

Supplementary data and methods for:

Elevated Fibroblast growth factor 21 (FGF21) in obese, insulin resistant states is normalised by the synthetic retinoid Fenretinide in mice

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Supplementary table S1

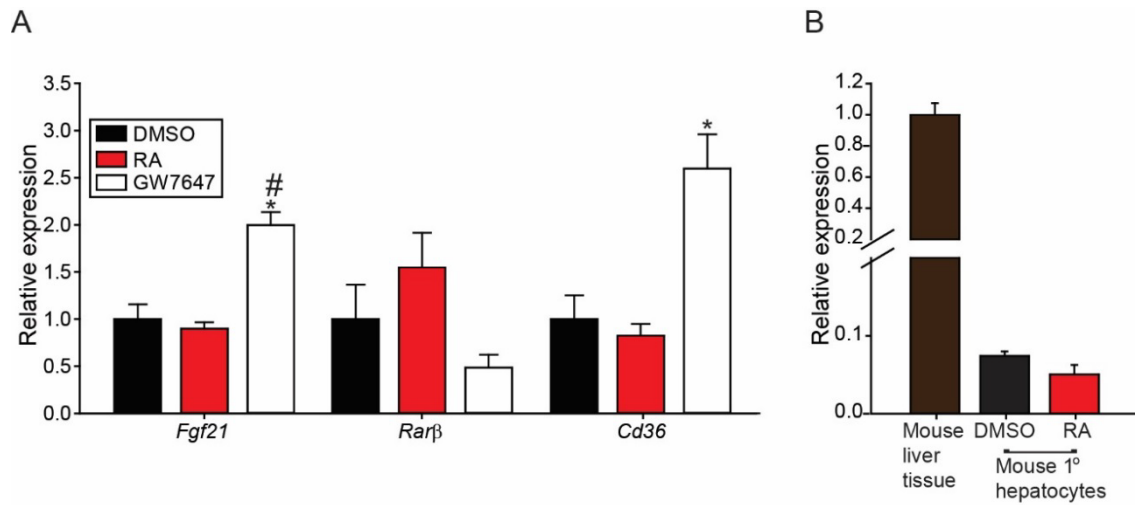
Measurement	Chow	HFD	FEN-HFD
20 weeks diet			
Final body weight (g)	31.8 ± 0.6	45.2 ± 0.87 *	39.8 ± 1.5 *#
Adiposity at 12 weeks (fat mass in g)	4.6 ± 0.2	13.4 ± 0.5 *	10.3 ± 0.9 *#
Serum glucose at 12 weeks (mmol/L)	11.9 ± 0.7	20.9 ± 0.2 *	11.6 ± 0.6 #
Serum insulin at 20 weeks (ng/mL)	0.7 ± 0.01	2.8 ± 0.3 *	1.6 ± 0.2 *
Serum leptin at 20 weeks	1.0 ± 0.3	1.5 ± 0.3 *	1.6 ± 0.2 *#
Food intake from chronic cumulative study (kcal/ day)	-	14.2 ± 0.2	13.8 ± 0.3
7 days diet			
Final body weight (g)	24.7 ± 0.7	26.2 ± 0.2	25.6 ± 0.3
Serum leptin	1.0 ± 0.2	4.6 ± 0.8 *	2.4 ± 0.5 #
Food intake from acute study (average change in energy intake from baseline in kcal/day)	- 0.3 ± 0.2	2.2 ± 0.8 *	1.8 ± 0.4 *

Supplementary table S1: Metabolic parameters measured in mice fed HFD for 7 days or 20 weeks

Data was previously published in McIlroy GD et al, Fenretinide Treatment Prevents Diet-Induced Obesity in Association With Major Alterations in Retinoid Homeostatic Gene Expression in Adipose, Liver and Hypothalamus, *Diabetes*, 2013 Mar; 62(3): 825- 836.

