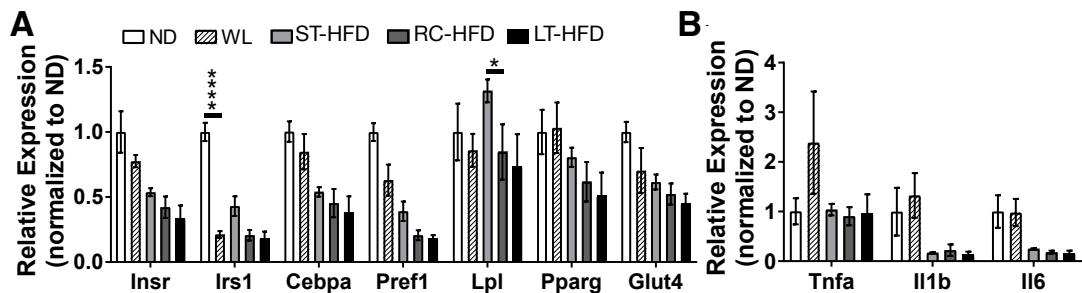
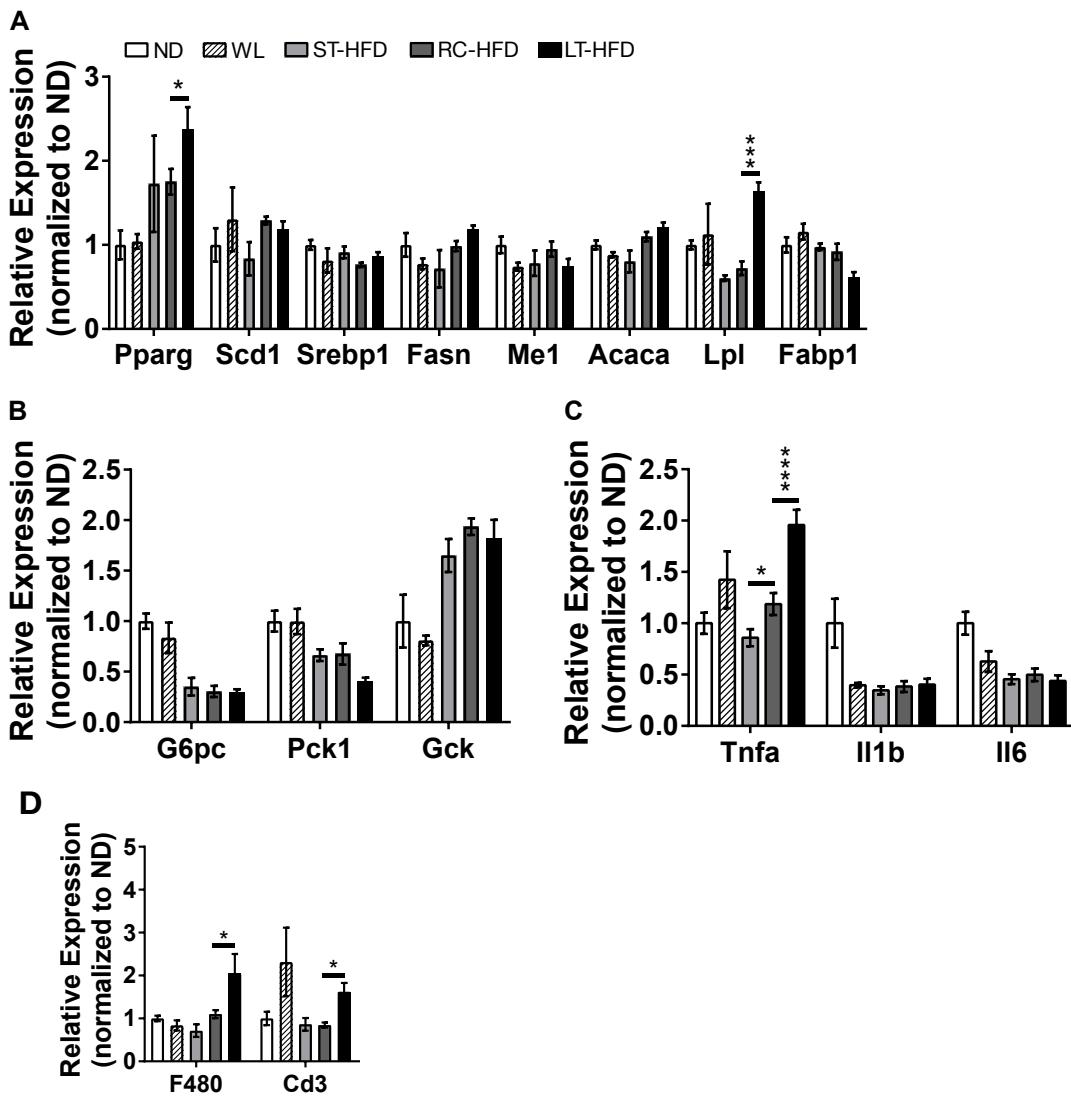


