

**The postnatal leptin surge in mice is variable in both time and intensity and reflects nutritional status.**

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### **Supplementary Material**

*Supplementary figure and table legends*

**Fig. S1.** Over- and undernutrition effects on mean plasma leptin concentrations during the first 4 weeks of life in females (A) compared to males (B). (C) AUC for plasma leptin for each sex. Differences in leptin concentration AUC among the groups were tested using two-way ANOVA with Sidak post hoc testing at the 5% level of significance; ns denotes non-significant difference. Means shown with  $\pm$  SEM.

**Fig. S2.** Body weight (A), body fat (B) and lean mass (C) in the replication cohort of mixed sexes pups raised in small (SL; 3 pups per dam; orange), normal (NL; 7-8 pups per dam; gray), and large (LL; 11-12 pups per dam; blue) litters that were monitored over the first 4 weeks of life and bled once per week. GEE analysis performed to compare changes in body weight and

composition during the pre-weaning period while pups were in their respective litter size groups; statistics in supplementary Table 3. Different lowercase letters indicate significant difference by the Wald Chi-Square test at the 5% level of significance. Data shown as mean  $\pm$  SEM.

**Fig. S3.** Maternal HFD-feeding effects on mean plasma leptin concentrations during the first 4 weeks of life in females (A) compared to males (B). (C) AUC for plasma leptin for each sex. Differences in leptin concentration AUCs among the groups were tested using two-way ANOVA with Sidak post hoc testing at the 5% level of significance; ns denotes non-significant difference. Means shown with  $\pm$  SEM.

**Fig. S4.** Over- and undernutrition effects on leptin surge in plasma in the replication cohort bled once per week. (A) Mean plasma leptin concentration, (B) mean leptin concentration AUC over P6-P17, (C) mean leptin concentration AUC over P6-P17 normalized to mean fat mass over that time period, (D) mean peak leptin concentration (peak defined as the maximum leptin concentration measured during the first 4 weeks of postnatal life), (E) mean peak leptin concentration adjusted for body weight and age at which the peak occurred in mixed sexes pups raised in small (SL; 3 pups per dam; orange), normal (NL; 7-8 pups per dam; gray), and large (LL; 11-12 pups per dam; blue) litters that were monitored over the first 4 weeks of life. Differences in leptin AUC and peak leptin concentrations among the groups were tested using one-way ANOVA with Tukey post hoc (or Sidak for ANOVA with covariates) at the 5% level of significance; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . Means shown with  $\pm$  SEM.

**Fig. S5.** Plasma leptin (A) and relative expression of *Lepr* (B), *Adrb3* (C), *Ucp1* (D), *Hsl* (E), and *Adipoq* (F) in SCAT, PGAT, PR, Mes, iBAT, and aBAT in P10 pups reared in small, normal or large litters. Relative expression was normalized to a housekeeping gene, *Actb*. Differences among the groups were tested with one- or two-way ANOVA with Tukey post hoc testing at the 5% level of significance; \* $p < 0.05$ ; \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . Means shown with  $\pm$  SEM.

**Table S1.** Sequences for primer-probe mixes used for gene expression assays.

**Table S2.** Statistical results of the Generalized Estimating Equation (GEE) used to compare body weight, fat mass and lean mass in the small (SL), normal (NL) and large (LL) litters pups of the primary cohort.

**Table S3.** Statistical results of the Generalized Estimating Equation (GEE) used to compare body weight, fat mass and lean mass in the small (SL), normal (NL) and large (LL) litters pups of the replication cohort.

**Table S4.** Statistical results of the Generalized Estimating Equation (GEE) used to compare body weight, fat mass and lean mass in the pups raised by dams fed HFD at parturition, or for 20 weeks pre-conception, or control dams fed breeder chow during the same periods.

