

SUPPLEMENTARY DATA

Lactational delivery of Triclosan promotes non-alcoholic fatty liver disease in newborn mice

André A. Weber¹, Xiaojing Yang¹, Elvira Mennillo¹, Jeffrey Ding², Jeramie D. Watrous², Mohit Jain², Shujuan Chen¹, Michael Karin³, Robert H. Tukey^{1,*}.

Short Title: *Triclosan drives neonatal NAFLD*

¹Laboratory of Environmental Toxicology, Department of Pharmacology, University of California, San Diego, La Jolla, CA 92093, USA.

²Departments of Medicine and Pharmacology, University of California, San Diego, La Jolla, CA 92093, USA.

³Laboratory of Gene Regulation and Signal Transduction, Department of Pharmacology, University of California, San Diego, La Jolla, CA 92093.

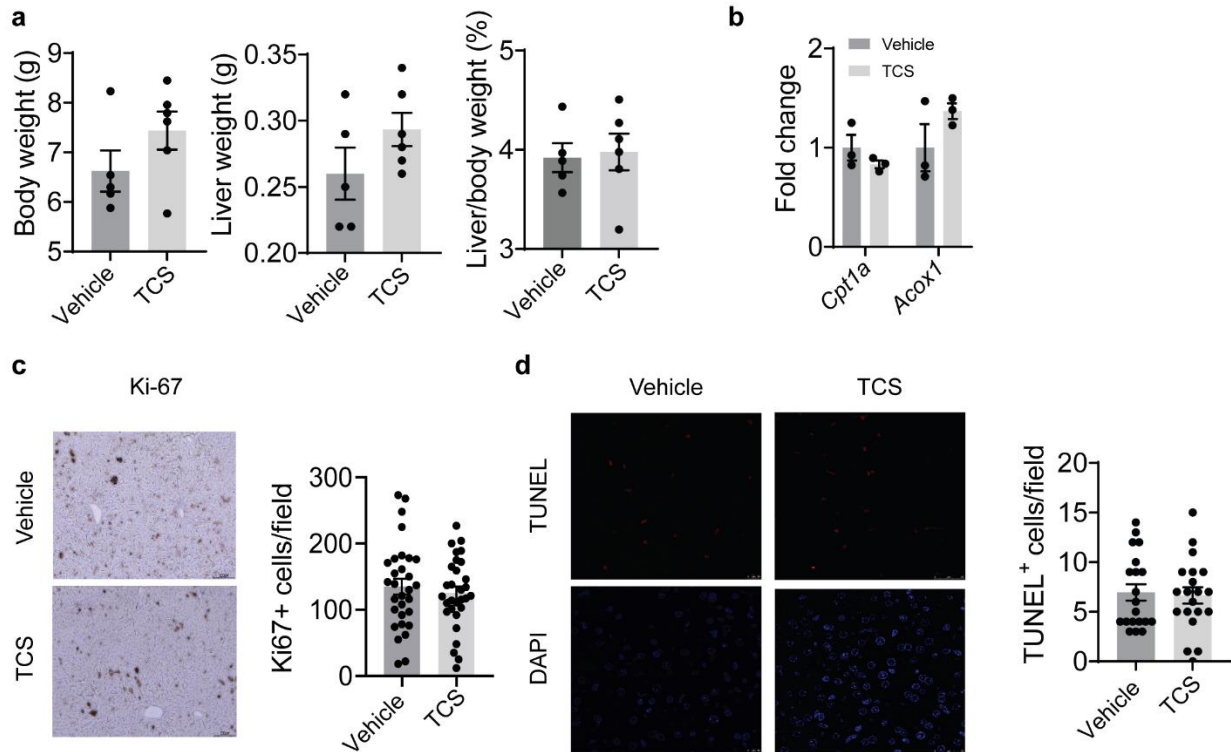


Figure S1. TCS has no effect on body and liver weight and hepatic proliferation and cell death. WT mice breastfed with vehicle or TCS for 21-days-old. **a**, Body weight, liver weight and Liver/Body Weight ratio (%) (n=5-6 mice per group). **b**, Expression of genes associated with fatty acid oxidation *Cpt1a* and *Acox1* (n=3 mice per group). **c**, Hepatocyte's proliferation was determined by Ki-67 immunostaining and quantitation of positive cells (n=30 images/5 mice per group). **d**, Liver cell apoptosis was determined by TUNEL staining and quantitation of positive cells (n=20 images/5 mice per group). Scale bars: **c**, 200 μ m and **d**, 20 μ m. **a-d** show mean \pm S.E., determined by two-tailed Student's test.

