

Figure S1: The placental weight and litter characteristics among the four groups (A) litter size (B) fetal weight (C) placenta weight.

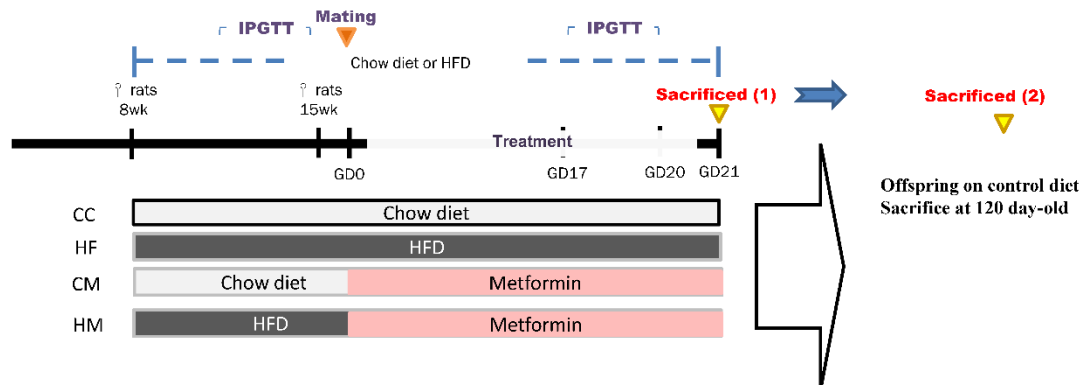


Figure S2: Schematic diagram of the experimental design.

Supplementary Table S1. Primer sequences for RT-PCR in this study.

Rat Gene	primer	sequence (5' to 3')
<i>GLUT1</i>	Forward	TGGCCAAGGACACACGAATACTGA
	Reverse	TGGAAGAGACAGGAATGGGCGAAT
<i>GLUT2</i>	Forward	TAGTCAGATTGCTGGCCTCAGCTT
	Reverse	TTGCCCTGACTTCCTCTTCCAAC
<i>GLUT3</i>	Forward	TGGCTACAACACCGGAGTCATCAA
	Reverse	CTGCCAAAGCGGTTGACAAAGAGT
<i>GLUT4</i>	Forward	TTTCCAGTATGTTGCGGATG
	Reverse	TCAGTCATTCTCATCTGGCC
<i>Leptin</i>	Forward	CGGTTCTGTGGCTTTGGT
	Reverse	CCGACTGCGTGTGTGAAATG
<i>LepR</i>	Forward	ACTGGGACATAGAGTGCTGGAT
	Reverse	GTTGCACTGGACAGTCTGAAAG
<i>ACC1</i>	Forward	TGAGGAGGACCGCATTATC
	Reverse	GCATGGAATGGCAGTAAGGT
<i>ACC2</i>	Forward	CGCTGCGGTCAAGTGT
	Reverse	CGTTGGCGTAGTTGTTATT
<i>ACL</i>	Forward	ACCCAGAGGAAGCCTACATTGC
	Reverse	TTCGCCAGTTCGTTGACACC
<i>FAS</i>	Forward	AGATCCTGGAACGTGAACATGA
	Reverse	GCCGTACTTCACGAATGGGT
<i>LPL</i>	Forward	GTACAGTCTTGGAGCCCATGC
	Reverse	GCCAGTAATTCTATTGACCTTCTTGTT
<i>ATGL</i>	Forward	TGTGGCCTCATTCTCCTAC
	Reverse	TCGTGGATGTTGGTGGAGCT
<i>HSL</i>	Forward	GCTGGGCTGTCAAGCACTGT
	Reverse	GTAAGTGGGTAGGCTGCCAT
<i>MGL</i>	Forward	TGACCAACTCTGTCCTCCAT
	Reverse	GCACTGCCCTTCCTCTT
<i>18S</i>	Forward	GCGATGCGGCGGCGTTAT
	Reverse	AGACTTTGGTTTCCCGGAAGC
<i>GAPDH</i>	Forward	TCTTGTGCAGTGCCAGCCTC
	Reverse	GTCACAAGAGAAGGCAGCCCTGG

Abbreviation: GLUT1, glucose transporter 1 ; GLUT2, glucose transporter 2 ; GLUT3, glucose transporter 3 ; GLUT4, glucose transporter 4 ; LepR, leptin receptor ; ACC1, acetyl-CoA carboxylase 1 ; ACC2, acetyl-CoA carboxylase 2 ; ACL, acylcholesterol

lipase; FAS, fatty acid synthetase ; LPL, lipoprotein lipase; ATGL, adipose triglyceride lipase ; HSL, hormone-sensitive lipase ; MGL, monoacylglycerol lipase ; GAPDH, glyceraldehyde-3-phosphate dehydrogenase

