

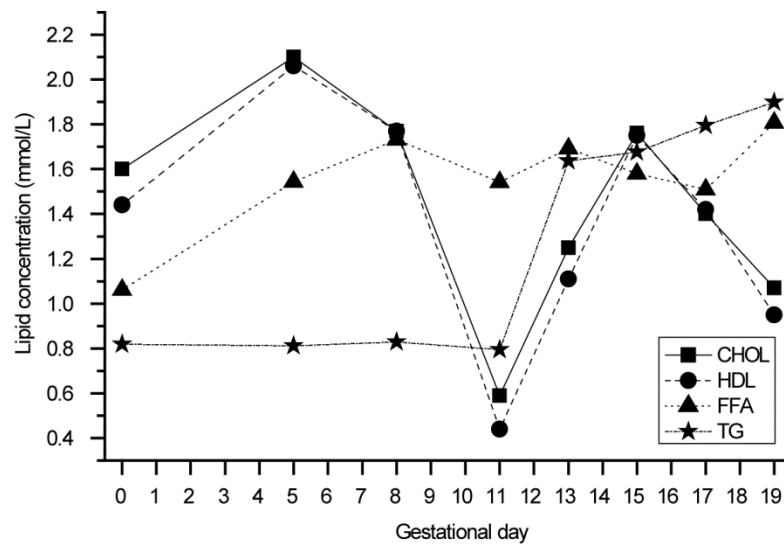
A progesterone-brown fat axis is involved in regulating fetal growth

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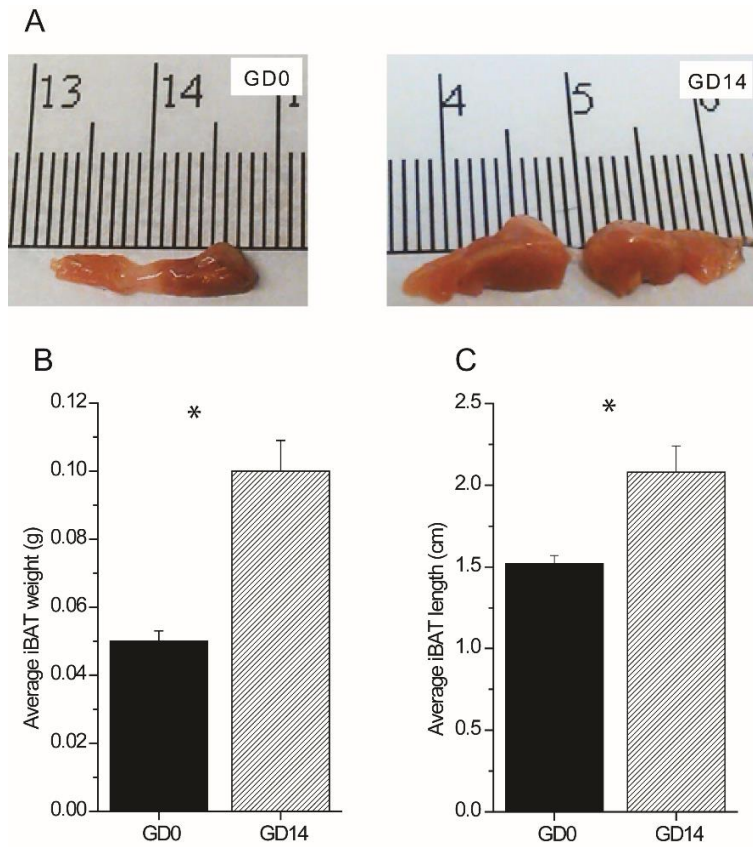
Jansen, Shadi Abu-Hayyeh and Catherine Williamson

