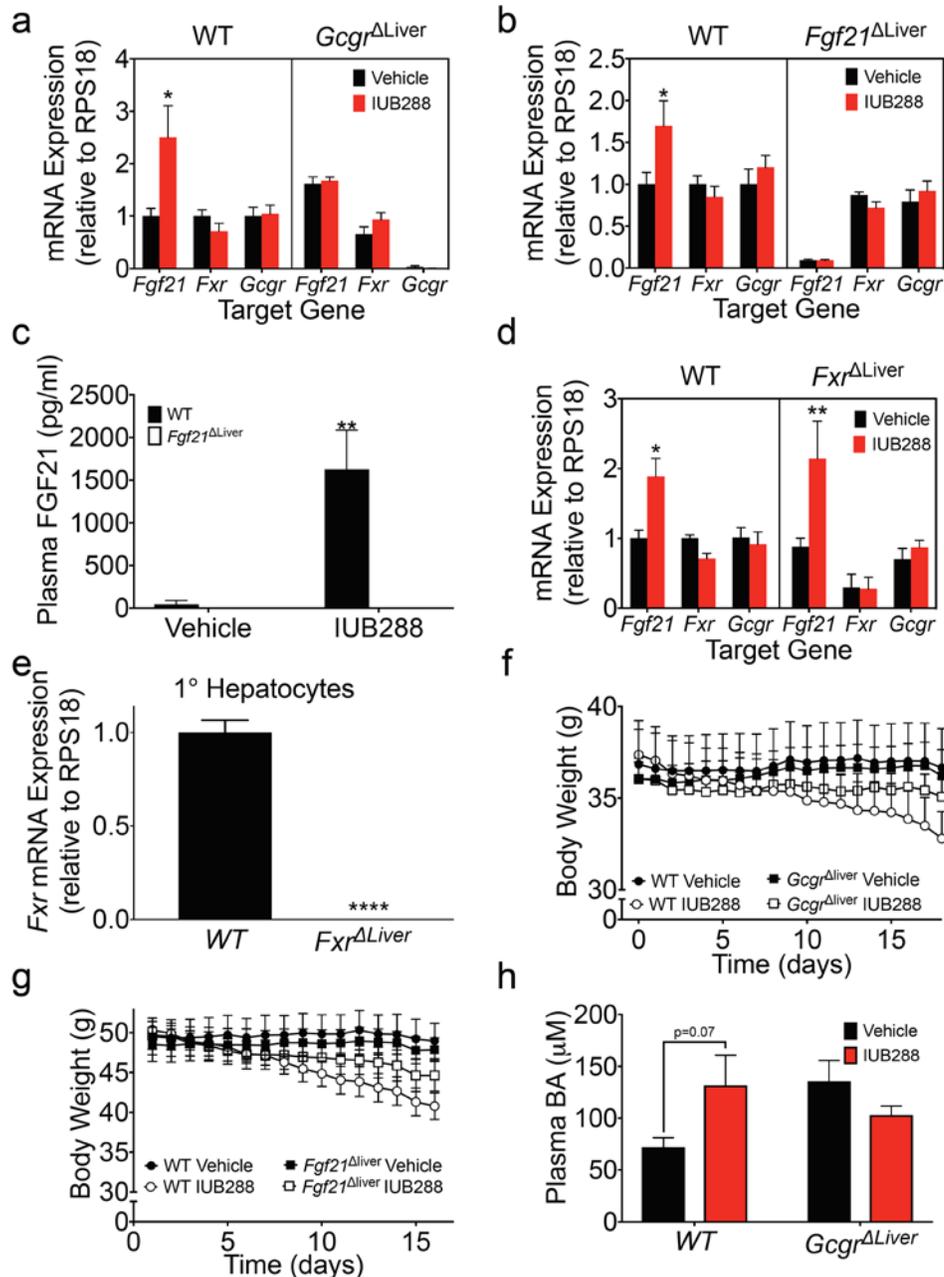


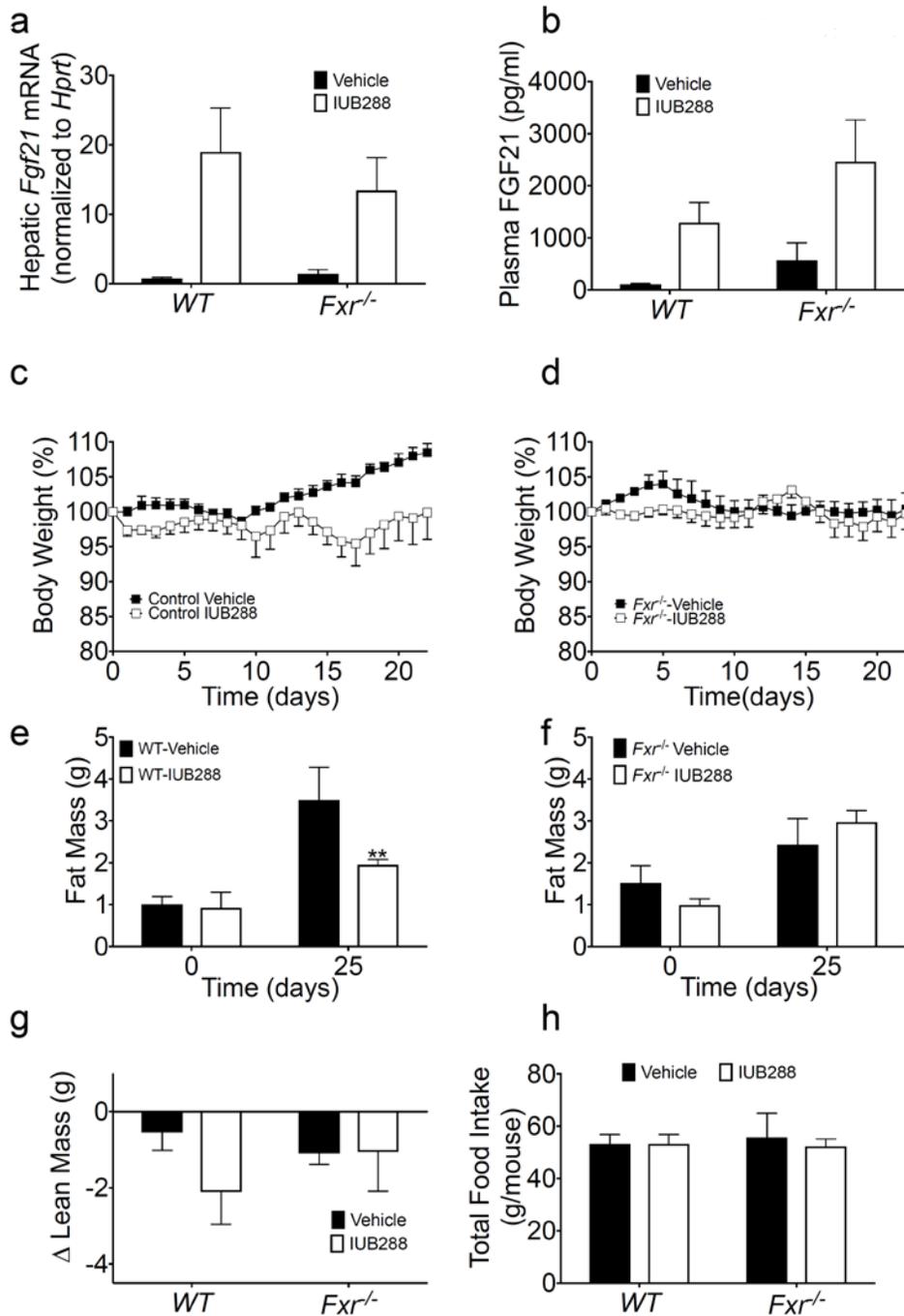
SUPPLEMENTARY DATA

Supplementary Figure S1. Model validation and GcgR agonism in GcgR and Fgf21 mice. mRNA expression of *Gcgr*, *Fgf21*, and *Fxr* in *GcgR* Δ Liver, *Fgf21* Δ Liver, *Fxr* Δ Liver, and littermate Control mice (a, b, and d, n=6-10 mice/group, see Figures 2-4). Plasma FGF21 in 8-week-old, chow fed *Fgf21* Δ Liver and littermate Control (WT) mice following 5d Vehicle or IUB288 treatment (c, 10 nmol/kg IUB288, n=4- 10 mice/group). *Fxr* mRNA