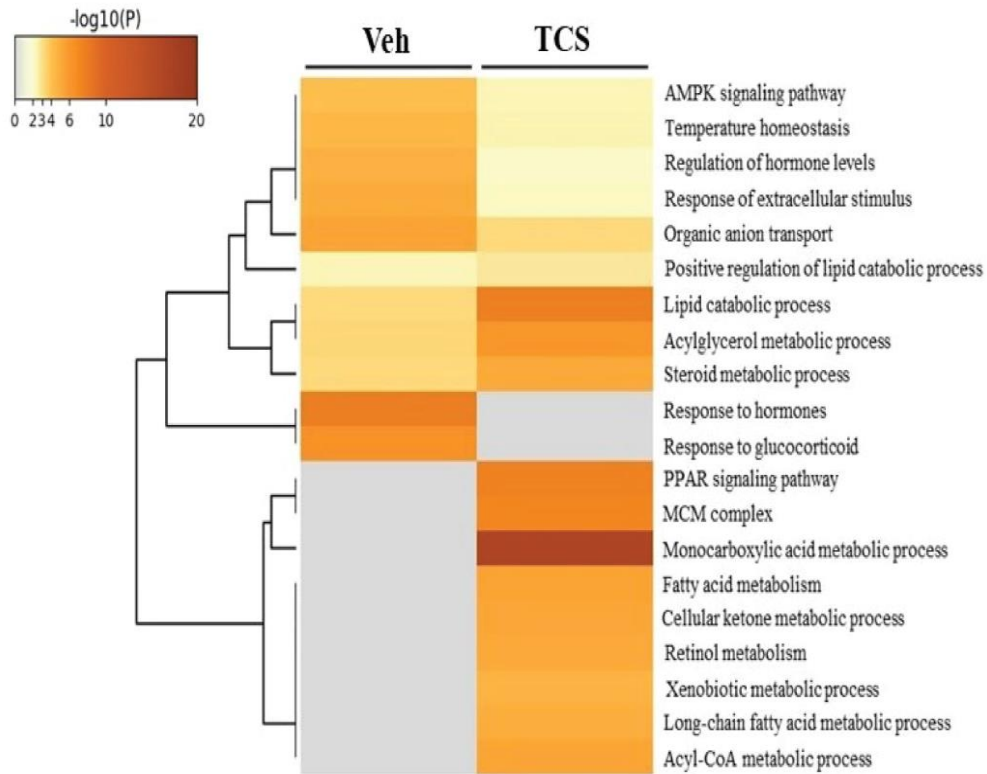


Figure S2. TCS-induced alterations in the liver transcriptome. WT mice breastfed with vehicle or TCS for 21-days-old. Heatmap depicting expression of HCC-related genes in liver tissue of neonatal mice treated with vehicle or TCS (n=3 mice per group). All genes showed in this figure had a p -adjusted value above 0.05, showing no differences between groups.