Fig. S1

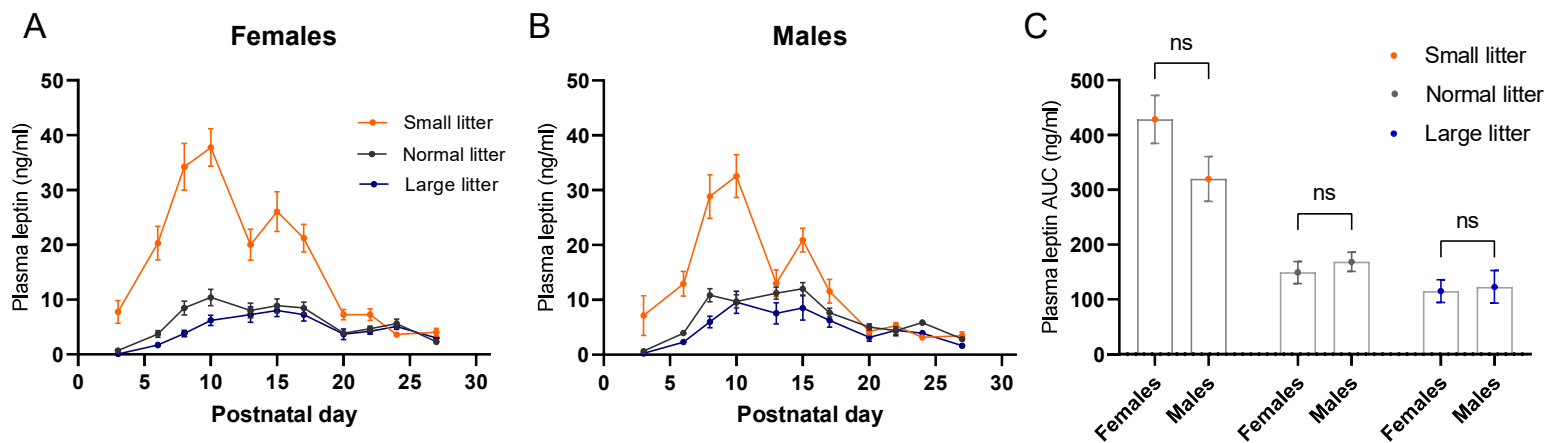


Fig. S2

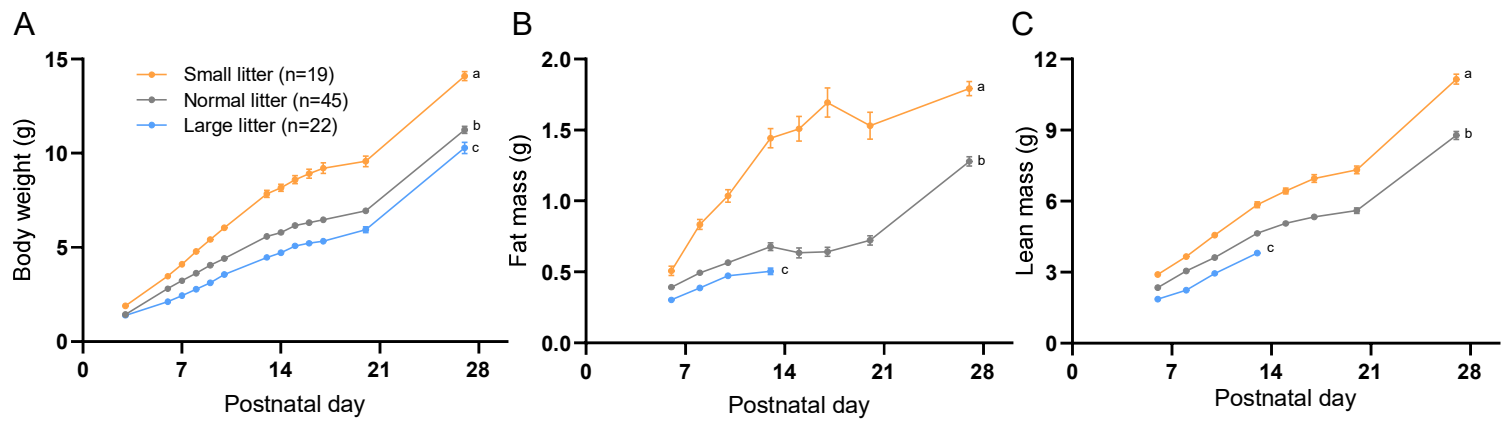