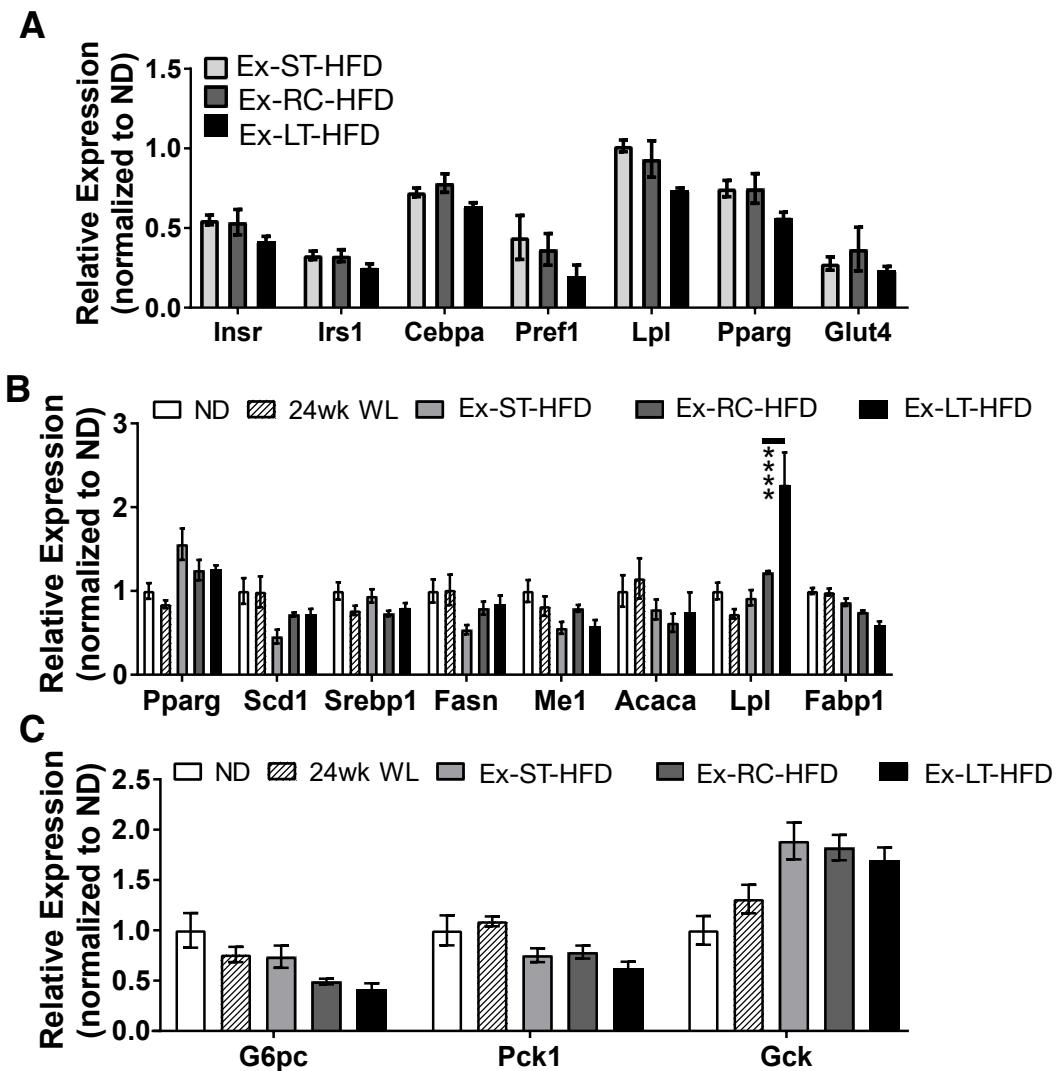
Figure S1



Inguinal adipose tissue gene expression. (A) Adipogenic genes. (B) Inflammatory genes. * p < 0.05, ** p < 0.01, **** p <0.0001, significance was only compared for ST HFD versus RC HFD, LT HFD versus RC HFD or ND versus WL.

Figure S2

Liver lipid metabolism and lipogenesis gene expression is not significantly different between ST HFD and RC HFD mice groups. Gene expression for select liver (A) lipid metabolism, (B) glucose metabolism, (C) inflammatory cytokine genes, and (D) leukocyte genes. * p < 0.05, ** p < 0.001, *** p < 0.0001, significance was only compared for ST HFD versus RC HFD, LT HFD versus RC HFD or ND versus WL.

Figure S3

Adipose tissue and liver gene expression after rechallenge after prolonged period off diet. (A)

Expression of select adipocyte maturation genes from whole eWAT. Gene expression for select liver (B) lipid metabolism and (C) glucose metabolism genes. * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001, significance was only compared for Ex-ST-HFD versus Ex-RC-HFD, Ex-LT-HFD versus Ex-RC-HFD, or ND versus WL.

