

Supplementary Materials: Hepatic Fasting-Induced PPAR α Activity Does Not Depend on Essential Fatty Acids

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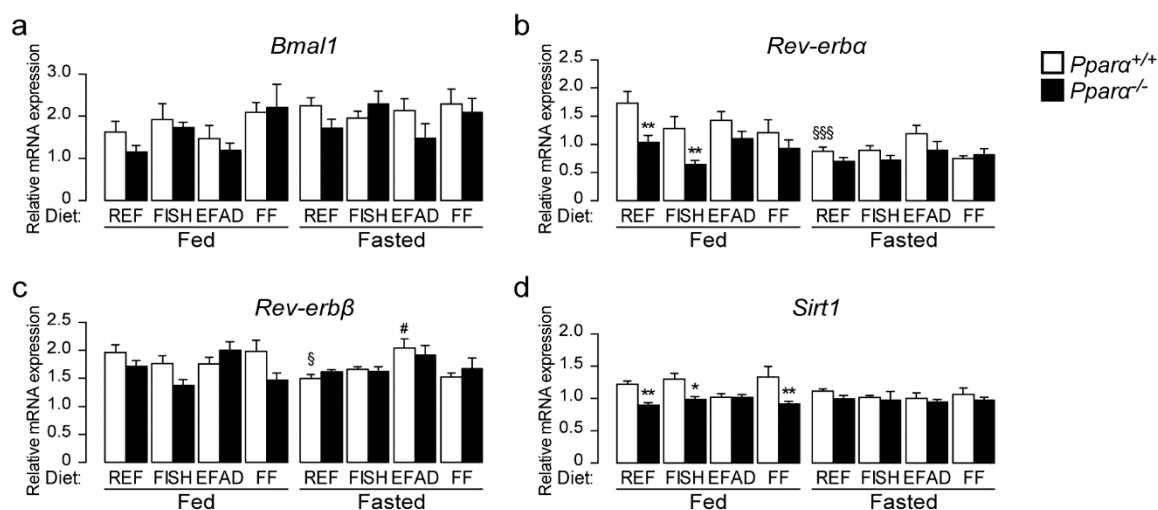


Figure S1. Effect of dietary fat on hepatic gene expression. Wild-type (*Ppara*^{+/+}) and total *Ppara* knockout (*Ppara*^{-/-}) mice were fed ad libitum or fasted for 24 h and then euthanized at ZT14. Quantification of *Bmal1* (a); *Rev-erba* (b); *Rev-erbβ* (c); and *Sirt1* (d) mRNA expression levels in the liver as determined by qRT-PCR. Data were normalized to the expression of TATA-binding protein (TBP). Data represent mean \pm SEM. * significant genotype effect, # significant effect of diet composition, § significant effect of fasting. *, #, § $p \leq 0.05$. ** $p \leq 0.01$. §§§ $p \leq 0.001$. REF: standard diet; FISH: diet enriched in n-3 essential fatty acids from fish oil; EFAD: essential fatty acid-deficient diet; FF: fat-free diet.

Table S1. Liver composition in FA from liver of wild-type (*Ppara*^{+/+}) and total *Ppara* knockout (*Ppara*^{-/-}) mice were fed ad libitum or fasted for 24 h and killed at ZT14. Transmethylated fatty acids from liver lipid extract were quantified by gas chromatography. Data shown as mean ± SEM. * significant genotype effect, # significant effect of diet composition, § significant effect of fasting. *, #, § $p \leq 0.01$, **, ##, §§ $p \leq 0.001$. REF: standard diet; FISH: diet enriched in n-3 essential fatty acids from fish oil; EFAD: essential fatty acids deficient diet; FF: fat free diet.