Fig. S3

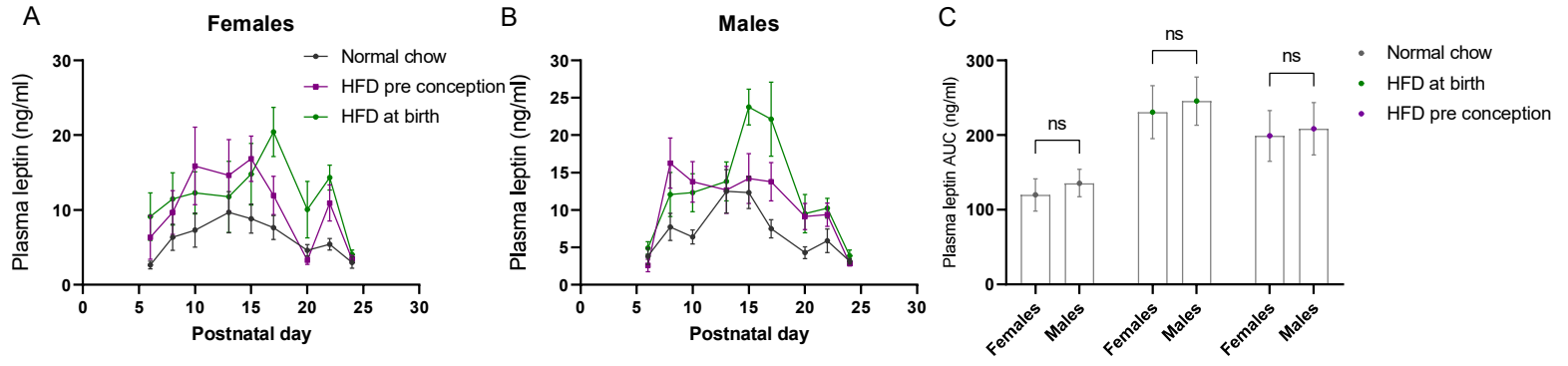


Fig. S4