Data are shown as mean ± SEM and significance was determined by one-way ANOVA followed by *post-hoc* tests. Differences are marked * p<0.05 vs chow; #p<0.05 vs FEN-HFD.

Supplementary figure S1



Supplementary figure S1: Downregulation of retinoid signalling in mouse primary hepatocytes prevents the inhibition of *Fgf21* expression by RA

A: qPCR analysis of gene expression in mouse primary hepatocytes treated overnight with DMSO followed by DMSO (black bars) or 1 μ M RA (red bars) the following day for 60 min or 1 μ M GW7647 overnight followed by 60 min DMSO treatment (white bars); n = 3 per treatment. Gene expression was normalised to *yWhaz*.

Significant differences were determined by student's t-test. Differences are shown as * p<0.05 vs DMSO + DMSO; # p<0.05 vs DMSO + RA.

B: Semi-quantitative analysis of *Rara* gene expression in mouse primary hepatocytes treated overnight with DMSO followed by DMSO (black bar) or 1 μ M RA (red bar) the for 60 min (gene expression for both normalised to *yWhaz*), compared to gene expression levels in liver from chow-fed lean control mice (n=10; gene expression normalised to *Nono*).

Supplementary table S2

Gene	Forward	Reverse
<i>18s</i>	CGTTCCTTACCTGGTTGAT	GAGCGACCAAAGGAACCATTA
<i>Adiponectin</i>	TGTTCCCTCTTAATCCTGCCCA	CCAACCTGCACAAGTTCCCTT
<i>Beta-actin</i>	GATCTGGCACCACACACCTTC	GGGGTGTGGAAGGTCTCAA
<i>Beta- klotho</i>	TGTTCTGCTGCGAGCTGTTAC	CCGGACTCACGTAAGTGTTC
<i>C/ebp-alpha</i>	CAAGAACAGCAACGAGTACCG	GTCACTGGTCAACTCCAGCAC
<i>Cd11b</i>	CCCCACACTAGCATCAAGGG	GAGGCAAGGGACACATGAC
<i>Cd36</i>	GAACCACTGCTTTCAAAAAGTGG	TGCTGTTCTTTGCCACGTCA
<i>Chrebp1</i>	AGATGGAGAACCGACGTATCA	ACTGAGCGTGCTGACAAGTC
<i>Crbp1</i>	CTGAGCAATGAGAATTTTCGAGGA	GCGGTCTGTCTATGCCTGTC
<i>Cyp26a1</i>	TTCGGGTTGCTCTGAAGACT	TCCTCCAAATGGAATGAAGC
<i>Cyp26a1 RARE</i>	CCGAATTAAGATGAACCTTTG	GCACGCTTCAGCCTCCCGCGCC
<i>F4/80</i>	CCCAGCTTATGCCACCTGCA	TCCAGGCCCTGGAACATTGG
<i>Fas</i>	AAGCTCAGTGTGCCACCTA	ATGGCAACGTGACACTGCTG
<i>Fgf21</i>	ACCTGGAGATCAGGGAGGAT	CACCCAGGATTTGAATGACC
<i>Fgf21 RARE</i>	CCTCAGACCCAAGAGCCAGA	AAGGAGGAGGCTGGGGTCTA
<i>Fgfr1</i>	GGTGAACGGGAGTAAGATCGG	CCCCGCATCCTCAAAGGAG
<i>Glut 4</i>	GTGACTGGAACACTGGTCCTA	CCAGCCACGTTGCATTGTAG
<i>Hprt</i>	GTTAAGCAGTACAGCCCCAAA	AGGGCATATCCAACAACAACTT
<i>Insulin receptor</i>	AGACCAACTGTCCTGCCACT	ACACACTTGGTGGGGTTCATC
<i>Leptin</i>	GAGACCCCTGTGTCCGGTTC	CTGCGTGTGTGAAATGTCATTG
<i>Lrat</i>	CCGTCCCTATGAAATCAGCTC	ATGGGCGACACGGTTTTTCC
<i>Mc4r</i>	CCCGGACGGAGGATGCTAT	TCGCCACGATCACTAGAATGT
<i>NoNo</i>	GCCAGAATGAAGGCTTGACTAT	TATCAGGGGGGAAGATTGCCCA
<i>Npy</i>	ATGCTAGGTAACAAGCGAATGG	TGTCGCAGAGCGGAGTAGTAT
<i>Pepck</i>	GAGATAGCGGCACAAT	TTCAGAGACTATGCGGTG
<i>Pomc</i>	GTGCCAGGACCTCACCACGG	CGTTGCCAGGAAACACGGGC
<i>Ppara</i>	ACGATGCTGTCTCCTTGATG	GTGTGATAAAGCCATTGCCGT
<i>Pparγ</i>	AGTGGAGACCGCCAGG	GCAGCAGGTTGTCTTGATGT
<i>Ptp1b</i>	GGAAGTGGGCGGCTATTTACC	CAAAGGGGCTGACATCTCGGT
<i>Rarβ</i>	CGCGAGCCCTTCTCCTGC	AAAAGCCCTTGCACCCCTCGC
<i>Rary</i>	GGAGCAGGCTTCCCATTG	CATGGCTTATAGACCCGAGGA
<i>Raldh1</i>	GGTGGAGGACGCTGGGGGAA	CCGAAGGGGCACTGGGCTGA
<i>Raldh2</i>	CAAGGAGGCTGGCTTTCCACCC	GGGCTCTTCCCTCCGAGTTCCA
<i>Rbp4</i>	ACGAGTCCGTCTTCTGAGCAACTG	GCACAGCTCCTCCTGCCGTT
<i>Resistin</i>	AAGAACCCTTTCATTTCCCTCCT	GTCCAGCAATTTAAGCCAATGTT
<i>Scd1</i>	TTCTTGCGATACACTCTGGTGC	CGGGATTGAATGTTCTTGTCTG
<i>Stat3</i>	CAATACCATTGACCTGCCGAT	GAGCGACTCAAAGTCCCT
<i>Tgfβ1</i>	AGCCCGAAGCGGACTACTAT	CTGTGTGAGATGTCTTTGGTTTTT
<i>yWhaz</i>	GAAAAGTTCTTGATCCCCAATGC	TGTGACTGGTCCACAATTCCTT