expression in primary hepatocytes isolated from 8-10 week old, chow fed *Fxr* Δ Liver or littermate control (WT) mice (e). Body weight of DIO WT and *GcgR* Δ Liver mice (f, n=8-12 mice/group) or WT and *Fgf21* Δ Liver mice (g, n=5-7 mice/group) following daily GcgR agonism (10 nmol/kg IUB288). Plasma bile acid levels in IUB288-treated DIO WT and *GcgR* Δ Liver mice after 2 h fast (h, n=8-12 mice/group, see Figure 2). All data are represented as mean +/- SEM.



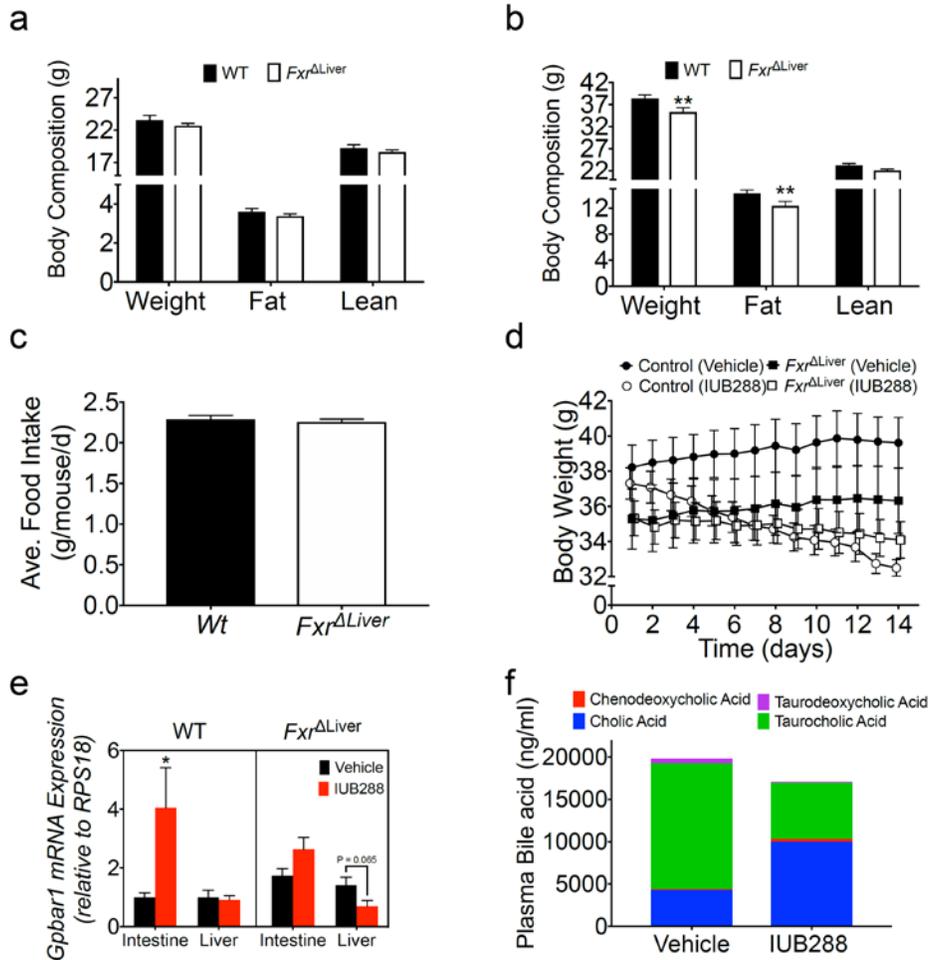
SUPPLEMENTARY DATA

Supplementary Figure S2: GcgR agonism and energy balance in FXR^{-/-} mice. Hepatic *Fgf21* mRNA expression (a) and plasma levels (b) in HF-fed WT and FXR^{-/-} mice. Body weight (%) and fat mass of HF-fed WT (c and e) or FXR^{-/-} mice (d and f) following daily GcgR agonism (10 nmol/kg IUB288). Change in lean mass (g) and total food intake (h) in HF-fed WT or FXR^{-/-} mice. All data are represented as mean +/- SEM (n=3-8 mice/group). **p< 0.01. Male, WT and *Fxr*^{-/-} mice were placed on HFD at 8-10 weeks old concurrent with IUB288 treatment..