SUPPLEMENTARY INFORMATION

Supplementary Data

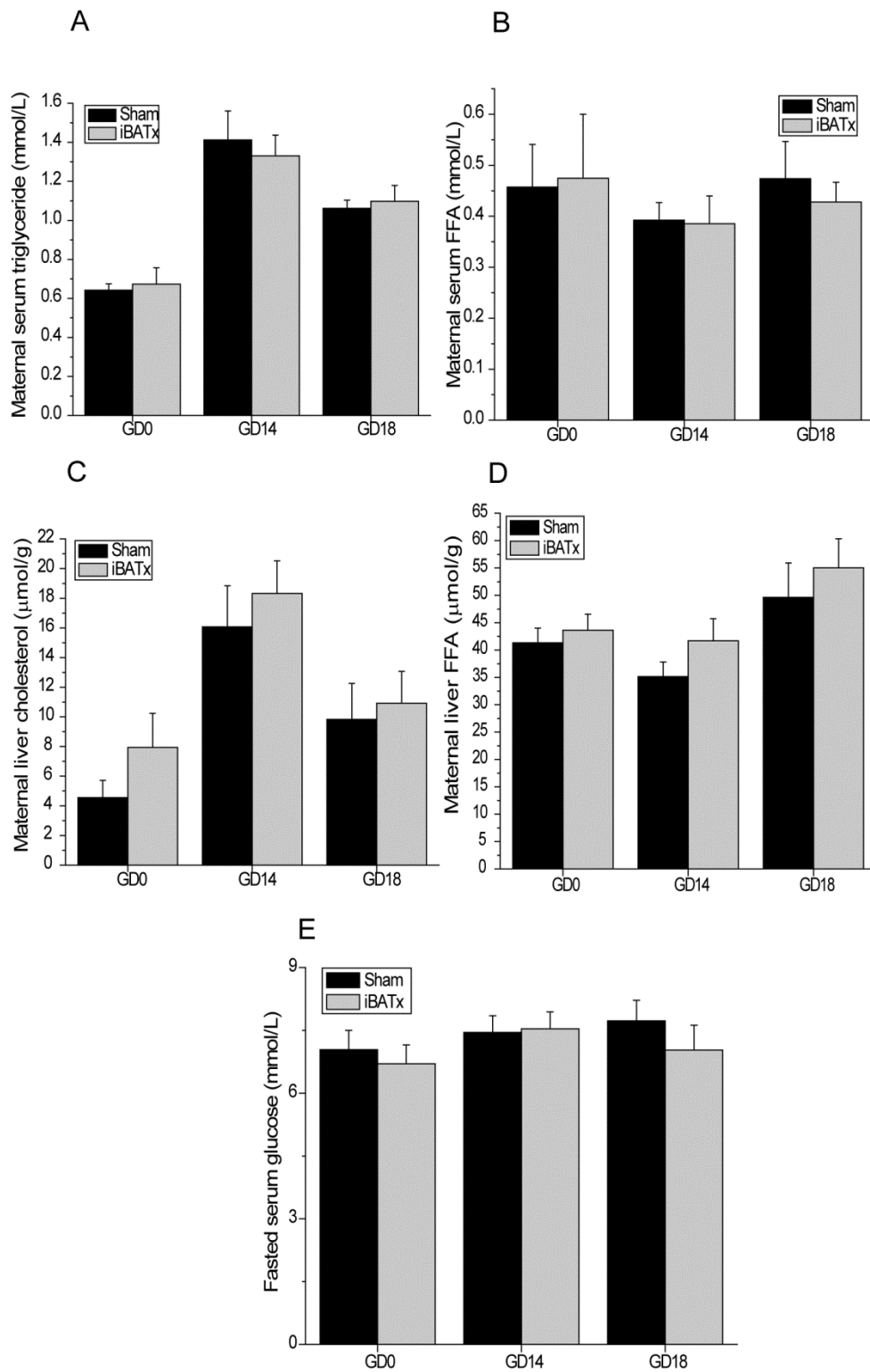


Supplementary Figure S1. Switching of serum lipid biochemistry takes place at GD11 onwards
Serum concentrations of cholesterol (CHOL), HDL-cholesterol (HDL), free fatty acids (FFA) and triglycerides (TG). Pooled sampled serum from n=11 pregnant mice.