	Fed							
	REF		FISH		EFAD		FF	
	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}
C14:0	0.19 ± 0.044	0.51 ± 0.029 **	0.02 ± 0.008 ##	0.35 ± 0.082 **	0.61 ± 0.11 ##	1.99 ± 0.07 **,#	0.12 ± 0.022	0.61 ± 0.039 **
C16:0	19.9 ± 0.99	21.8 ± 0.85	18.4 ± 1.01	24.9 ± 1.76 **	20.9 ± 0.56	23.1 ± 0.72	18 ± 0.6	23.1 ± 0.86 **
C16:1 n-9	0.85 ± 0.049	1.19 ± 0.121	0.99 ± 0.069	1.09 ± 0.147	0.93 ± 0.156	1.28 ± 0.069	1.36 ± 0.082	1.56 ± 0.205
C16:1 n-7	2.1 ± 0.18	4.5 ± 0.32 **	1.8 ± 0.43	3.8 ± 0.73 **	4.6 ± 0.31 ##	10.5 ± 0.48 **,#	4.1 ± 0.23 ##	8.7 ± 0.78 **,#
C18:0	10.2 ± 0.43	4.4 ± 0.57 **	10.7 ± 0.98	8.3 ± 1.84	7.3 ± 0.68	2.7 ± 0.3 **	7.3 ± 0.64	5.5 ± 1.61
C18:1 n-9	23.2 ± 1.27	34.9 ± 0.9 **	18.9 ± 2.38	26.4 ± 3.51 *,##	34.5 ± 1.71 ##	38.1 ± 0.76	38.9 ± 1.58 ##	38.7 ± 1.47
C18:1 n-7	4.6 ± 0.29	6.4 ± 0.31 *	2.8 ± 0.34 ##	3.7 ± 0.45 ##	10.3 ± 0.43 ##	14.8 ± 0.52 **,#	11.5 ± 0.26 ##	16.5 ± 0.88 *,##
C18:2 n-6	13 ± 0.34	17.4 ± 1.07	11.3 ± 0.8	15.4 ± 1.09	3.5 ± 0.37 ##	2.4 ± 0.26 ##	2.1 ± 0.3 ##	2 ± 0.3 ##
C18:3 n-6	0.26 ± 0.014	0.15 ± 0.027 *	0.1 ± 0.018 ##	0.06 ± 0.004 ##	0.07 ± 0.01 ##	0.03 ± 0.006 **,#	0.04 ± 0.006 ##	0.02 ± 0.004 ##
C18:3 n-3	0.21 ± 0.021	0.67 ± 0.042 **	0.33 ± 0.017 #	0.95 ± 0.104 **	0.08 ± 0.017 ##	0.13 ± 0.01 **,#	0.2 ± 0.03	0.11 ± 0.011 *,##
C20:0	0.74 ± 0.171	0.81 ± 0.161	0.26 ± 0.054	0.28 ± 0.027	0.81 ± 0.184	0.77 ± 0.174	0.06 ± 0.009 #	0.02 ± 0.002 ##
C20:1 n-9	0.41 ± 0.084	0.55 ± 0.058 *	0.55 ± 0.034	0.71 ± 0.055	0.93 ± 0.068 ##	0.92 ± 0.088 #	1.41 ± 0.1 ##	0.78 ± 0.085 *
C20:2 n-6	0.47 ± 0.048	0.32 ± 0.042	0.34 ± 0.03	0.21 ± 0.035	2.88 ± 0.424 ##	0.41 ± 0.074 **	4.34 ± 0.415 ##	0.23 ± 0.041 **
C20:3 n-9	0.0081 ± 0.0017	0.0094 ± 0.0024	0.0046 ± 0.0016	0.0012 ± 0.0005	0.11 ± 0.016 ##	0.032 ± 0.0064 #	0.1506 ± 0.0126 ##	0.0097 ± 0.0016 *
C20:3 n-6	1.68 ± 0.081	0.41 ± 0.064 **	1.46 ± 0.119	0.33 ± 0.109 **	1.09 ± 0.144 #	0.19 ± 0.031 **,#	0.66 ± 0.104 ##	0.13 ± 0.02 **,#
C20:4 n-6	13.9 ± 0.71	3.2 ± 0.65 **	6.3 ± 0.8 ##	1.7 ± 0.99 **,#	6.9 ± 0.77 ##	1.4 ± 0.2 **,#	5.6 ± 0.64 ##	1 ± 0.2 **,#
C20:5 n-3	0.1 ± 0.011	0.09 ± 0.014	4.49 ± 0.509 ##	2.47 ± 0.603 ##	0.03 ± 0.007 ##	0.01 ± 0.003 ##	0.01 ± 0.006 ##	0.01 ± 0.003 ##
C22:4 n-6	0.48 ± 0.063	0.18 ± 0.04 **	0.09 ± 0.015 ##	0.07 ± 0.014 #	0.15 ± 0.028 ##	0.03 ± 0.005 **,#	0.09 ± 0.016 ##	0.02 ± 0.006 **,#
C22:5 n-6	0.76 ± 0.195	0.13 ± 0.031 **	0.08 ± 0.016 ##	0.05 ± 0.007 #	0.85 ± 0.178	0.19 ± 0.019 **	0.87 ± 0.07	0.12 ± 0.028 **
C22:5 n-3	0.23 ± 0.013	0.12 ± 0.021 *	1.92 ± 0.082 ##	1.94 ± 0.311 ##	0.04 ± 0.008 ##	0.01 ± 0.002 **,#	0.01 ± 0.006 ##	0 ± 0.001 ##
C22:6 n-3	6.3 ± 0.41	1.7 ± 0.37 **	18.7 ± 1.75 ##	6.9 ± 2.28 **,#	2.9 ± 0.3 ##	0.6 ± 0.09 **,#	2.7 ± 0.32 ##	0.4 ± 0.09 **,#

