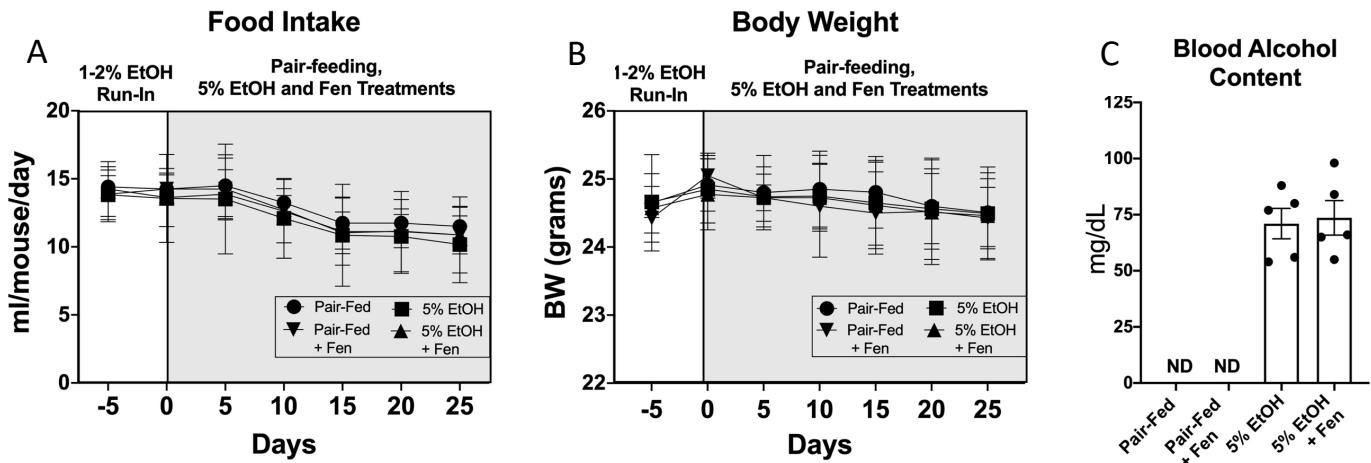