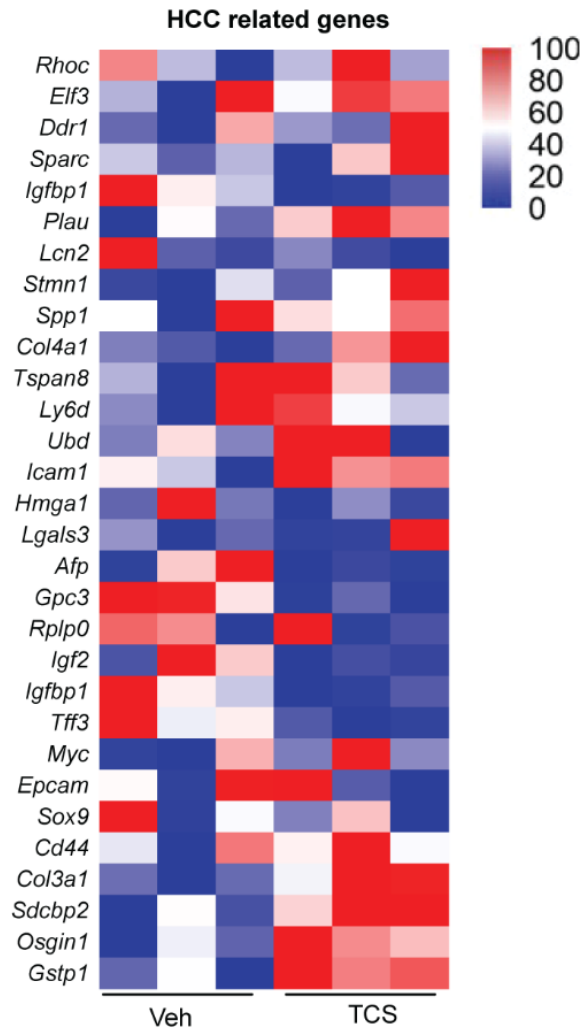


Figure S3. TCS has little effect on HCC-related genes expression. WT mice breastfed with vehicle or TCS for 21-days-old. Heatmap depicting expression of HCC-related genes in liver tissue of neonatal mice treated with vehicle or TCS (n=3 mice per group). All genes showed in this figure had a *p*-adjusted value above 0.05, showing no differences between groups.

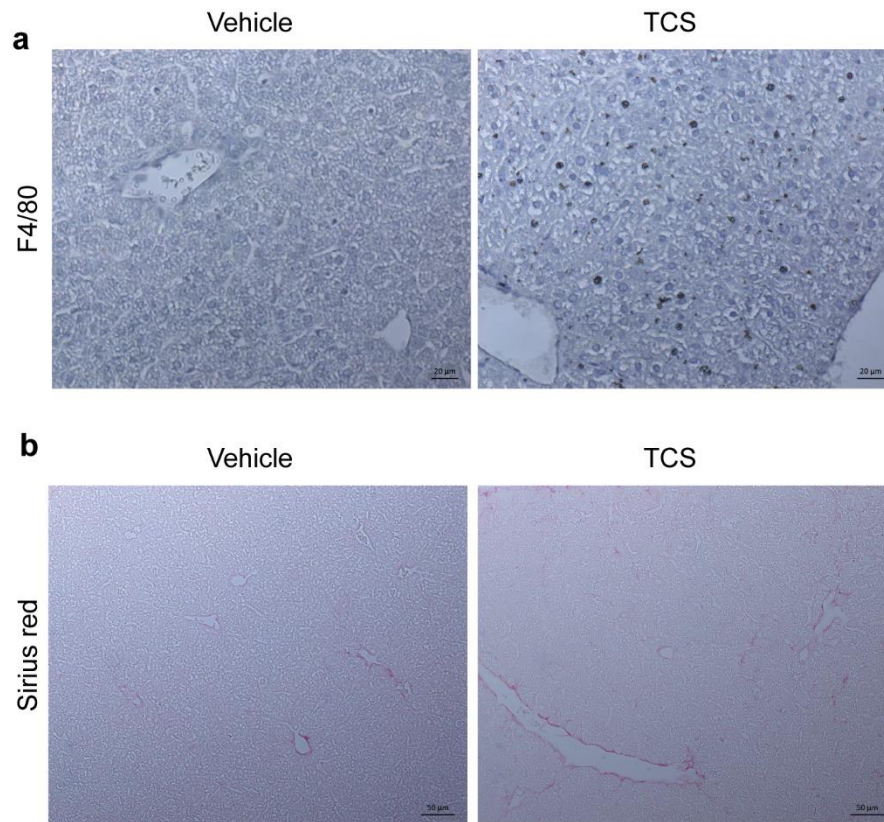


Figure S4. TCS increases immune infiltration and fibrosis. WT mice breastfed with vehicle or TCS milk for 21-days-old. **a**, IHC of formalin-fixed paraffin-embedded liver sections from vehicle and TCS group F4/80 immunostaining show macrophage infiltration (n=4 mice per group). **b**, Frozen sections were stained by Sirius Red to visualize collagen deposition (n=4 mice per group). Scale bars: **a**, 20µm and **b**, 50µm.