Table S2. Cont.

	Fasted							
	REF		FISH		EFAD		FF	
	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}	<i>Ppara</i> ^{+/+}	<i>Ppara</i> ^{-/-}
C14:0	0.35 ± 0.043 §§	0.53 ± 0.077	0.36 ± 0.033 §§	0.55 ± 0.027	1.3 ± 0.084 ##§	1.96 ± 0.126 ##	0.47 ± 0.052 §§	0.83 ± 0.037
C16:0	20.8 ± 0.62	18 ± 0.78 §	22.7 ± 0.57 §	25.4 ± 1.15 ##	20.7 ± 0.43	19.9 ± 0.21	20.4 ± 0.29	20.3 ± 1.1
C16:1 n-9	1.14 ± 0.138	1.32 ± 0.113	1.13 ± 0.068	1.13 ± 0.132	1.26 ± 0.063	1.1 ± 0.115	1.53 ± 0.121	1.74 ± 0.12
C16:1 n-7	4.6 ± 0.34 §§	7.2 ± 0.52 *§	4.4 ± 0.08 §§	4.6 ± 0.28 #	11.7 ± 0.44 ##,§§	17.4 ± 1.02 *,##,§§	8.3 ± 0.68 ##,§§	11.6 ± 1.68 #
C18:0	3.8 ± 0.3 §§	3.5 ± 0.57	3.9 ± 0.21 §	5.7 ± 1.27	3.4 ± 0.17 §	2.9 ± 0.65	4.4 ± 0.81	5.7 ± 2.38
C18:1 n-9	34.6 ± 1.46 §§	36.5 ± 0.71	23.9 ± 0.87 ##	29.4 ± 2.02	40.5 ± 0.52	38.8 ± 0.72	41.3 ± 1.23	40.5 ± 1.66
C18:1 n-7	3.4 ± 0.22 §	4.9 ± 0.4 *	1.9 ± 0.25 ##,§	3.1 ± 0.36 *,##	6.1 ± 0.48 ##,§§	11.5 ± 0.98 **,##	6.5 ± 0.36 ##,§§	14.5 ± 0.54 *,##
C18:2 n-6	18 ± 1.11	21.8 ± 1.08	18.9 ± 0.73 §	16.3 ± 0.91	3.8 ± 0.27 ##	2.8 ± 0.22 ##	3.8 ± 0.3 ##,§	2.4 ± 0.72 *,##
C18:3 n-6	0.78 ± 0.125 §§	0.46 ± 0.05 §§	0.28 ± 0.046 ##,§§	0.1 ± 0.017 **,##	0.39 ± 0.046 ##,§§	0.18 ± 0.02 **,##,§§	0.31 ± 0.037 ##,§§	0.12 ± 0.037 **,##,§§
C18:3 n-3	0.55 ± 0.038 §§	1.02 ± 0.064 **,§	0.88 ± 0.056 ##,§§	1.14 ± 0.067	0.07 ± 0.007 ##	0.11 ± 0.013*,##	0.09 ± 0.007 ##,§§	0.12 ± 0.007##
C20:0	0.37 ± 0.1	0.47 ± 0.115	0.05 ± 0.004	0.35 ± 0.066	0.49 ± 0.151	0.42 ± 0.107	0.02 ± 0.005 #	0.01 ± 0.003 #
C20:1 n-9	0.25 ± 0.028	0.37 ± 0.039	0.31 ± 0.034 §	0.67 ± 0.052 *,#	0.32 ± 0.014 §§	0.62 ± 0.038 *,#	0.45 ± 0.072 #,§§	0.59 ± 0.083
C20:2 n-6	0.24 ± 0.01 §§	0.22 ± 0.017	0.12 ± 0.012 #,§§	0.15 ± 0.014	1.95 ± 0.176 ##	0.27 ± 0.02 **	2.16 ± 0.338 ##,§	0.16 ± 0.011 **
C20:3 n-9	0.002 ± 0.0003	0.0039 ± 0.0009	0.0021 ± 0.0005	0.001 ± 0.0006	0.0576 ± 0.0153 ##	0.0129 ± 0.0019	0.0508 ± 0.0192 ##	0.0042 ± 0.0007
C20:3 n-6	0.39 ± 0.021 §§	0.17 ± 0.012 **,§§	0.41 ± 0.042 §§	0.18 ± 0.016 **	0.42 ± 0.033 §§	0.06 ± 0.004 **,##,§§	0.38 ± 0.069 §	0.03 ± 0.004 *,##,§§
C20:4 n-6	5.7 ± 0.41 §§	1.6 ± 0.14 **	2.6 ± 0.14 ##,§	0.7 ± 0.05 *,##	4 ± 0.14	0.9 ± 0.05 **,#	5.6 ± 0.77	0.6 ± 0.11 **,##
C20:5 n-3	0.15 ± 0.012	0.08 ± 0.007	3.46 ± 0.152 ##	2.27 ± 0.366 ##	0.02 ± 0.002 ##	0.01 ± 0.001 ##	0.02 ± 0.002 ##	0 ± 0.001 *,##
C22:4 n-6	0.19 ± 0.014 §	0.13 ± 0.018	0.04 ± 0.004 ##	0.05 ± 0.007	0.14 ± 0.036	0.02 ± 0.004 **,##	0.08 ± 0.011 #	0.01 ± 0.005 **,##
C22:5 n-6	0.2 ± 0.02 §§	0.08 ± 0.018 **	0.06 ± 0.006 ##	0.05 ± 0.006	0.73 ± 0.146 ##	0.1 ± 0.006 **	0.59 ± 0.104 ##	0.07 ± 0.007 **
C22:5 n-3	0.26 ± 0.025	0.11 ± 0.012 *	1.65 ± 0.094 ##	1.94 ± 0.332 ##	0.09 ± 0.013 ##,§	0.01 ± 0.001 **,##	0.07 ± 0.001 ##,§§	0.01 ± 0.003 **,##
C22:6 n-3	3.8 ± 0.26	1 ± 0.09 **	12.5 ± 0.7 ##	5.7 ± 0.89 *,##	2.1 ± 0.09 #	0.3 ± 0.02 **,##	3.1 ± 0.39	0.2 ± 0.03 **,##