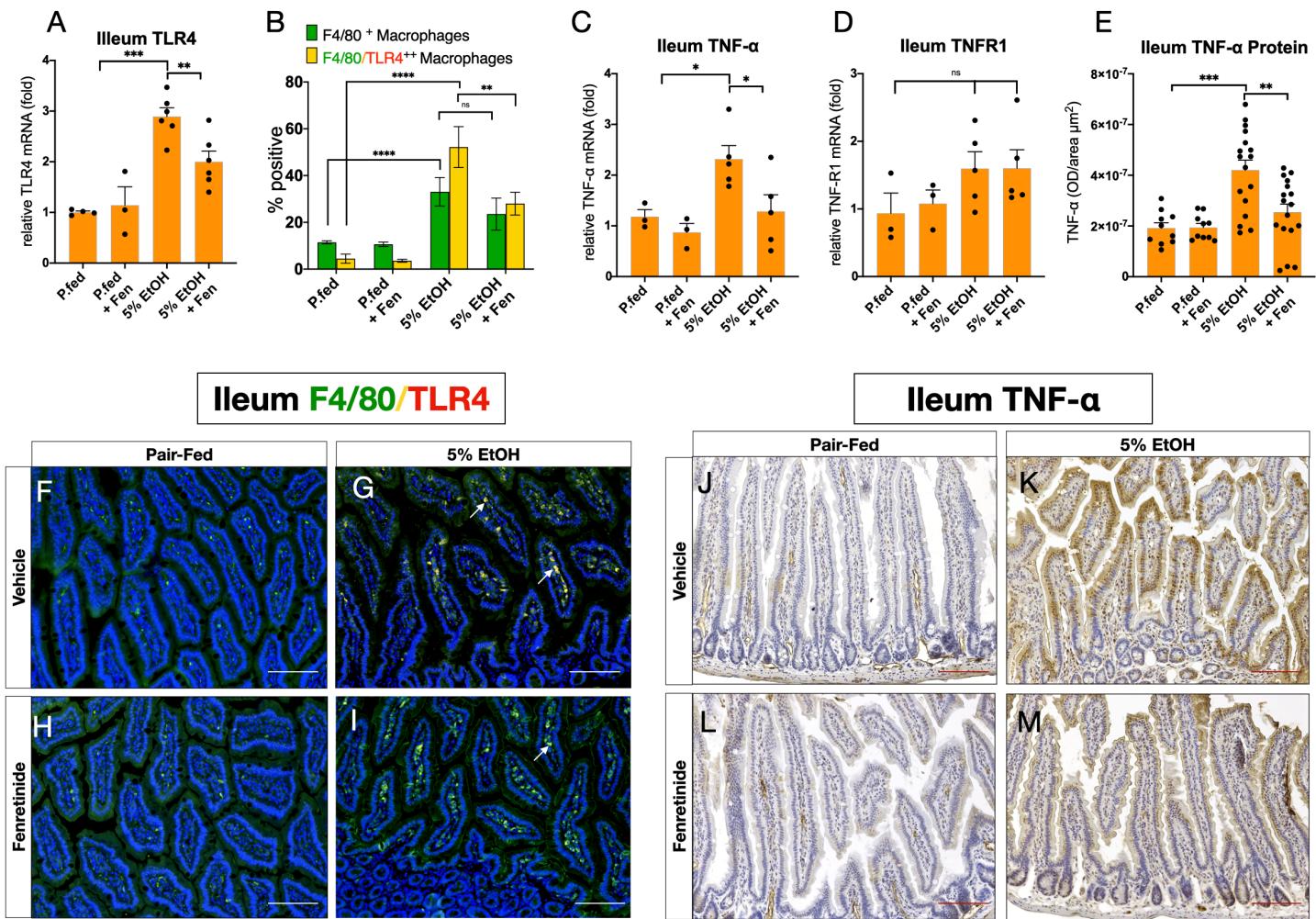