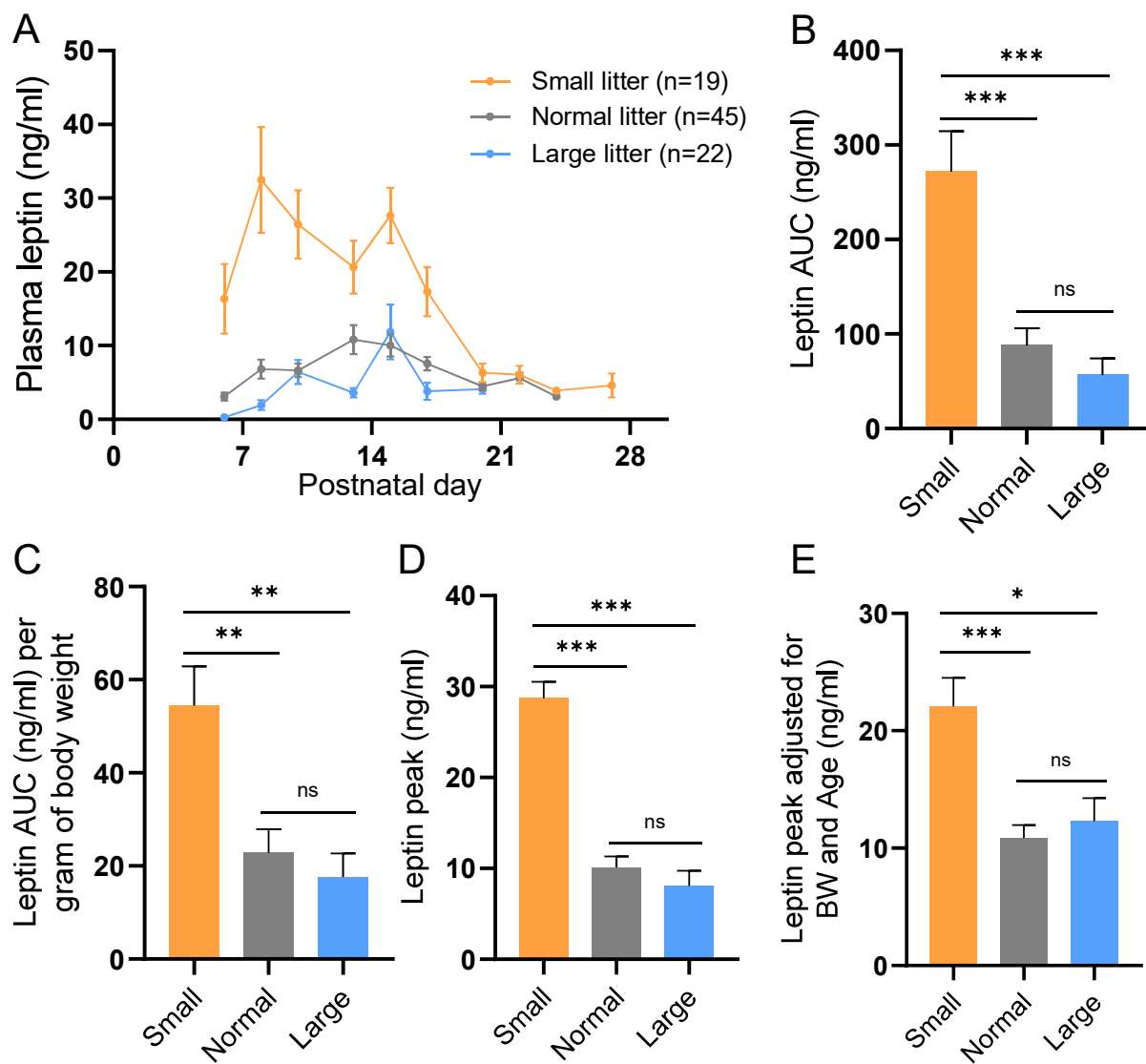
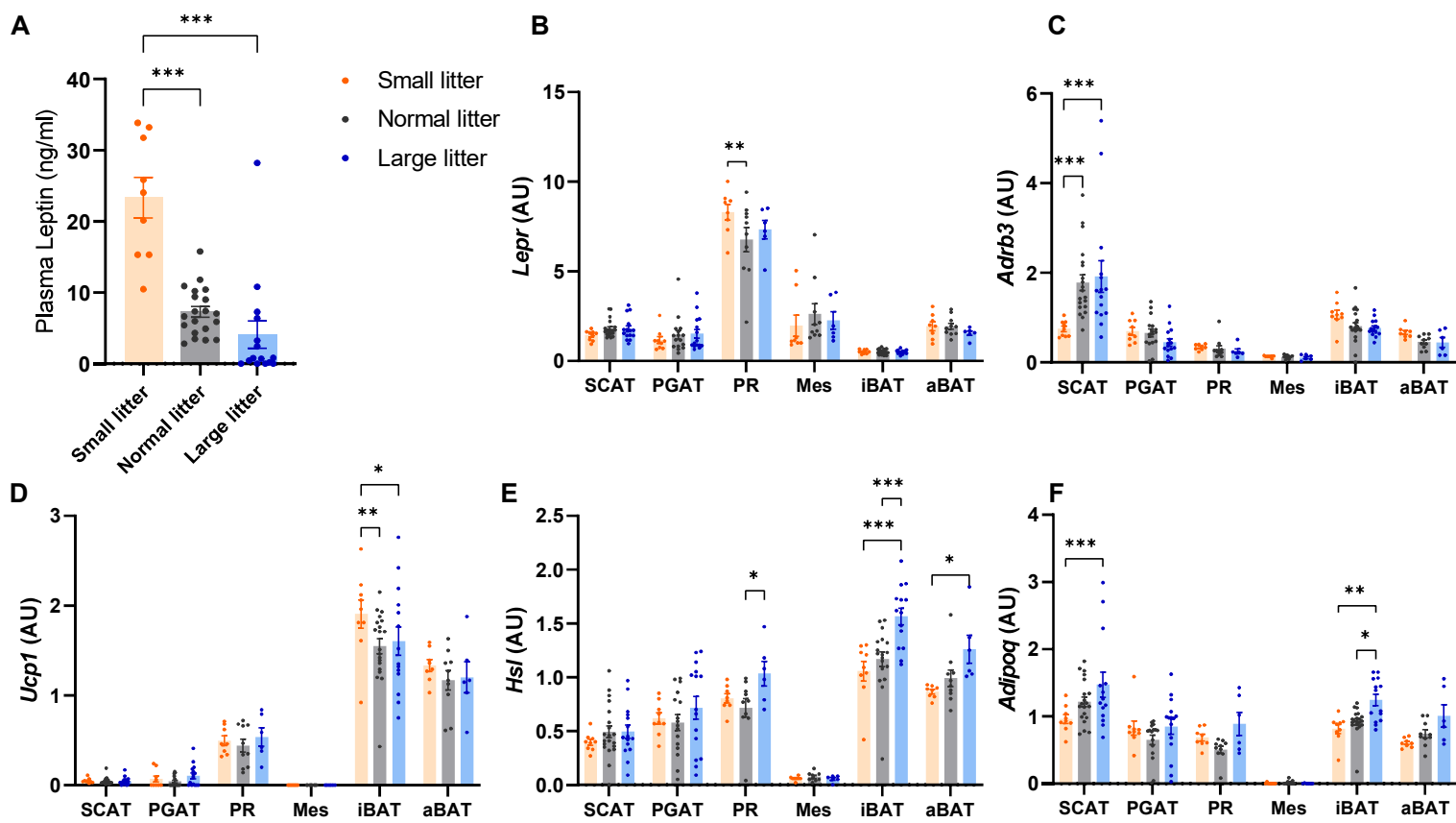