Table S2. Diet compositions.

% (w/w)	Fat Free Diet (FF)	CTRL, FISH, EFAD Diets
Cellulose	2	2
Casein	22	22
Starch	46.2	43.5
Methionin	0.2	0.2
Sucrose	23.1	21.8
Minerals	4.5	4.5
Vitamins	1	1
Oil	0	5
Agar-Agar	1	0

Table S3. Relative fatty acid composition of dietary oils.

Diet	REF	EFAD	FISH
12:00	0	47.7	0
14:00	0	23	0
16:00	5.9	13.1	4.4
18:00	2.8	15.3	2.1
18:1w9	39.7	0.9	30.5
18:2w6	47.0	0	34.7
18:3w3	4.6	0	3.4
20:5w3	0	0	5.3
22:6w3	0	0	19.5

Table S4. Oligonucleotide sequences for the real time PCR. Oligonucleotides (Sigma Aldrich, St. Louis, MO, USA) were designed with Primer Express 2.0 software (Applied Biosystems, Foster City, CA, USA).

Gene	NCBI Refseq	Forward Primer (5'-3')	Reverse Primer (5'-3')
<i>Acly</i>	NM_134037	AAAGCTTGGCCTCGTCGG	GGGACGAAGGGTTCAATGAGA
<i>Acox1</i>	NM_015729	CAGACCCTGAAGAAATCATGTGG	CAGGAACATGCCAAAGTGAAG
<i>Alas1</i>	NM_020559	CAAAGAAACCCCTCCAGCC	GCTGTGTGCCGTCTGGAGT
<i>Bien</i>	NM_023737	CGTCTCCTCGGTGTGGTTC	ATTATCTTCTTGCAGTATCTAGCTGCTT
<i>Bmal1</i>	NM_007489	CAAACACTACAAGCCAACATTTCATCAG	TCGGTCACATCCTACGACAAAC
<i>Cd36</i>	NM_007643	GITAAACAAAGAGGTCCTTACACATACAG	CAGTGAAGGCTCAAAGATGGC
<i>Chrebp</i>	NM_021455	ACTCAGGGAATACAGCCTACAG	GAAGAAGGAATTCAGAGCTCAGAAA
<i>Cpt1a</i>	NM_013495	GAAGAAGAAGTTCATCCGATTCAAG	GATATCACACCCACCACCAG
<i>Cyp4a10</i>	NM_010011	TCCAGCAGTTCATCACCT	TTGCTTCCCCAGAACCATCT
<i>Cyp4a14</i>	NM_007822	TCAGTCTATTCTGGTGCTGTC	GAGCTCCTTGCTTCAGATGGT
<i>Eci</i>	NM_010023	GTTCCACATCAGCCTGGAGAAG	AGAAGATACCCGGGCATCC
<i>Fasn</i>	NM_007988	AGTCAGCTATGAAGCAATTGTGGA	CACCCAGACGCCAGTGTTC
<i>Fgf21</i>	NM_020013	AAAGCCTCTAGGTTCTTTGCCA	CCTCAGGATCAAAGTGAGGGC