Supplementary Material

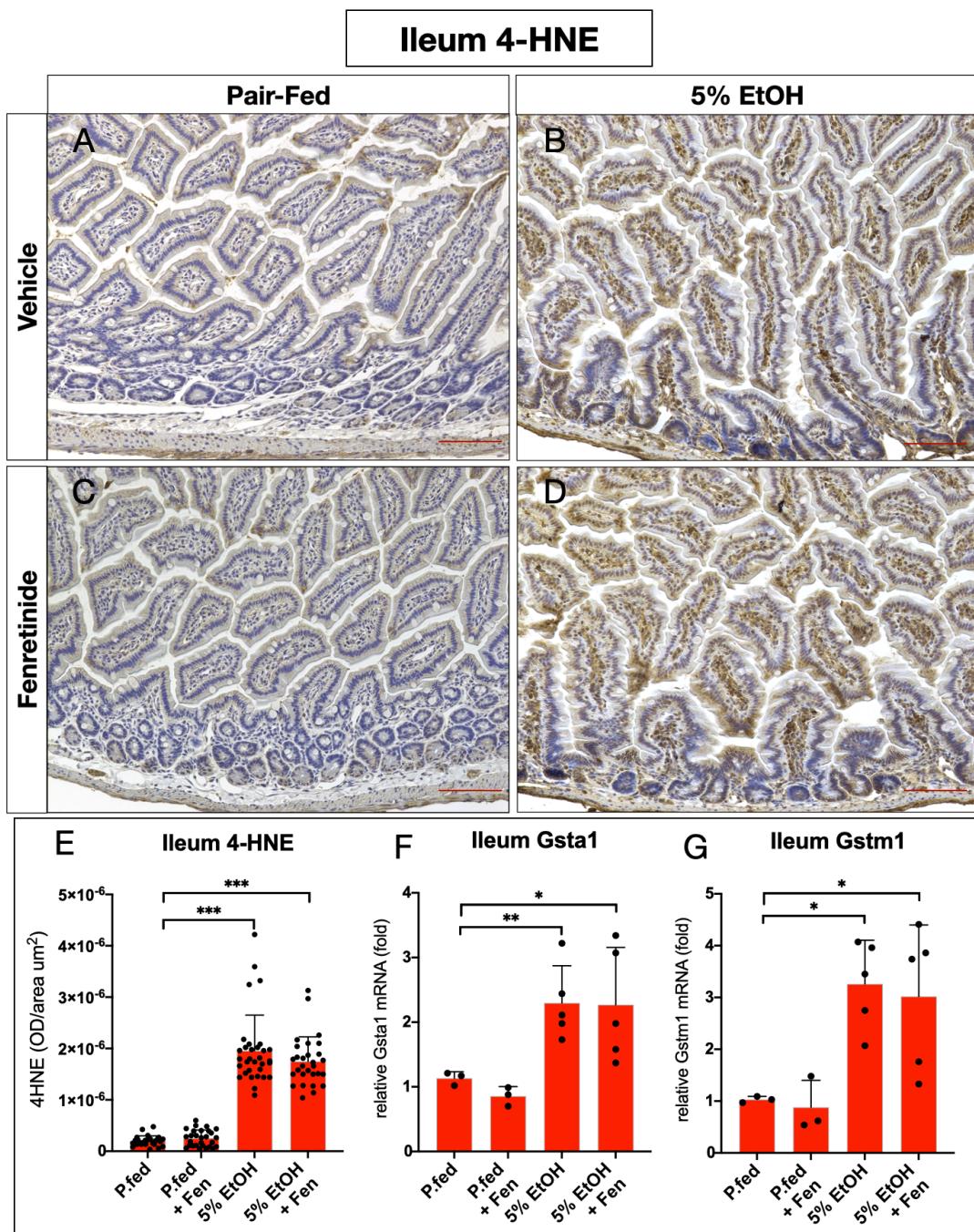


Supplementary Figure 1. A) Food intake in pair-fed (no EtOH) and 5% ethanol (EtOH)-fed C57BL/6 mice treated with either vehicle (0.1% DMSO) [pair-fed, EtOH-untreated] or fenretinide (10mg/kg/bw/day) [pair-fed-Fen, EtOH+Fen for 25 days]. B) Body weight of mice described in A). C) Blood alcohol content (BAC) in mice 2 hours after peak blood alcohol content relative to pair-fed mice described in A). All data errors bars represent \pm SD, with ND=not detected.