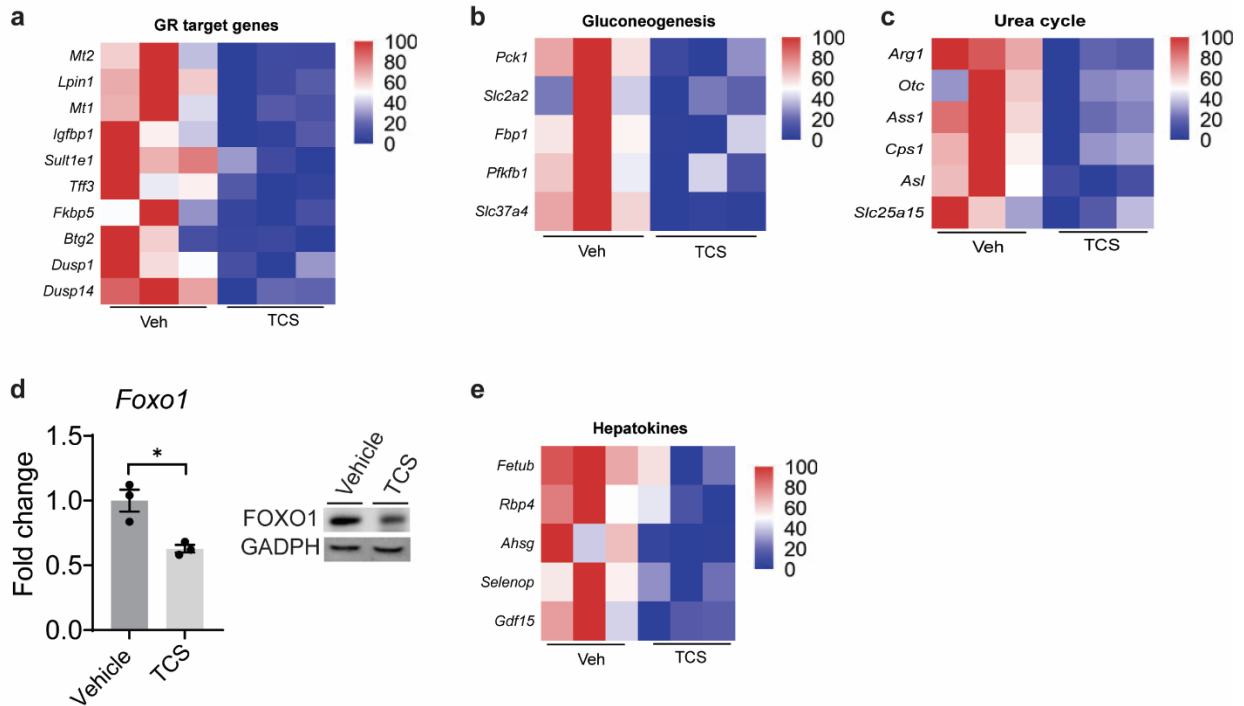


Figure S5. TCS downregulates glucocorticoid receptor and downstream pathways. WT mice breastfed with vehicle or TCS for 21-days-old. **a-c** Heatmaps showing differential expression of genes associated with glucocorticoid receptor (GR) signaling (**a**), and genes associated with downstream GR pathways, gluconeogenesis (**b**) and urea cycle (**c**) (n=3 per group). **d**, gene expression and representative IB image of FOXO1 in liver. **e**, Heatmap showing differential expression of hepatokines in liver (n=3 per group). **d** show mean \pm S.E., determined by two-tailed Student's test; * $P < 0.05$. All genes used are significant and p adjusted value below the cut-off level of 0.05.

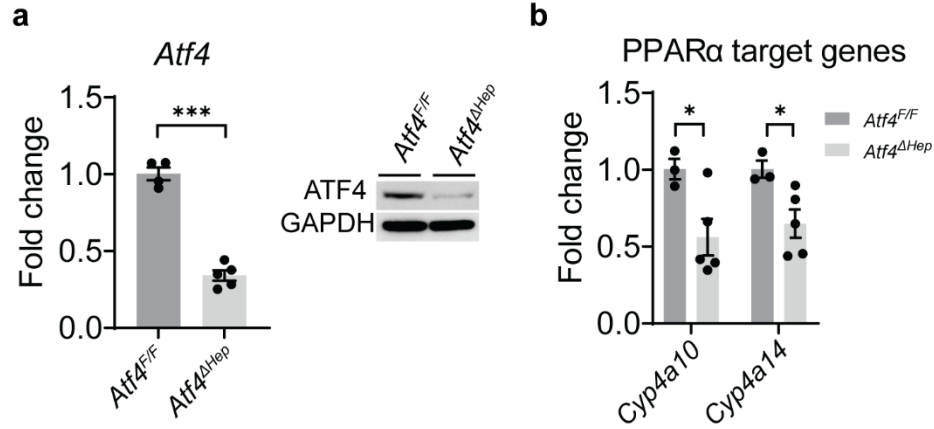


Figure S6. Downstream **PPAR α** target genes are downregulated in ***Atf4^{ΔHep}*** mice treated with **TCS**. *Atf4^{F/F}* and *Atf4^{ΔHep}* mice breastfed with TCS milk 21-days-old. **a**, Hepatic expression of ATF4 gene in both *Atf4^{F/F}* and *Atf4^{ΔHep}* livers and IB of ATF4 (n=4-5 mice per group). **b**, Hepatic expression of downstream PPAR α target genes in *Atf4^{F/F}* and *Atf4^{ΔHep}* (n=3-5 mice per group). **a**, **b** show mean \pm S.E., determined by two-tailed Student's test; *P<0.05, ***P<0.001.