Fig. S5





**Supplemental Table 1**

Sequences for primer-probe mixes used for gene expression assays

Gene	Sequence	
<i>Actb</i>	Forward	GACTCATCGTACTCCTGCTTG
	Reverse	GATTACTGCTCTGGCTCCTAG
	Probe:	/5HEX/CTGGCCTCA/ZEN/CTGTCCACCTTCC/3IABkFQ/
<i>Lep</i>	Forward	TTCTCCAAGAGCTGCTCC C
	Reverse	CCTGGTGGCCTTTGAAACTT
	Probe:	/5Cy5/AAC AGT TGG /TAO/ATG TTA GCC CTG AAT GCT /3IAbRQSp/
<i>Ptgfr</i>	Forward	GATCTGATTCCACGTTGCCA
	Reverse	GCCATAATGTGCGTCTCCT
	Probe:	/56-FAM/TGGAGTCCC/ZEN/TTTCTGGTAACAATGGC/3IABkFQ/
<i>Ucp1</i>	Forward	CACACCTCCAGTCATTAAGCC
	Reverse	CAAATCAGCTTTGCCTCACTC
	Probe:	/56-FAM/AAACACCTG/ZEN/CCTCTCTCGGAAACAA/3IABkFQ/
<i>Adrb3</i>	Forward	CCAGAAGTCCTGCAAAAACG
	Reverse	CCACCGCTCAACAGGTTT
	Probe:	/5Cy5/CCGTGAAGATCCAGCAAGGAAGCT/3IAbRQSp/
<i>Lepr</i>	Forward	GCTCAGACGTAGGATGAATAGATG
	Reverse	TCACCCAGCACAATCCAAT
	Probe:	/56-FAM/TGAGGTATC/ZEN/ACAGGCGCAGCC/3IABkFQ/
<i>Lipe</i>	Forward	CCATATTGTCTTCTGCGAGTGT
	Reverse	GGCGAAAAGGCAAGATCAAAG
	Probe:	/5Cy5/CGTGCGTAAATCCATGCTGTGTGAGA/3IAbRQSp/
<i>Adipoq</i>	Forward	GCAGGATTAAGAGGAACAGGAG
	Reverse	TGTCTGTACGATTGTCAGTGG
	Probe:	/56-FAM/ACGACACCA/ZEN/AAAGGGCTCAGGAT/3IABkFQ/