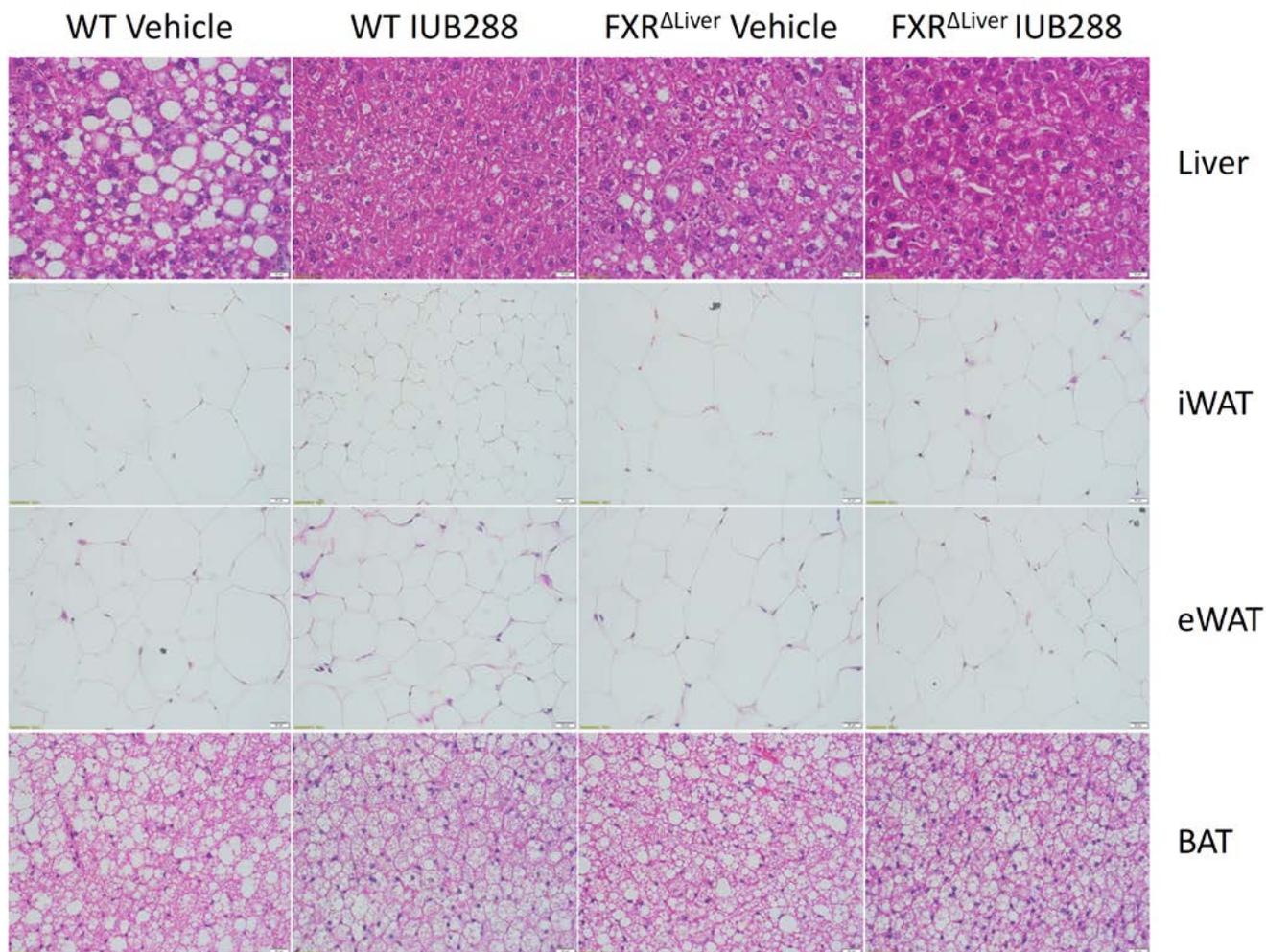
SUPPLEMENTARY DATA

Supplementary Figure S3. DIO and GcgR agonism in FXR Δ Liver mice. Body composition before (a) and after (b) HF-feeding in WT and FXR Δ Liver mice (n=13-15 mice/group). Average food intake (c) during HF-fat feeding in WT and FXR Δ Liver mice (n=13-15 mice/group). Body weight (d) during daily GcgR agonism (10 nmol/kg IUB288) in WT and FXR Δ Liver mice (n=8-10 mice/group). Intestine and liver *Gpbar1/Tgr5* mRNA expression (e) in 14d IUB288-treated DIO WT and Fxr Δ Liver mice. Plasma bile acid profile (f) in male Fxr Δ Liver mice following 16d GcgR agonism. *p< 0.05, **p< 0.01. Male, WT and Fxr Δ Liver mice were placed on HFD at 8-10 weeks old and maintained on HFD for 10 weeks to induce DIO prior to treatment.