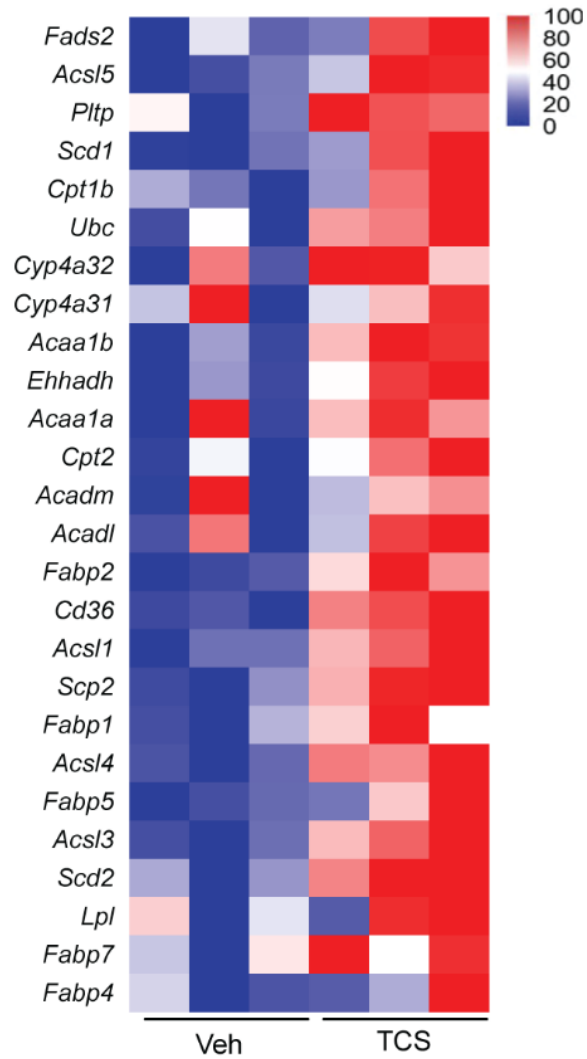


Figure S7. TCS has strong effect on PPAR α target genes. WT mice breastfed with vehicle or TCS for 21-days-old. Heatmap depicting expression of downstream PPAR α target genes in liver tissue of neonatal mice treated with vehicle or TCS (n=3 mice per group). All genes showed in this figure had a p-adjusted value above 0.05, showing no differences between groups.