Supplementary Material

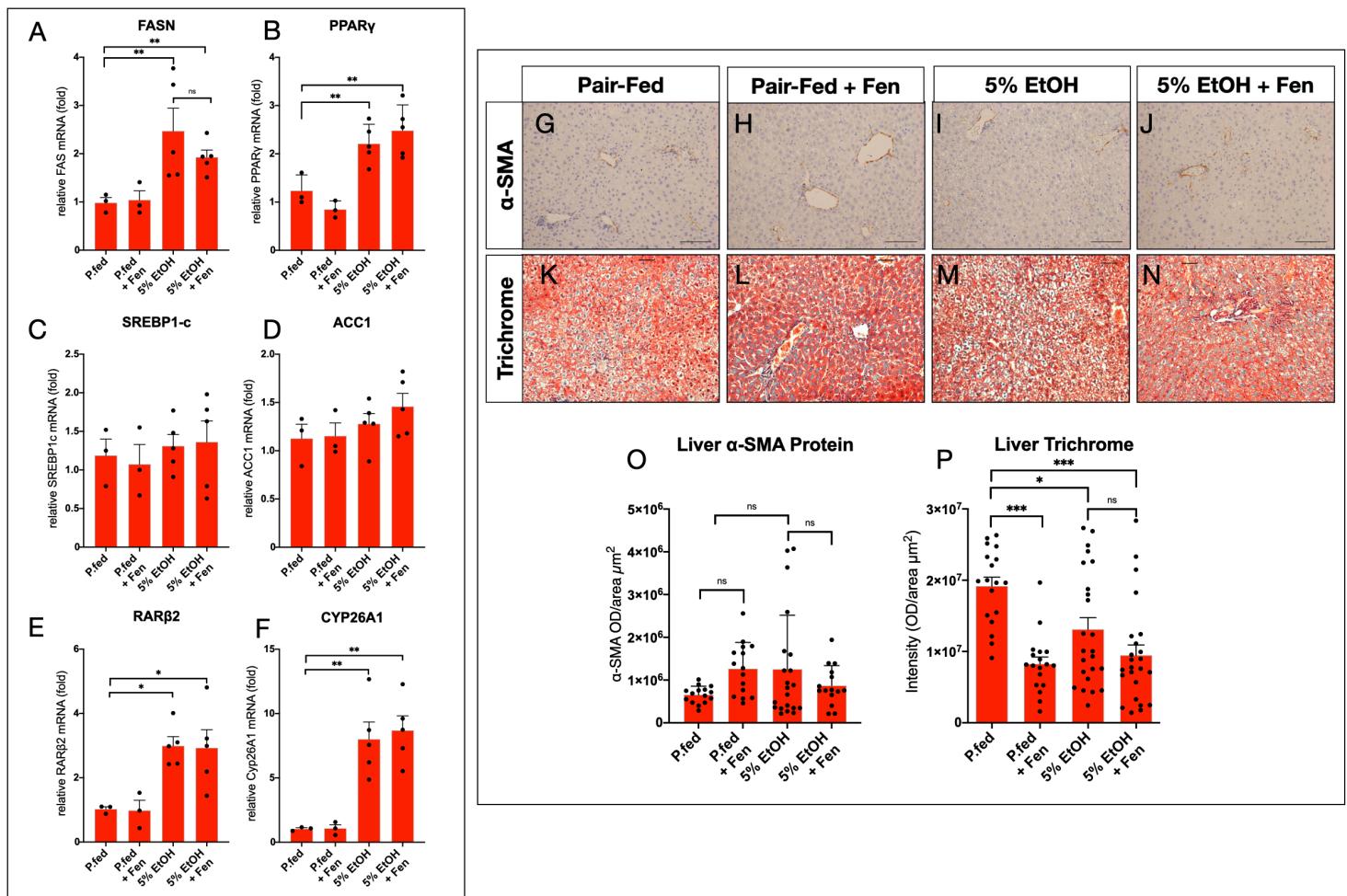


Supplementary Figure 2. A) Ileum mRNA levels of *TLR4*, B) Quantification of ileum immunofluorescence (IF) optical density of F4/80 and F4/80:TLR4 double positive macrophages. Ileum mRNA levels of C) *TNF- α* , and D) *tumor necrosis factor receptor 1* (*TNFR1*). E) Quantification of ileum immunohistochemistry (IHC) for TNF- α protein levels. F-I) Representative double immunofluorescence (IF) images of ileum sections stained with antibodies against F4/80 (green channel) and TLR4 (red channel) with double positive cells (yellow/orange [white arrows]) Magnification: 100X; Scale Bar=50 μm . J-M) Representative immunohistochemistry (IHC) images of ileum sections stained with an antibody against tumor necrosis factor α (TNF- α) Magnification: 100X; Scale Bar=50 μm . All data errors bars represent \pm SD, with *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001, ns=not significant.