Supplementary table S2: Mouse primer sequences used in qPCR reactions.

Supplementary methods

Mouse primary hepatocyte cell culture

Mouse primary hepatocytes were extracted from a C57BL/6 wild-type mouse using a modified version of the collagenase method¹. Tissue was perfused via the inferior vena cava with 137mM NaCl, 7mM KCl, 0.7mM Na₂HPO₄, 10mM Hepes, 0.5mM EDTA, pH 7.65 (perfusion buffer) followed by 137mM NaCl, 7mM KCl, 0.7mM Na₂HPO₄, 10mM Hepes, pH 7.65 (wash buffer) and finally with perfusion buffer plus added 5.1mM CaCl₂ and 20mg collagenase (from *C.histolyticum*). All perfusions were performed at 5ml/min flow rate. Cells were strained and seeded at a density of 2.5×10^5 cells/ well (6 well plate) in M199 + glutamax, with added 1% (v/v) Pen-strep, 0.1% (v/v) BSA, 10% (v/v) FCS, 10nM insulin Actrapid, 200nM 3,3',5-triiodo-L-thyronine and 500nM dexamethasone, and incubated at 37°C, 5% CO₂. After 4 hours, media was replenished and supplemented with compounds as per appropriate figure legends. The following day, plating media was replaced with M199+ glutamax with added 1% (v/v) Pen-strep and 100nM dexamethasone, supplemented with compounds as per respective figure legends.

References

1. Berry, M. N. & Friend, D. S. High-yield preparation of isolated rat liver parenchymal cells: a biochemical and fine structural study. *J. Cell Biol.* **43**, 506-520 (1969).