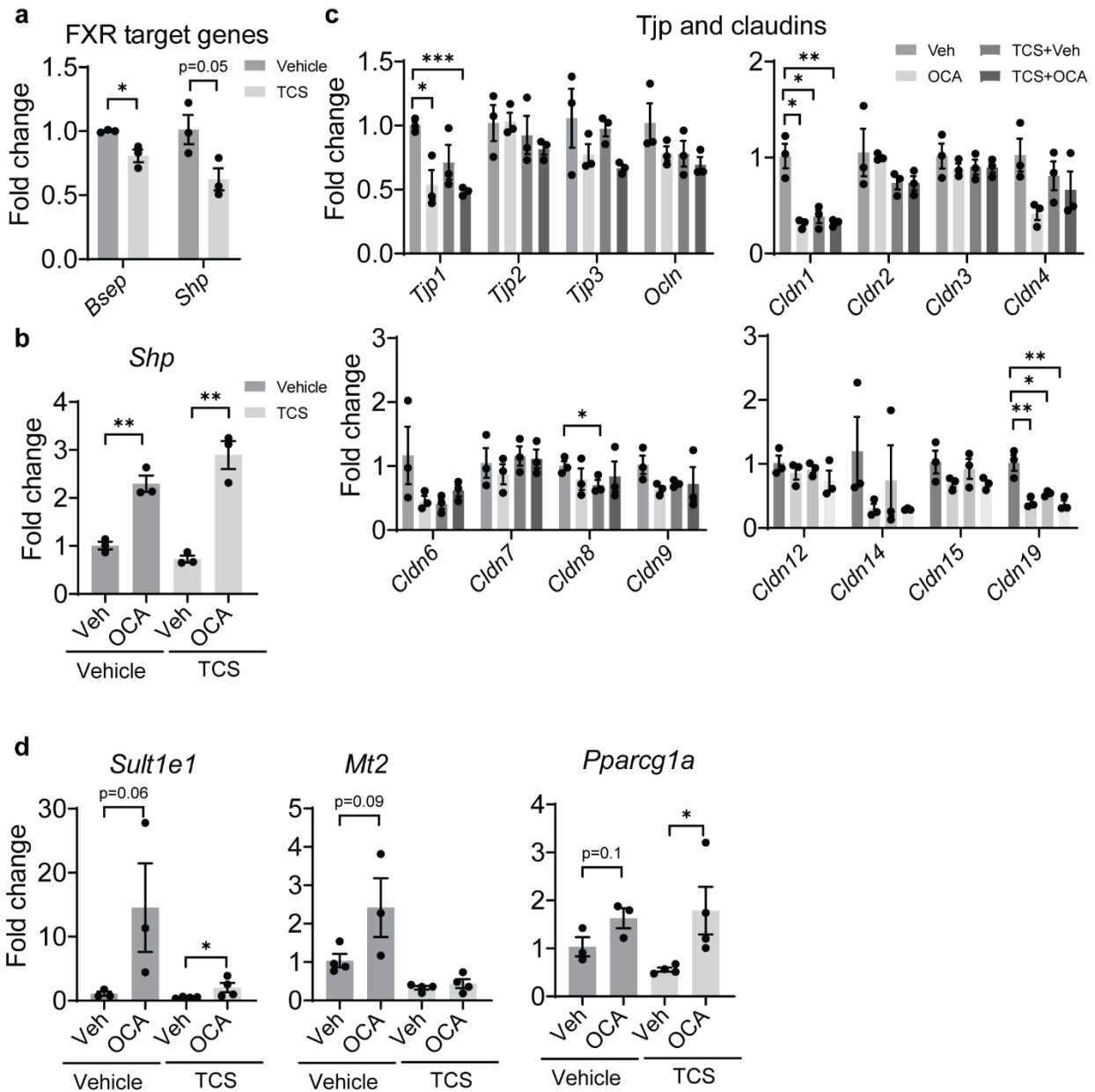


Figure S8. OCA has no effect in intestinal barrier genes but increases GR response in liver in the absence of TCS. WT neonatal mice breastfed with vehicle or TCS milk for 21 days and treated with vehicle or OCA. **a**, Hepatic expression of genes associated with FXR (n=3 mice per group). **b**, Intestinal expression of *Shp* (n=3 mice per group). **c**, Intestinal expression of TJPs, occludin and claudins (n=3 mice per group). **d**, Hepatic expression GR related genes (n=3-4 mice per group). **a-d** show mean \pm S.E., determined by two-tailed Student's test; *P<0.05, **P<0.01, ***P<0.001.