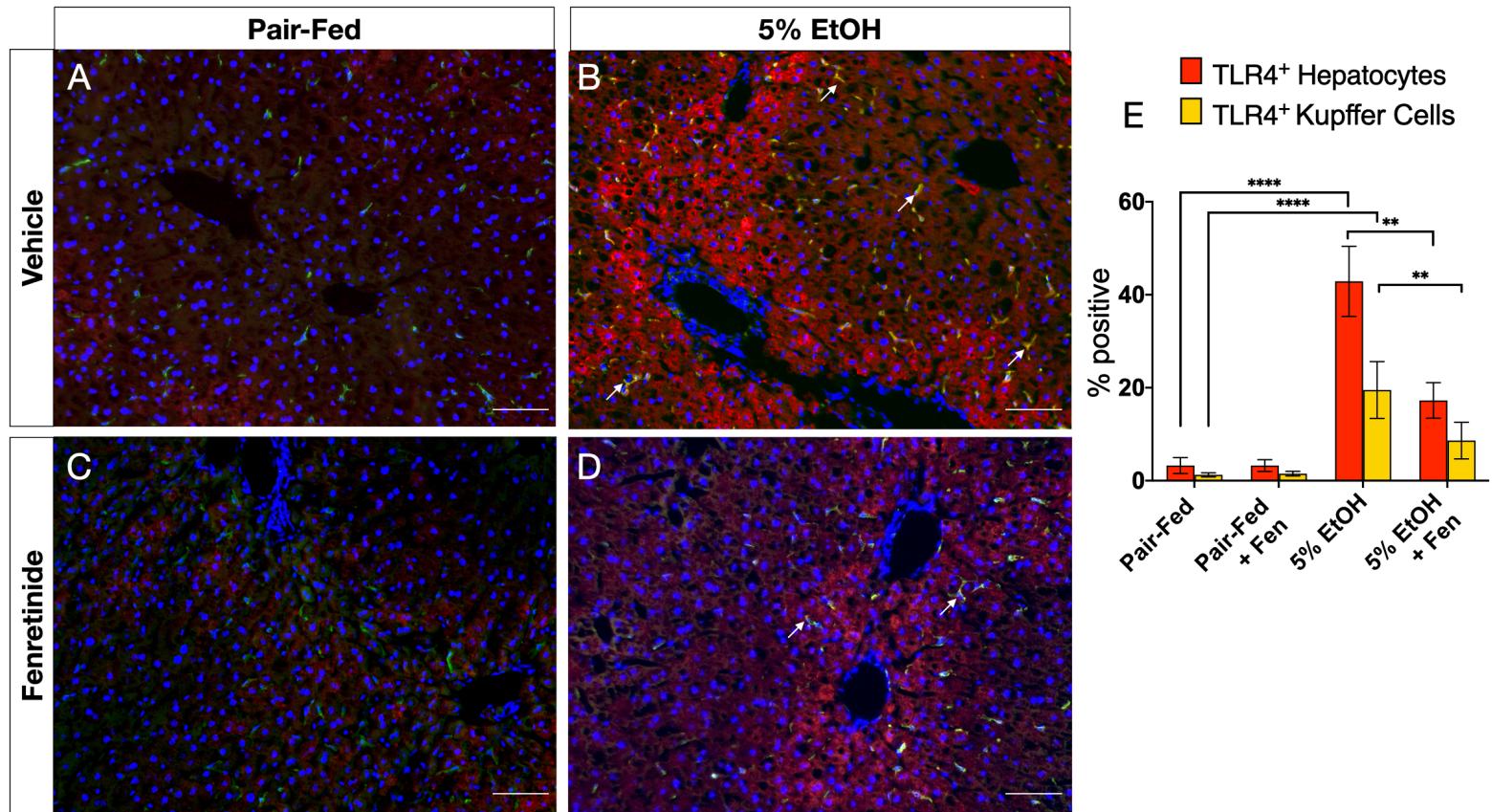
Supplementary Figure 3. A-D) Representative immunohistochemistry (IHC) images of ileum sections stained with an antibody against 4 hydroxyneonenal (4-HNE) Magnification: 100X; Scale Bar=50 μm . E) Quantification of ileum IHC for 4-HNE protein levels. Ileum mRNA levels of Nrf-2 target genes F) Gsta1, and B), Gstm1. All data errors bars represent $\pm \text{SD}$, with * $p<0.05$, ** $p<0.01$, *** $p<0.001$, **** $p<0.0001$.

Supplementary Material



Supplementary Figure 4. Hepatic mRNA levels of lipogenic genes A) *FASN*, B) *PPAR- γ* , C) *SREBP1-c*, D) *ACC1* and retinoid target genes, E) *RAR β 2*, and F) *Cyp26A1*. G-J) Representative immunohistochemistry (IHC) images of liver sections stained with an antibody against smooth muscle actin α (α -SMA). Magnification: 100X; Scale Bar=50 μm . K-N) Representative images of liver sections stained with Masson's trichrome for collagen. Magnification: 100X; Scale Bar=50 μm . Quantification liver M) IHC for α -SMA protein levels and N) Masson's trichrome for collagen. All data errors bars represent \pm SD, with * $p<0.05$, ** $p<0.01$, *** $p<0.001$, ns=not significant.