Table S1 - Primers for qRT-PCR analysis

Gene	Forward Primer	Reverse Primer
Insr	TTTGTCAATGGATGGAGGCTA	CCTCATCTTGGGGTTGAAC
Irs1	CGATGGCTTCTCAGACGTG	CAGCCCGCTTGTGATGTTG
Cebpa	CAAGAACAGCAACGAGTACCG	GTCACTGGTCAACTCCAGCAC
Pref1 (Dlk1)	TTCGGGCTTGCACCTCAA	GGAGCATTCTGACTGGCCTT
Lpl	TGTGTCTTCAGGGTCCTTAG	GGGAGTTGGCTCCAGAGTTT
Pparg	GGAAGACCCTCGCATTCTT	TCGCACTTGGTATTCTGGAG
Akt1	ATGAACGACGTAGCCATTGTG	TTGTAGCCAATAAGGTGCCAT
Glut4	GTGACTGGAACACTGGTCCTA	CCAGCCACGTTGCATTGTAG
Tnfa	ACGGCATGGATCTCAAAGAC	AGATAGCAAATCGGCTGACG
Il1b	AAATACCTGTGGCCTTGGC	CTTGGGATCCACACTCTCCAG
Il6	AGTGAGGAACAAGCCAGAGC	CATTGTGGTTGGTCAGG
Scd1	GCTGGAGTACGTCTGGAGGAA	TCCCAGAGAGGCAGGTGTAG
Srebp1	GATGTGCGAACTGGACACAG	CATAGGGGGCGTAAACAG
Fasn	GGAGGTGGTGATAGCCGGTAT	TGGGTAATCCATAGAGCCCAG
Me1	GTCGTGCATCTCACAGAAG	TGAGGGCAGTTGGTTTATCTT
Acaca	TAATGGGCTGCTCTGTGACTC	CTCAATATGCCATCAGCTTG
Fabp1	ATGAACTTCTCCGGCAAGTACC	CTGACACCCCCCTGATGTCC
G6pc	CGACTCGCTATCTCAAGTGA	GTTGAACCAGTCTCCGACCA
Pck1	CTGCATAACGGTCTGGACTTC	CAGCAACTGCCGTACTCC
Gck	ATGGCTGTGGATACTACAAGGA	TTCAGGCCACGGTCCATCT
Fabp4	AAGGTGAAGAGCATCATAACCCT	TCACGCCTTCATAACACATTCC