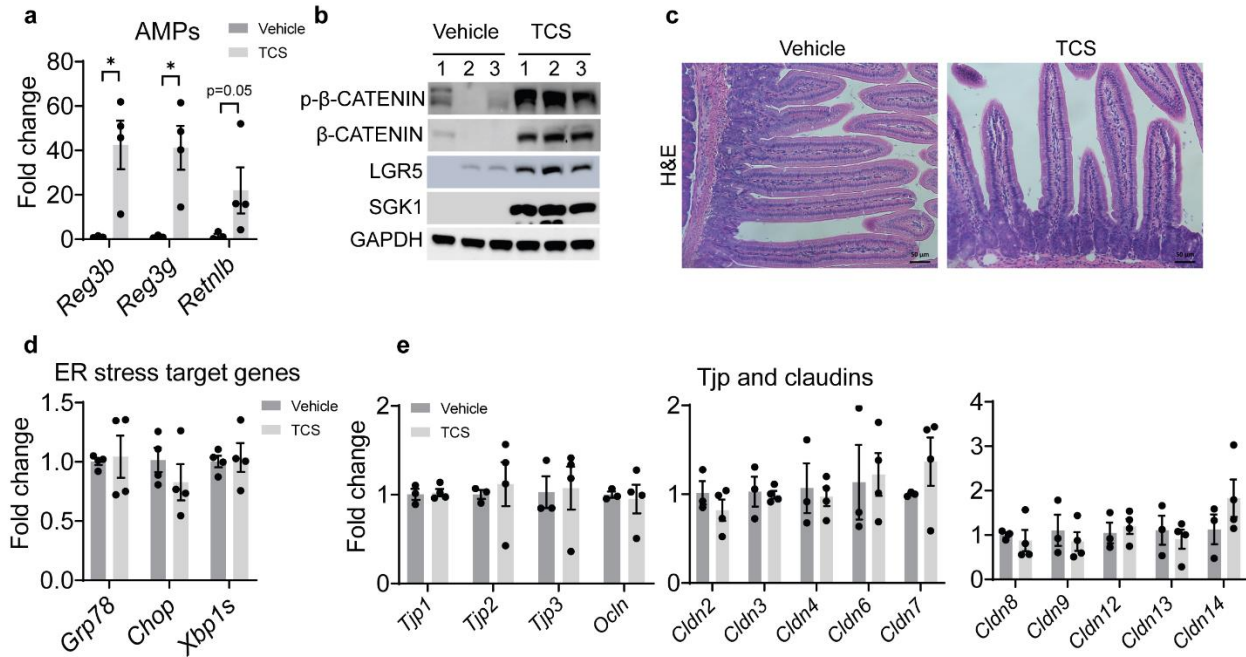


Figure S9. TCS treatment upregulates AMPs and β -catenin pathway in intestines. WT neonatal mice breastfed with vehicle or TCS milk for 21 days. **a**, Intestinal expression of *Reg3b*, *Reg3g* and *Retnlb* (n=3-4 mice per group). **b**, Intestinal IB of total and phosphorylated β -catenin and LGR5 and SGK1 (n=3 mice per group). **c**, Jejunum histological sections were stained with H&E (n=3 per group), Scale bars=50 μ m. **d**, Intestinal expression of genes associated with ER stress (n=4 mice per group). **e**, Intestinal TJP and claudins genes (n=3 mice per group). **a**, **d** and **e** show mean \pm S.E., determined by two-tailed Student's test; *P<0.05.

Table

Table S1. NAFLD semi-quantitative score in neonatal mice breastfed with TCS, according to Brunt (Brunt, E.M. Nonalcoholic steatohepatitis: pathologic features and differential diagnosis. *Semin Diagn Pathol* **22, 330-338, 2005)**

Histopathology	Vehicle	TCS
Steatosis	0	1
Inflammation	0	1
Fibrosis	0	1
Balloning	0	0
Necrosis	-	-
Total score	0	3

Steatosis: hepatocytes exhibiting macro/microvacuolization, 0 (<5%); 1 (5-33%); 2 (33-66%); 4 (>66%). **Inflammation:** foci per 200x field, 0 (no foci); 1 (<2 foci per field); 2 (2-4 foci per field); 3 (>4 foci per field). **Fibrosis:** 0 (none); 1 (mild); 2 (moderate); 3 (severe); 4 (cirrhosis). **Ballooning:** 0 (none); 1 (few balloon cells); 2 (many balloon cells). **Necrosis:** not reported in NAFLD.

Table S2. Primer sequences used for Reverse Transcription Quantitative-PCR (RT-qPCR).