Liver F4/80/TLR4



Supplementary Figure 5. A-D) Representative double immunofluorescence (IF) images of liver sections stained with antibodies against F4/80 (green channel) and TLR4 (red channel) with double positive cells (yellow/orange [white arrows]) Magnification: 100X; Scale Bar=50 µm. Quantification of liver immunofluorescence (IF) optical density of F4/80 and F4/80:TLR4 double positive macrophages. All data errors bars represent \pm SD, with ** p <0.01, *** p <0.0001.

Gene Name	Forward Primer (5'-3')	Reverse Primer (5'-3')
Pathway: Tight Junction Proteins		
Claudin 1	TGGGTTTCATCCTGGCTTCT	TGTATCTGCCCGGTGCTTT
Occludin	GCAGCCTCTGCTTCATCG	CGTCGGGTTCACTCCCATT
Zona-Ocludins 1 (ZO-1)	TCACGATCTCCTGACCAACG	GGCTGACGGGTAAATCCACA
Cingulin	AGCAGAGCAGCAAGGAACCT	GCTGTAGCTCCTTCACCTGG
Fodrin/Spectrin alpha	CGCATCTTTTCCTCAGCAG	CCAGGACTTGCTGTCGTCTC
Symplekin	CGGAGTGTGGCATCACAGTT	CGCACTCAATGGATTGTCTG
Pathway: Inflammation		
Interleukin-1 β (IL1 β)	GAAATGCCACCTTTGACAGTG	CTGGATGCTCTCATCAGGACA
Interleukin-6 (IL-6)	TAGTCCTCCTACCCCAATTCC	TTGGTCCTTAGCCACTCCTTC
Toll-like receptor 4 (TLR4)	ATGGCATGGCTTACACCACC	GAGGCCAATTTGTCTCCACA
Tumor necrosis factor receptor 1(TNFR1)	CCGGGAGAAGAGGGATAGCTT	TCGGACAGTCACTCACCAAGT
Tumor necrosis factor- α (TNF- α)	CCTGTAGCCCACGTCGTAG	GGGAGTAGACAAGGTACAACCC
Pathway: Oxidative Stress		
Glutathione S-transferase, alpha 1(Gsta1)	AGCCCGTGCTTCACTACTTC	TCTTCAAACCTCCACCCCTGC
Glutathione S-transferase, m1 (Gstm1)	GAGGGCCTCAAGAAGATCTCTG	TTACTCCAGTGGCCATCTTG
Pathway: Retinoid Signaling		
Retinoic acid receptor β (isoform 2) (RAR β 2)	GCGAGAGTTTGATGGAGTTC	TGGTAGCCCGAGACTTGTCCCT
Cytochrome P450 26A1 (CYP26A1)	GAAACATTGCAGATGGTGTTCAG	CGGCTGAAGGCCTGCATAATCAC
Pathway: Lipid Metabolism		
Acetyl-Coenzyme A carboxylase alpha (ACC1)	ATGGGCGGAATGGTCTCTTC	TGGGGACCTTGTCTTCATCAT
Fatty Acid Synthase (FASN)	GGAGGTGGTGTAGCCGGTAT	TGGGTAATCCATAGAGCCCAG
Peroxisome proliferator-activated receptor γ (PPAR γ)	CTCCAAGAACATCAAAGTGCAGA	GCCTGATGCTTATCCCCACA
Sterol regulatory element binding transcription factor 1-c (SREBP1-c)	CGGAAGCTGTCGGGGTAG	GTTGTTGATGAGCTGGAGCA
Pathway: Housekeeping		
36B4	AGAACAAACCCAGCTCTGGAGAAA	ACACCCTCCAGAAAGCGAGAGT