

Supplementary data

Supplementary Table 1: Composition of the diets.

	C1000	Cholesterol	Cholesterol + Formononetin	Cholesterol + C7F
Energy, kcal/kg	3518	3444	3444	3444
Crude protein, g/kg	176	176	176	176
Crude fat, g/kg	50.8	66.6	66.6	66.6
Crude fibre, g/kg	40.5	31.0	31.0	31.0
Disaccharides, g/kg	111	111	111	111
Polysaccharides, g/kg	472	454	454	454
Cholesterol, g/kg	0	20	20	20
Formononetin, mg/kg	0	0	1000	0
C7F, mg/kg	0	0	0	1300

Supplemental Table 2: A list of functional gene groupings included in the gene expression array analysis of phase I and II enzymes on C57BL/6 mice fed cholesterol supplemented formononetin or 2-heptyl formononetin (CF7) for three weeks (experiment 1).

Drug Transporters:

Metallothioneins: Mt2, Mt3.

P-Glycoprotein Family: Abcb1a, Abcb1b, Abcb4, Abcc1, Gpi1.

Phase I Metabolising Enzymes:

P450 Family: Cyp11b2, Cyp17a1, Cyp19a1, Cyp1a1, Cyp1a2, Cyp27b1, Cyp2b10, Cyp2C29, Cyp2e1, Cyp4b1.

Phase II Metabolising Enzymes:

Carboxylesterases: Ces1, Ces2.

Decarboxylases: Gad1, Gad2.

Dehydrogenases: Adh1, Adh4, Adh5, Alad, Aldh1a1, Hsd17b1, Hsd17b2, Hsd17b3.

Glutathione Peroxidases: Gpx1, Gpx2, Gpx3, Gpx5, Gsta1, Gsta3, Gsta4, Gstm1, Gstm2, Gstm3, Gstm4, Gstm5, Gstp1, Gstt1, Gstz1, Lpo, Mpo.

Hydrolases: Ephx1, Faah, Fbp1.

Kinases: Hk2, Pklr, Pkm2.

Lipoxygenases: Alox12, Alox15, Alox5, Apoe.

Oxidoreductases: Blvra, Blvrb, Cyb5r3 (Dia1), Gpx1, Gpx2, Gsr, Mthfr, Nos3, Nqo1, Srd5a1, Srd5a2.

Paraoxonases: Pon1, Pon2, Pon3.

Sulfotransferases: Chst1, Gsta3, Gstm2, Gstm3, Gstm5, Gstp1, Gstt1, Mgst1, Mgst2, Mgst3.

Transferases: Nat1, Nat2, Comt, Ggt1.

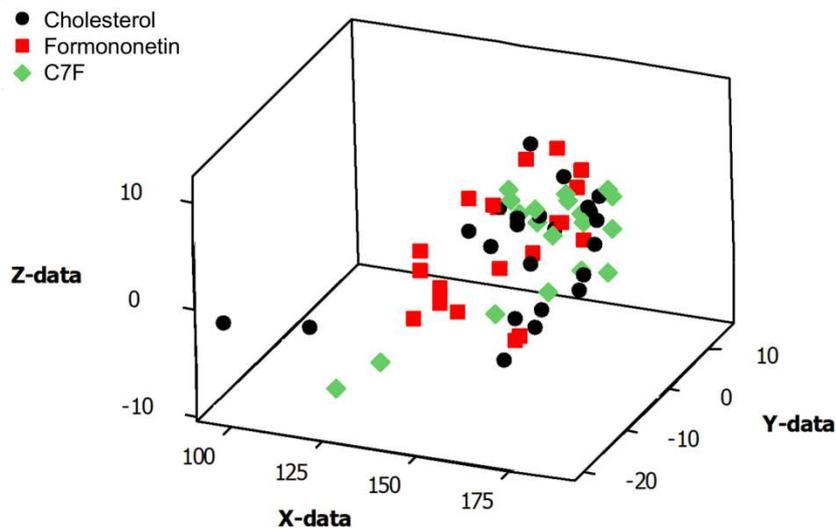
Other Genes Related to Drug Metabolism: Abp1, Ahr, Arnt, Asna1, Gckr, Marcks, Smarcal1, Snn.

Supplemental Table 3: Primer sequences.