<i>Fsp27</i>	NM_178373	CATGAAGTCTCTCAGCCTCTGTA	CAGCTGTTGGGTCACCACTG
<i>Hmgcr</i>	NM_008255	CTGTGGAAATGCCTTGTGATTG	GAAGAATGTCATGAACACAAAGTAGTGG
<i>Hmgcs2</i>	NM_008256	TGCAGGAAACTTCGCTCACA	AAATAGACCTCCAGGGCAAGGA
<i>Insig2</i>	NM_133748	TGTATATTTTTTGTGGAGGCATAAC	TTCAGCAATAACTTTGCATTCATACAT
<i>Lpk</i>	NM_013631	TCGACTCAGAGCCTGTGGC	AGTCGTGCAATGTTTCATCCT
<i>Lxra</i>	NM_013839	GGAGTGTGCACTTCGCAAATG	TCAAGCGGATCTGTTCTTCTGAC
<i>Lxrb</i>	NM_009473	ATGCGGGAGCAGTGCCG	CCGGCCTGAGCTGCTG
<i>Me1</i>	NM_008615	CATTTCGAGGCGTTTCGTTG	CAGGTAGGATCTGGTCATAATTAGTGC
<i>Pdk4</i>	NM_013743	ATCGCCAGAATTAACCTCACAC	TGGATTGGTTGGCCTGGA
<i>Pepck</i>	NM_011044	GAACCCAGCCTGCCC	GAGCAACTCCAAAAAACCCG
<i>Plin2</i>	NM_007408	CCATTTCTCAGTCCACTCCAC	GTGTCGTAGCCGATGC
<i>Pnpla2</i>	NM_001163689	AGTGTCTTACCATCCGCTT	GGATATCTTCAGGACATACAGGC
<i>Pnpla3</i>	NM_054088	ACGCGGTACCTTCGTGT	AGCCCGTCTCTGATGCACTT
<i>Ppara</i>	NM_011144	CCCTGTTTGTGGCTGCTATAATTT	GGGAAGAGGAAGGTGTCATCTG
<i>Pparb</i>	NM_011145	AAGTGGCCATGGGTGACG	TGGTCCAGCAGGGAGGAAG
<i>Pparg2</i>	NM_011146	ATGGGTGAAACTCTGGGAGATTCT	CTTGGAGCTTCAGGTCATATTGTA
<i>Reverba</i>	NM_145434	CAGCTGGTGAAGACATGACGAC	GGAGGAGCCACTAGAGCCAA
<i>Reverbb</i>	NM_011584	CGCCATGGAGCTGAACG	GACAAGAGGCAGGGCTGGA
<i>Scd1</i>	NM_009127	CAGTGCCGCGCATCTCTAT	CTGACTGGCAAATATAGCTGTATCCT
<i>Spot14</i>	NM_009381	AACGGAGGAGGCCGAAGAAG	GTTGATGCACCTCGGGTCT
<i>Tbp</i>	NM_013684	ACTTCTGTCAAGAAATGCTGAA	GCAGTTGTCCGTTGGCTCTCT
<i>Srebp1c</i>	NM_011480	GGAGCCATGGATTGCACATT	GCTTCCAGAGAGGAGGCCAG
<i>Thiolb</i>	NM_146230	ACATCTCCGTGGCAATGTG	TGTCGTGACCGTAGACAAAGGT