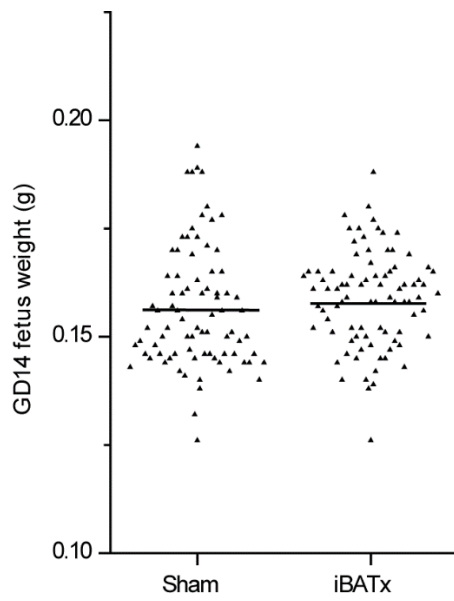
Supplementary Figure S2. Pregnancy alters the morphometry of BAT

Images of iBAT taken from non-pregnant (GD0) or day 14 pregnant (GD14) mice (A). Average iBAT weight (B) and length (C). Data represented as mean \pm SEM, n=5 per group. * $p < 0.05$ as determined by Student's t-test.



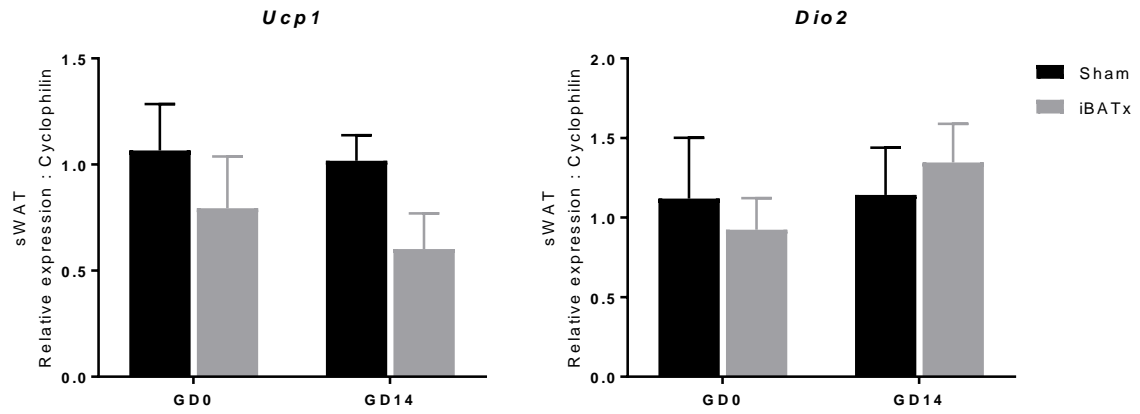
Supplementary Figure S3. The effect of surgically ablating interscapular BAT on maternal biochemistry

Lipid profiles of maternal (A) serum triglycerides, (B) serum free fatty acids (FFA), (C) liver cholesterol and (D) liver FFA. (E) Maternal fasted serum glucose. Data represented as mean \pm SEM, n=6-11 per group.

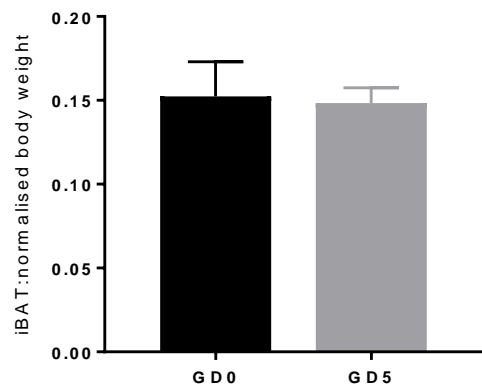


Supplementary Figure S4. Surgical ablation of maternal interscapular BAT does not alter fetal weight at gestational day 14

Sham / iBATx fetal weight: n=82 / 86 fetuses from 11 / 11 mothers.

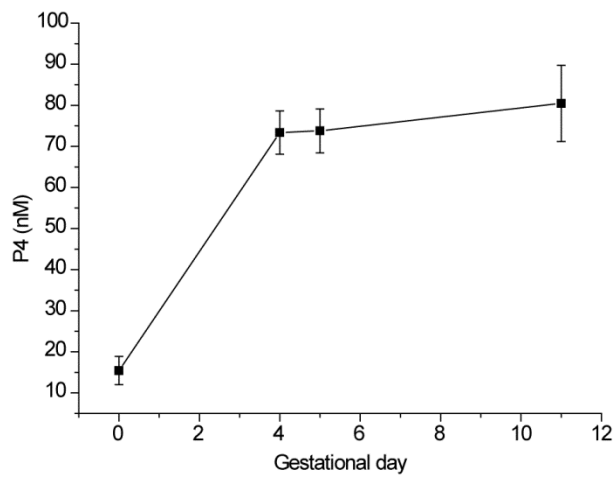


Supplementary Figure S5. Surgical ablation of maternal interscapular BAT does not alter maternal subcutaneous WAT expression of BAT marker genes *Ucp1* and *Dio2*
Data represented as mean \pm SEM, n=3-6 per group.