Gene name	Forward (5' - 3')	Reverse (5' - 3')
<i>Acaca</i>	TTCCACGTAGGAGGAGCTTCC	CCTCCGGTGCCTTCTCATTAC
<i>Acly</i>	CTGTGCCACCATGTTCTCCTC	AGGCCTGGTTCTTGGCTACTG
<i>Atf4</i>	TTGTCCGTTACAGCAACACTG	GCAGCAGCACCAGGCTCT
<i>Bsep</i>	AAGGACAGCCACACCAACTC	CCAGAACATGACAAACGGAA
<i>Ccne1</i>	TTGCAAGACCCAGATGAAGA	TCCACGCATGCTGAATTATC
<i>Cldn1</i>	ATCACCTTCGGGAGCTCAGGT	TGATGGGGGTCAAGGGGTCAT
<i>Cldn2</i>	ATGCCTTCTTGAGCCTGCTT	CAGTGTCTCTGGCAAGCTGA
<i>Cldn3</i>	CGTACCGTCACCACTACCAG	CAGCCTAGCAAGCAGACTGT
<i>Cldn4</i>	CCATGGAACCCTTCCGTTGA	ACCCGTCCATCCACTCTACA
<i>Cldn6</i>	CCCTTGGTGGCTGATGCTCAA	AGGTGGAGCTTGGACTCAGGT
<i>Cldn7</i>	TGTACCTACCTGGTCCTGGG	TCTCAGAAAAGACGGGACGC
<i>Cldn8</i>	GGAATGCCAATCCATCACGC	CTCTTTTATCCCCAGGCCCC
<i>Cldn9</i>	AAGTGGTATGGGAGGGGCTGT	CGCAGGTGGAAGCTTCTGGAA
<i>Cldn12</i>	TGTGTGCAGATGTGCTCCTGT	GCAGGAGGGCTTGAGCTGTAT
<i>Cldn14</i>	CTAACCCAGAGGGCATGTGTGC	AGTCCCATCCACCTTGATGCT
<i>Cldn15</i>	GGTGGCTATCTCGTGGTACG	GCACTCCAGCCCAAGTAGAG
<i>Cldn19</i>	GGAATTCTTCAACCCAGCAC	ATAGGGCTGTGGGATGCTGTT
<i>Chop</i>	CCACCACACCTGAAAGCAGAA	AGGTGAAAGGCAGGGACTCA
<i>Ctgf</i>	CAGACTGGAGAAGCAGAGCC	GCTTGGCGATTTTAGGTGTC
<i>Cyclophiline</i>	CAGACGCCACTGTGCTT	TGTCTTTGGAACCTTGTCTGC
<i>Cyp4a14</i>	ACCCCTCTAGATTTGCACCA	AGCAAACCTGTTTCCCAATGC
<i>Cyp4a10</i>	CACACCCTGATCACCAACAG	TCCTTGATGCACATTGTGGT
<i>Edem1</i>	CTACCTGCGAAGAGGCCG	GTTTCATGAGCTGCCCACTGA
<i>Ehhadh</i>	CTATGATCCGCCTCTGCAA	TGGCTCTAACCGTATGGTCC
<i>Fabp1</i>	TGCAGAGCCAGGAGAACTTT	GATTTCTGACACCCCTTGA
<i>Fgf21</i>	CTCCAGCAGTTCTCTGA	CCTGGGTGTCAAAGCCTCTA
<i>Fasn</i>	AGAAGAGCCATGGAGGAGGTG	ATGTCCACACCACCAATGAGG
<i>Grp78</i>	TTCAGCCAATTATCAGCAAACCTCT	TTTTCTGATGTATCCTCTTACCAGT
<i>Mlxip1</i>	CGGATACGGACTTGGAGGATC	GAAGTGTCCGCTGTGGATGAC
<i>Mt2</i>	TTGCGCTCGACCCAATACTC	CATTGTTTGCATTTGCAGGCG
<i>Ocln</i>	TCCGGATCCTGTCTATGCTCA	ATAGCCACCTCCGTAGCCAAA
<i>Perk</i>	TCATCCAGCCTTAGCAAACC	ATGCTTTCACGGTCTTGGTC

<i>Pparcg1a</i>	AAGTGGTGTAGCGACCAATCG	AATGAGGGCAATCCGTCTTCA
<i>Reg3b</i>	CCCAAGGGCTCCCAGGCTTA	GAGGTGTCCTCCAGGCCTCT
<i>Reg3g</i>	TGGCGCTGAAGCTTCCTTCC	TCATAGCCCAGTGTCTGGGTCA
<i>Retnlb</i>	CCATTTCTGAGCTTTCTGG	AGCACATCCAGTGACAACCA
<i>Scd1</i>	GCTCTACACCTGCCTCTTCG	CAGCCGAGCCTTGTAAGTTC
<i>Shp</i>	CACGATCCTCTTCAACCCAG	AGACTTCACACAGTGCCCA
<i>Srebf1</i>	GGCTCTGGAACAGACACTGG	TGGTTGTTGATGAGCTGGAG
<i>Sult1e1</i>	AAACTCACCTGCCACCCAAG	TTGGCGTTCGGCAAAGATA
<i>Xbp1s</i>	CTGAGTCCGAATCAGGTGCAG	GTCCATGGGAAGATGTTCTGG
<i>Tjp1</i>	GAAACTCTGCTGAGCCCCCTA	GTTTTAGGGTCACCCGACGAG
<i>Tjp2</i>	CGAAGCAGTCTGGGTCTCTGA	CCGGCTCCTCTAGCTCATTGT
<i>Tjp3</i>	ATGGTATGCCATTTCGGAACC	CCGGGTACAACGTGTCCACTA