Supplementary Table S2. The weekly body weight change of dams

	8w	9w	10w	11w	12w	13w
<b>CC</b>	210.22±3.33	208.80±2.68	214.90±2.61	227.55±2.95	231.84±2.77	240.28±2.93
<b>HC</b>	217.21±4.17	234.51±5.59 <sup>*</sup>	255.14±8.26 <sup>*</sup>	272.96±8.95 <sup>*</sup>	287.96±9.84 <sup>*</sup>	298.01±9.88 <sup>*</sup>
<b>CM</b>	205.38±2.68	199.85±1.94 <sup>#</sup>	205.90±1.73 <sup>#</sup>	217.40±2.42 <sup>#</sup>	225.60±1.87 <sup>#</sup>	232.00±2.19 <sup>#</sup>
<b>HM</b>	215.01±4.30	232.80±4.45 <sup>*†</sup>	253.25±4.64 <sup>*†</sup>	269.21±5.36 <sup>*†</sup>	279.28±5.62 <sup>*†</sup>	289.60±7.40 <sup>*†</sup>
	14w	15w	16w	17w	18w	19w
<b>CC</b>	252.42±3.03	247.00±4.24	255.40±4.75	276.89±4.81	292.05±5.39	312.47±2.52
<b>HC</b>	307.70±10.34 <sup>*</sup>	305.11±11.70 <sup>*</sup>	309.78±13.08 <sup>*</sup>	318.33±18.20 <sup>*</sup>	355.29±11.52 <sup>*</sup>	395.29±10.92 <sup>*</sup>
<b>CM</b>	241.03±2.48 <sup>#</sup>	253.07±2.36 <sup>#</sup>	261.47±2.37 <sup>#</sup>	265.25±2.29 <sup>#</sup>	278.15±4.83 <sup>#</sup>	302.00±9.24 <sup>#</sup>
<b>HM</b>	298.10±8.39 <sup>*†</sup>	300.75±8.40 <sup>*†</sup>	303.00±8.91 <sup>*†</sup>	291.44±7.99	292.95±6.91 <sup>#</sup>	300.19±6.25 <sup>#</sup>

The results are presented as mean ± standard error. ;\*compared with CC,  $P < 0.05$ ; # compared with HC,  $P < 0.05$ ; † compared to the CM group,  $P < 0.05$ .

Abbreviation: CC, maternal control diet ; HC, maternal high-fat diet ; CM, maternal control diet plus metformin treatment ; HM, maternal high-fat diet plus metformin treatment

Supplementary Table S3. Intraperitoneal glucose tolerance test (IPGTT) of dams

Min.	CC	HC	CM	HM
0	71.00 ± 4.70	64.40 ± 4.23	61.60 ± 4.70	57.20 ± 2.27
15	183.80 ± 39.97	386.20 ± 57.93 <sup>*</sup>	290.40 ± 24.06	254.80 ± 49.53 <sup>*#</sup>
30	166.00 ± 30.68	319.20 ± 48.24 <sup>*</sup>	206.60 ± 6.80 <sup>#</sup>	220.80 ± 48.94
60	114.00 ± 20.71	142.00 ± 14.20	90.00 ± 3.45	127.80 ± 31.18
120	82.80 ± 9.96	76.20 ± 6.87	69.00 ± 6.28	58.40 ± 6.26
AUC (x1000)	14.64 ± 1.76	22.13 ± 2.10	15.59 ± 0.54	16.72 ± 2.70

The results are presented as mean ± standard error. ;\*compared with CC,  $P < 0.05$ ; # compared with HC,  $P < 0.05$ .

The integrated AUC values of the IPGTT were calculated using the trapezoidal method and a repeated-measure analysis of variance (ANOVA) model.

Abbreviation: CC, maternal control diet ; HC, maternal high-fat diet ; CM, maternal control diet plus metformin treatment ; HM, maternal high-fat diet plus metformin treatment ; AUC, area under curve

Supplementary Table S4. The monthly body weight change of offspring

	CC	HC	CM	HM
1M	54.36±2.38	86.92±3.67 <sup>*</sup>	53.53±1.19 <sup>#</sup>	64.37±6.30 <sup>*#†</sup>
2M	202.40±5.09	268.04±8.48 <sup>*</sup>	213.09±4.01 <sup>#</sup>	226.20±11.92 <sup>*#</sup>
3M	322.32±4.92	397.19±12.38 <sup>*</sup>	351.55±6.11 <sup>*#</sup>	351.31±10.65 <sup>*#</sup>
4M	384.15±7.22	459.63±10.85 <sup>*</sup>	416.15±9.61 <sup>*#</sup>	427.92±12.89 <sup>*</sup>

\* compared to the CC group,  $P < 0.05$ ; # compared to the HC group,  $P < 0.05$ ; † compared to the CM group,  $P < 0.05$ .

Abbreviation: CC, maternal control diet ; HC, maternal high-fat diet ; CM, maternal control diet plus metformin treatment ; HM, maternal high-fat diet plus metformin treatment ; AUC, area under curve

Supplementary Table S5. Intraperitoneal glucose tolerance test (IPGTT) of offspring

Min.	CC	HC	CM	HM
0	80.92 ± 2.33	80.75 ± 2.97	78.00 ± 2.24	85.75 ± 1.87
15	240.31 ± 18.47	308.58 ± 20.63 *	313.85 ± 16.99 *	238.75 ± 24.33 #†
30	209.46 ± 14.75	235.33 ± 19.30	278.77 ± 12.07 *	213.00 ± 25.33 †
60	152.15 ± 8.62	171.83 ± 9.00	188.23 ± 8.08 *	138.50 ± 20.03 #†
120	90.92 ± 4.89	99.33 ± 6.06	99.69 ± 4.59	98.13 ± 4.08
AUC (x1000)	18.50 ± 0.92	21.24 ± 1.07	23.03 ± 0.84 *	18.19 ± 1.70 †

The results are presented as mean ± standard error. ;\*compared with CC,  $P < 0.05$ ; # compared with HF,  $P < 0.05$ ; † compared to the CM group,  $P < 0.05$ .

Abbreviation: CC, maternal control diet ; HC. maternal high-fat diet ; CM, maternal control diet plus metformin treatment ; HM, maternal high-fat diet plus metformin treatment ; AUC, area under curve