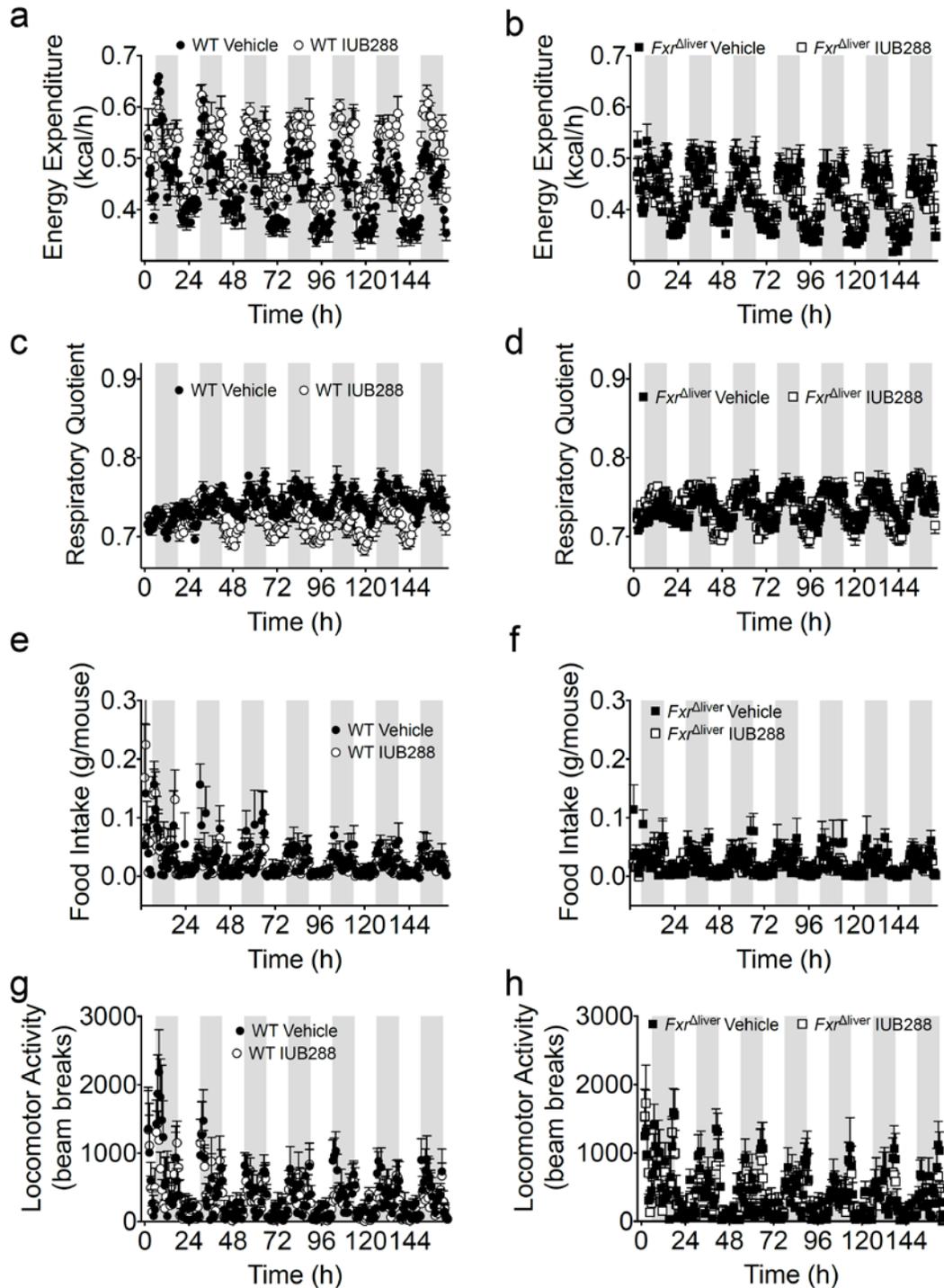
SUPPLEMENTARY DATA

Supplementary Figure S4. Liver and Adipose Tissue morphology following GcgR agonism in FXR Δ Liver mice. Representative haematoxylin and eosin (H & E) staining of liver, inguinal white adipose tissue, ependymal white adipose tissue, and interscapular brown adipose tissue following 14d IUB288 treatment. Male, WT and Fxr Δ Liver mice were placed on HFD at 8-10 weeks old and maintained on HFD for 10 weeks to induce DIO prior to treatment. Scale bars are 20 μ m in length.