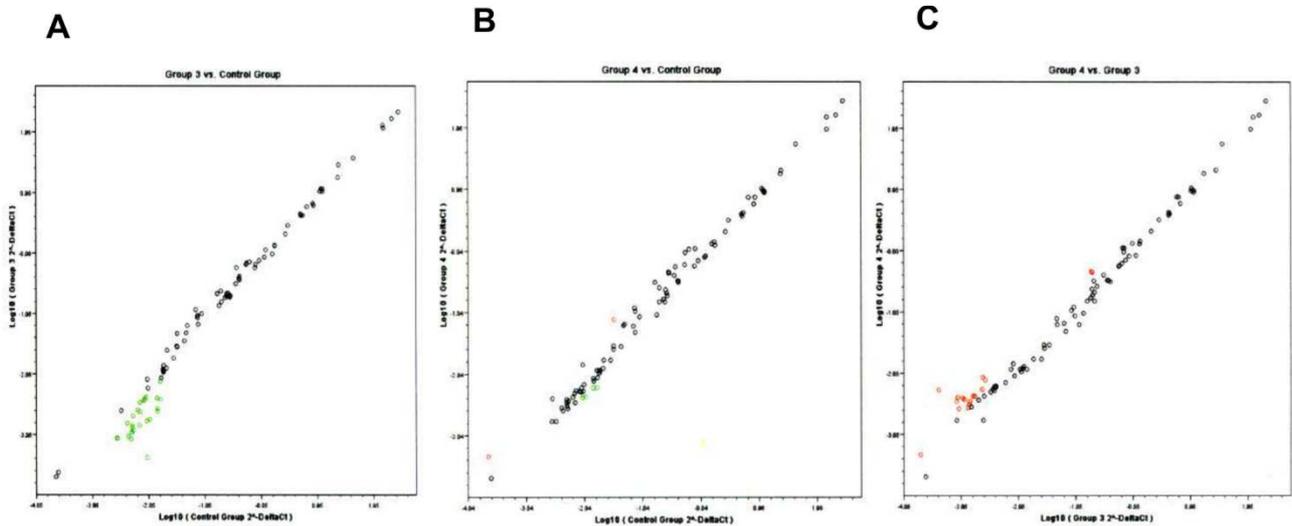
Gene name	Trivial name	5' prime	3' prime
<i>Rn18S</i>	Ribosomal 18S	GTA ACC CGT TGA ACC CCA TT	CCA TCC AAT CGG TAG TAG CG
<i>Acat2</i>	Acetyl-coA acetyltransferase 2	GGT GGA ACT ATG TGG CCA AGA	CCA GGA TGA AGC AGG CAT AGA
<i>Acaca</i>	Acetyl-coA carboxylase alpha	TGC TGC CCC ATC CCC GGG	TCG AAC TCT CAC TGA CAG G
<i>Acox1</i>	Acyl-Coenzyme A oxidase 1	GAT CAC GGG CAC TTA TGC G	CGA AGA TGA GTT CCA TGA CCC
<i>Atgl</i>	Adipose triglyceride lipase	CAT CTC CCT GAC TCG TGT TTC	CAA GTT GTC TGA AAT GCC GC
<i>Cebpa</i>	CCAAT/enhancer binding protein alpha	CAA GAA CAG CAA CGA GTA CCG	GTC ACT GGT CAA CTC CAG CAC
<i>Cpt1a</i>	Carnitine palmitoyltransferase 1a	TAC TGL TGT ATC GTC GCA CG	GAC GAA TAG GTT TGA GTT CCT CAC
<i>Cpt1b</i>	Carnitine palmitoyltransferase 1b	GAT CTG GGC TAT CTG TGT CCG T	ACG TTT GGA AGC TGT AGA GCA TG
<i>Cyp7a1</i>	Cholesterol 7 alpha-hydroxylase	AGC AAC TAA ACA ACC TGC CAG TAC TA	GTC CGG ATA TTC AAG GAT GCA
<i>Dgat2</i>	Diglyceride acyltransferase 2	GGG TCC AGA AGA AGT TCC AGA AG	CCC AGG TGT CAG AGG AGA AGA G
<i>Emr1</i>	EGF-like module containing, mucin-like, hormone receptor-like sequence 1	CTT TGG CTA TGG GCT TCC AGT	GCA AGG AGG ACA GAG TTT ATC TGT
<i>Fasn</i>	Fatty acids synthase	ATT GGT GGT GTG GAC ATG	CCCAGC CTT CCA TCT CCT G
<i>Gpat1</i>	Glycerol phosphate acyltransferase 1	GCT ATC ATG TCC ACC CAC ATT G	ACT TCC TCC TTC ATC ACA AAG AAG TC
<i>Hmgcr</i>	3-hydroxy-3-methylglutaryl-Coenzyme A Reductase	CCC TGA GTT TAG CCT TCC TTT TG	GCT TTC TTT GAG GTC ACG ACG G
<i>Ldlr</i>	Low density lipoprotein receptor	GCA TCA GCT TGG ACA AGG TGT	GGG AAC AGC CAC CAT TGT TG
<i>Mlxipl</i>	MLX interacting protein-like	CTT CTT CCG TTG CAC ATA CTG	GTT GCT ATG CCG GGA CAA GA
<i>Mttp</i>	Microsomal triglyceride transfer protein	CAA GCT CAC GTA CTC CAC TGA AG	TCA TCA TCA CCA TCA GGA TTC CT
<i>Nr1h3</i>	Nuclear receptor subfamily 1, group H, member 3	CCG ACA GAG CTT CGT CC	CCC ACA GAC ACT GCA CAG
<i>Nr1h4</i>	Nuclear receptor subfamily 1, group H, member 4	CCA ACC TGG GTT TCT ACC C	CAC ACA GCT CAT CCC CTT T
<i>Ppara</i>	Peroxisome proliferator-activated receptor alpha	AGA GAG GAC AGA TGG GGC TC	CGT TTG TGG CTG GTC AAG TT
<i>Pparg2</i>	Peroxisome proliferator-activated receptor gamma 2	ACAGCAAATCTCTGT TTTATGC	TGCTGGAGAAATCAAC TGTGG
<i>Scd1</i>	Stearoyl-CoA desaturase-1	ACA CCT GCC TCT TCG GGA TT	TGA TGG CCA GAG CGC TG
<i>Srebf1</i>	Sterol regulatory element-binding protein-1c	GGA GCC ATG GAT TGC ACA TT	GCT TCC AGA GAG GAG GCC AG
<i>Tnf</i>	Tumour necrosis factor	CCC TCA CAC TCA GAT CAT CTT CT	GCT ACG ACG TGG GCT ACA G
<i>Ucp1</i>	Uncoupling protein 1	AGCCGGCTTAATGACT GGAG	TCTGTAGGCTGCCCAA TGAAC