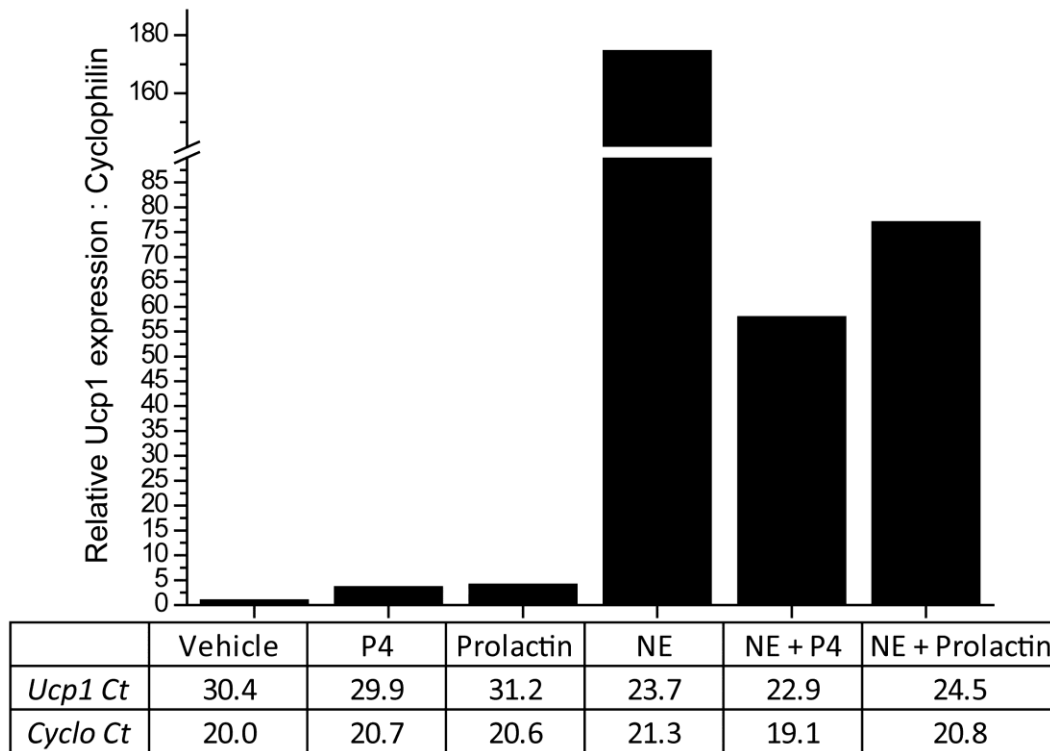
Supplementary Figure S6. No changes in normalised iBAT weight at GD5

Data represent mean \pm SEM, GD0 n=4, GD5, n=7.



Supplementary Figure S7. Serum progesterone concentrations increase rapidly following coitus
Data represented as mean \pm SEM, n=9-11.

Supplementary Experimental Procedures



Supplementary Figure S8. Representative example of *Ucp1* mRNA expression in cultured primary brown adipocytes +/- treatment with progesterone (P4), prolactin and norepinephrine (NE) (n=1).

Serum Progesterone Measurements. The serum progesterone concentration in mice at GD0 to GD11 was measured using the HTRF progesterone assay kit (CisBio) following a 1:15 dilution in assay buffer, according to manufacturer's instructions.