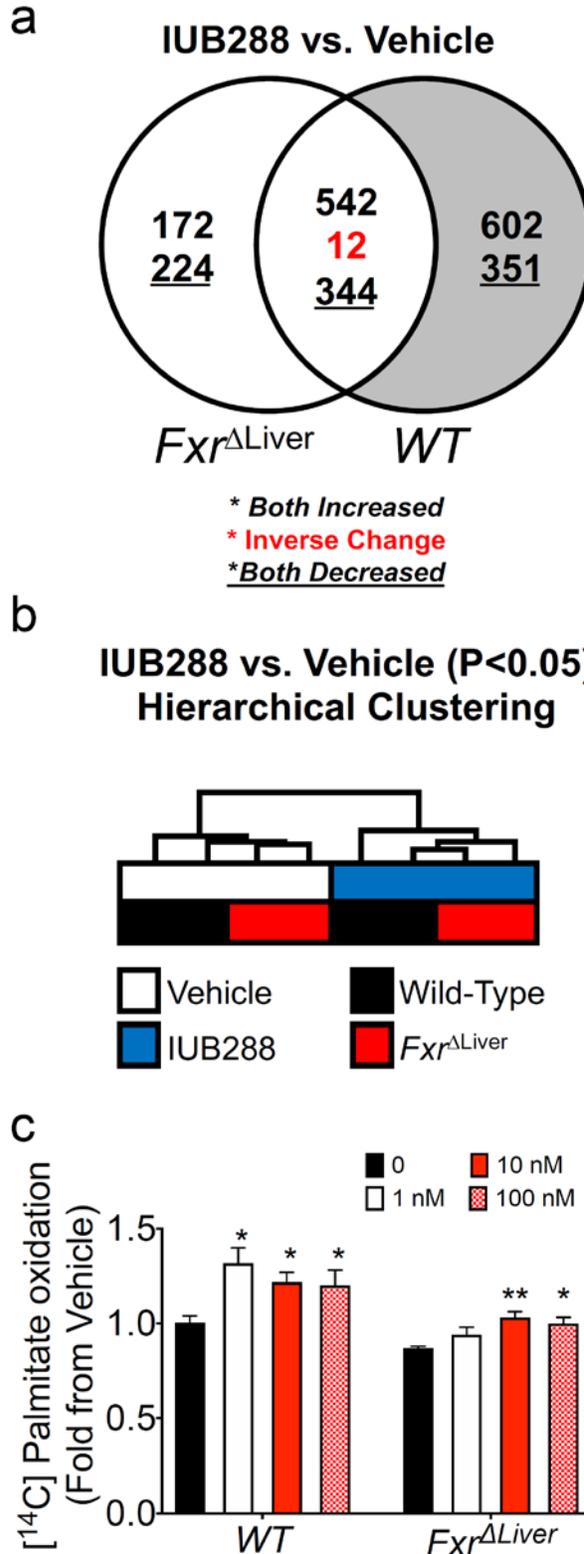
SUPPLEMENTARY DATA

Supplementary Figure S5. 7 d indirect calorimetry during GcgR agonism in *Fxr* Δ Liver mice. Energy expenditure (EE, a-b), respiratory quotient (RQ, c-d), food intake (e-f), and locomotor activity (g-h) measured during 7 d indirect calorimetry analysis (in DIO WT (a,c,e, and g) and *Fxr* Δ Liver mice (b,d,f, and h) during daily GcgR agonism (10 nmol/kg IUB288). IUB288 administered via subcutaneous injection 1hr prior to dark phase (ZT11). All data are represented as mean \pm SEM (n=6 mice/group, see Figure 5).



SUPPLEMENTARY DATA

Supplementary Figure S6. Transcriptional Analysis of IUB288 Treatment in *Fxr* Δ Liver and WT Mice. (a) Venn diagram illustrating the selection of FXR-dependent DEGs (shaded). (b) Hierarchical clustering of IUB288 vs. Vehicle comparison ($p < 0.05$). (c) [^{14}C] Palmitate oxidation in primary hepatocytes isolated from DIO WT and *Fxr* Δ Liver mice and treated with glucagon for O.N. treatment followed by 3 hr incubation with radioactive substrate.