Supplementary figure legends and figures

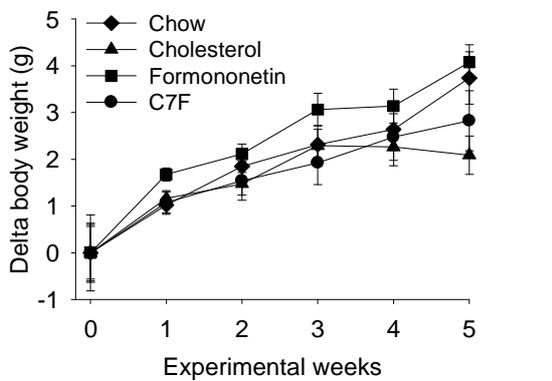
Supplementary Figure 1: Clustering of gut microbiota from C57BL/6 mice fed formononetin or 2-heptyl formononetin (C7F) (experiment 1) (n=23-25). Three-dimensional principal component analysis (3D-PCA) based on denaturation gradient gel electrophoresis data. Black circle: cholesterol, red square: formononetin, green square: C7F. The clustering was significant on the Y-axis and the Z-axis. On the Y-axis C7F fed mice clustered significantly differently from mice fed cholesterol and the formononetin. Also the formononetin fed mice clustered significantly differently from mice fed cholesterol. On the Z-axis the C7F fed mice clustered borderline differently from mice fed cholesterol.



Supplementary Figure 2: Gene expression of phase I and II metabolites (experiment 1). Scatter plot of liver gene expression (n=23-25). Gene expression in **A** mice fed chow compared to formononetin, **B** chow compared to C7F and **C** formononetin compared to C7F.



Supplementary Figure 3: Body weight of C57BL/6 mice fed chow, cholesterol or cholesterol supplemented formononetin or 2-heptyl-formononetin (C7F) for five weeks (experiment 2) (n=8). Graphs shows mean \pm SEM.



Supplementary Figure 4: Total feed intake in C57BL/6 mice fed chow, cholesterol or cholesterol supplemented formononetin or 2-heptyl-formononetin (C7F) for five weeks (experiment 2) (n=8).

Graphs shows mean \pm SEM.