## Supplemental Table 2

Litter size effects on weight and composition of pups in the primary cohort

Primary Cohort P0-P22		95% Wald Confidence Interval			
Diet Group * Postnatal Day	Beta	Lower	Upper	Wald Chi-Square	Sig
<b>Body weight</b>					
SL vs NL	0.029	0.012	0.046	11.614	0.001
LL vs NL	-0.077	-0.091	-0.064	122.086	<0.001
LL vs SL	-0.107	-0.121	-0.092	196.400	<0.001
<b>Fat mass</b>					
SL vs NL	0.01	0.003	0.017	7.56	0.006
LL vs NL	-0.019	-0.025	-0.013	36.310	<0.001
LL vs SL	-0.029	-0.035	-0.022	78.972	<0.001
<b>Lean mass</b>					
SL vs NL	0.052	0.033	0.07	30.483	<0.001
LL vs NL	-0.035	-0.051	-0.02	20.136	<0.001
LL vs SL	-0.087	-0.103	-0.071	111.650	<0.001

Primary Cohort P0-P28		95% Wald Confidence Interval			
Diet Group * Postnatal Day	Beta	Lower	Upper	Wald Chi-Square	Sig
<b>Body weight</b>					
SL vs NL	-0.03	-0.047	-0.012	11.266	0.001
LL vs NL	-0.081	-0.095	-0.067	122.857	<0.001
LL vs SL	-0.051	-0.067	-0.036	42.349	<0.001
<b>Fat mass</b>					
SL vs NL	-0.007	-0.013	-0.002	6.214	0.013
LL vs NL	-0.017	-0.022	-0.012	42.824	<0.001
LL vs SL	-0.009	-0.014	-0.004	12.182	<0.001
<b>Lean mass</b>					
SL vs NL	-0.02	-0.038	-0.002	4.715	0.030
LL vs NL	-0.064	-0.079	-0.048	64.183	<0.001
LL vs SL	-0.044	-0.06	-0.027	27.612	<0.001

### Supplemental Table 3

Litter size effects on weight and composition of pups in the replication cohort

REPLICATION COHORT Diet Group * Postnatal Day	95% Wald Confidence Interval			Wald Chi-Square	Sig
	Beta	Lower	Upper		
<b>Body weight</b>					
SL vs NL	0.119	0.104	0.134	239.917	<0.001
LL vs NL	-0.026	-0.04	-0.011	12.735	<0.001
LL vs SL	-0.144	-0.161	-0.128	284.972	<0.001
<b>Fat mass</b>					
SL vs NL	0.014	0.008	0.02	21.133	<0.001
LL vs NL (analysis from P6 to P13)	-0.011	-0.021	-0.002	5.263	0.022
LL vs SL (analysis from P6 to P13)	-0.098	-0.112	-0.085	216.282	<0.001
<b>Lean mass</b>					
SL vs NL	0.094	0.077	0.11	126.988	<0.001
LL vs NL (analysis from P6 to P13)	-0.038	-0.057	-0.019	15.441	<0.001
LL vs SL (analysis from P6 to P13)	-0.16	-0.186	-0.134	148.242	<0.001

**Supplemental Table 4**

Maternal high fat diet (HFD) effects on weight and composition of pups

Diet Group * Postnatal Day	Beta	95% Wald Confidence Interval		Wald Chi-Square	Sig
		Lower	Upper		
<b>Body weight</b>					
HFD at parturition vs Control chow	0.051	0.04	0.063	75.318	<0.001
HFD pre-conception vs Control chow	0.032	0.021	0.043	31.562	<0.001
<b>Fat mass</b>					
HFD at parturition vs Control chow	0.022	0.017	0.027	81.632	<0.001
HFD pre-conception vs Control chow	0.015	0.011	0.02	44.618	<0.001
<b>Lean mass</b>					
HFD at parturition vs Control chow	0.044	0.022	0.067	14.732	<0.001
HFD pre-conception vs Control chow	0.029	0.009	0.049	8.144	0.004