SUPPLEMENTARY DATA

Supplementay Table 1. qPCR primers

Gene	Forward (5'-3')	Reverse (5'-3')
<i>Gcgr</i>	GCCAGCGAGGTCTCCATA	ACATCATTACCTTCTTGTGG
<i>Fgf21</i>	CTG CTG GGG GTC TAC CAA G	CTG CGC CTA CCA CTG TTC C
<i>Scl10a1</i>	GCCACACTATGTACCCTACGTCCTC	GAATGTAGCCCATCAGGAAGCCAGTG
<i>Cyp27a1</i>	GAAGGACCACCGAGACCACAAGG	CGT TTA AGG CAT CCG TGT AGA GCG
<i>Hmgcr</i>	GTGTTCAAGGAGCATGCAAAG	AGCCATCACAGTGCCACATAC
<i>Cyp7a1</i>	GGGATTGCTGTGGTAGTGAGC	GGTATGGAATCAACCCGTTGTC
<i>Fxr</i>	CACAGCGATCGTCATCCTCTCT	TCTCAGGCTGGTACATCTTGCA
<i>Gpbar1/Tgr5</i>	AAGAGCCAAGAGGGACAATC	GTAGCTGCTGCTTCCCTAAT
<i>Ppargc1a</i>	CCCTGCCATTGTTAAGACC	TGCTGCTGTTCCCTGTTTTC
<i>Ppara</i>	AGCAGTGCTGGCTACCTTCAA	AATATGTAGCCACCCCCTTGG
<i>Scd1</i>	TCAGAAACACATGCTGATCCTCAT	TGGGTGTTTGCGCACAAAG
<i>Srebp1c</i>	GAGGACCTTTGTCATTGGCTG	TACAGAGCAAGAGGGTGCCAT
<i>Hprt</i>	AAGGAGATGGGAGGCCAT	GTTGAGAGATCATCTCCACCAAT
<i>Rps18</i>	TTCTGGCCAACGGTCTAGACAAC	CCAGTGGTCTTGGTGTGCTGA
<i>Ppia</i>	CAGACGCCACTGTCGCTT T	TGTCTTTGGAACCTTTGTCTG