Table of Primer Sequences

Gene	Forward Primer	Reverse Primer
<i>β-Actin</i>	GATGACGATATCGCTGCGCTG	CTAGGGCGGCCACGAT
<i>Abca1</i>	TCCTCATCCTCGTCATTCAA	GGACTTGGTAGGACGGAACCT
<i>Abcg1</i>	GCTGTGCGTTTTGTGCTGTT	TGCAGCTCCAATCAGTAGTCC
<i>Abcg5</i>	TCAATGAGTTTTACGGCCTGAA	GCACATCGGGTGATTTAGCA
<i>Abcg8</i>	TGCCCACCTCCACATGTC	ATGAAGCCGGCAGTAAGGTAG
<i>Acc1</i>	GGCCAGTGCTATGCTGAGAT	AGGGTCAAGTGCTGCTCCA
<i>Acc2</i>	ACTTTGACCTGACCGCTGTG	CTGAGTGCCGGATAATGGC
<i>Adipoq</i>	AGATGCAGGTCTTCTTGCTC	CTGAGCGATACACATAAGCGG
<i>Adrb3</i>	GGTAATCATAGCCATCGCCC	CAGTTACACAGAGCACGTCC
<i>Adrp</i>	TGGCAGCAGCAGTAGTGGAT	AGCTCACCAAGGGCAGGTT
<i>Cd36</i>	GATGTGGAACCCATAACTGGA	GGCTTGACCAATATGTTGACC
<i>Cidea</i>	AGAGTCACCTTCGACCTATACAG	AACCTCAGCAGATTCTTAACAC
<i>Cyp7a1</i>	AGCAACTAAACAACCTGCCAGTACTA	GTCCGGATATTCAAGGATGCA
<i>Cyclophilin B</i>	TGGAGAGCACCAAGACAGACA	TGCCGGAGTCGACAATGAT
<i>Dgat2</i>	AGTGGCAATGCTATCATCATCGT	AAGGAATAAGTGGGAACCCAGATCA
<i>Dio2</i>	CCTACAAACAGGTTAAACTGGG	CTCTGCACTGGCAAAGTC
<i>El</i>	AGCGTCTATTGTTACTTCCC	TTTATGATGCTCATCTCGCAG
<i>Fabp-pm</i>	CAAAGATGCAGAAGAAGCC	CCTCTTGCAACCATTGCT
<i>Fas</i>	CCCAGAGGCTTGTGCTGACT	CGAATGTGCTTGGCTTGGT
<i>Fatp1</i>	CGTTTCGATGGTTATGTTAGTG	CTAGCACGTCACCTGAGAG
<i>Fatp4</i>	TTCCCTCATCCTCCTGCT	CGATGTTTCTGCTGAGTG
<i>Hmgcr</i>	TTGGCACCATGTGAGGCGTCC	AGCGACACACAGGCCGGGAA
<i>Hsl</i>	CTATTCAGGGACAGAGGCAG	TAGTTCCAGGAAGGAGTTGAG
<i>Lal</i>	TCTTCTCAAGGACATGTTTG	CAAAGCTCCTTCATGATGAC
<i>Lep</i>	GTTCAAGCAGTGCCTATCC	AAGTCCAAGCCAGTGACC
<i>Lpl</i>	CCAGGATGCAACATTGGAGA	CAACTCAGGCAGAGCCCTTT
<i>Lxra</i>	AGGAGTGTGACTTCGCAAA	CTCTTCTTGCCGCTTCAGTT
<i>Lxrβ</i>	AAGCAGGTGCCAGGGTTCT	TGCATTCTGTCTCGTGGTTGT
<i>mtCox2</i>	TGCTCTCCCCTCTCTAC	GGTGCCCTATGGTTTTAACG
<i>Nd5</i>	AGCATTGCGGAAGCATCTTTG	TTGTGAGGACTGGAATGCTG
<i>Pgc1a</i>	TGAAAGGGCCAAACAGAGAGA	TAAATCACACGGCGCTCTT
<i>Scd1</i>	CCCCTGCGGATCTTCCTTAT	AGGGTTCGGGTGTGTTTCT
<i>Scd2</i>	AGCGGGCTGCAGAACTTAG	GGCTGAGTAAGCGCCAGAGAT
<i>Srebp1c</i>	GCAGCCATGGATTGCACATT	GGCCCGGAAGTCACTGT
<i>Ucp1</i>	AATACTGGCAGATGACGTCC	TTACCACATCCACTGGAGAG
<i>Zic1</i>	CACATGAAGGTCCATGAGTCC	GGGTTGTCTGTTGTGGGAG