

which should be cited to refer to this work.

1 **Supplemental Fig 1.-** Hypothalamic gene expression of orexigenic (*Npy* and *Agrp*) and anorexigenic  
2 (*Cart* and *Pomc*) neuropeptides of *ob/ob* and C57BL6/J mice subcutaneously treated with oxytocin  
3 (50nmol/day) or vehicle (saline) during 14 days, considering the levels in the C57BL6/J saline-  
4 treated group as a log2 fold change of 0 (A) or 100% (B), N=7-8. \*\* $p < 0.01$ , \*\*\* $p < 0.001$  vs. C57 Sal;  
5 † $p < 0.05$ , †† $p < 0.01$ , ††† $p < 0.001$  vs. C57 Oxt.

6 **Supplemental Fig 2.-** Food intake-dependent and –independent effects of oxytocin, as determined  
7 by using a pair-fed control group. *Ob/ob* mice were treated with vehicle (saline), oxytocin  
8 (50nmol/day) or vehicle (saline), but receiving the same amount of food as oxytocin-treated mice  
9 (PF, pair-fed group) during 14 days. (A) Delta body composition between the beginning and the  
10 end of the treatment measured by magnetic resonance imaging, N=6-20. (B) Epididymal white  
11 adipose tissue (eWAT) weight, N=6-7. (C) eWAT gene expression of lipolytic (*Hsl*), lipogenic (*Fasn*)  
12 lipid uptake (*Lpl*) and glyceroneogenic (*Pepck*) enzymes, as well as of the oxytocin receptor (*Oxtr*),  
13 considering the levels in the *ob/ob* Sal group as 100%, N=6-7. (D) FASN activity in eWAT, N=6-7. (E)  
14 Expression of the macrophage marker *Emr1* in eWAT, considering the levels in the *ob/ob* Sal group  
15 as 100%, N=6. (F) Representative merged immunofluorescence images of the macrophage marker  
16 MAC-2 in the red channel and the nuclear marker, Hoechst 33258 in the blue channel and  
17 quantification of the immunofluorescence in percent of MAC-2 positive cells over all cells present  
18 on the slice, scale white bar: 100 um, N=6-7. (G) Representative hepatic hematoxylin and eosin  
19 staining from animals with the different treatments, scale black bar: 100 um. (H) Quantification of  
20 hepatic triglyceride content, N=6-7. (I) Representative hepatic oil red O staining from animals with  
21 the different treatments, scale black bar: 100 um. (J) Hepatic gene expression of gluconeogenic  
22 (*G6pc*, *Fbp1* and *Pepck*) and glycolytic (*Gck*, *Pfkb1* and *Pklr*) enzymes, N=6-7. (K-L) Glycemia (K,  
23 N=6-17) and insulinemia (L, N=6-7) during a glucose tolerance test. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$   
24 vs. *ob/ob* Sal; † $p < 0.05$ , †† $p < 0.01$  vs. *ob/ob* Oxt.

25 **Supplemental Fig 3** Hepatic glycogen content and related genes in C57BL6/J and *ob/ob* mice. (A-B)  
26 Quantification of hepatic glycogen content, (C-D) Hepatic gene expression of glycogen synthase  
27 (*Gys2*) and phosphorylase (*Pygl*). Mice were *ob/ob* (A and C) or C57BL6/J (B and D) subcutaneously  
28 treated with oxytocin (50nmol/day) or vehicle (saline) during 14 days. N=6-8. \* $p < 0.05$ ,  
29 \*\*\* $p < 0.001$ ..

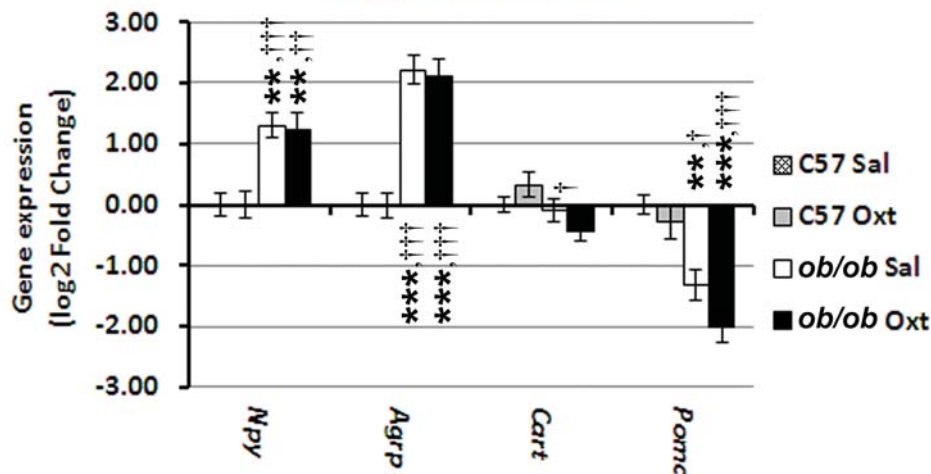
30 **Supplemental Fig 4** *Oxtr* mRNA expression and plasma oxytocin levels in C57BL6/J and *ob/ob* mice.  
31 (A) *Oxtr* levels in different mouse tissues, according to the BioGPS database (24). (B) eWAT gene  
32 expression of *Oxtr* in saline-treated C57BL6/J and *ob/ob* mice, considering the levels in the  
33 C57BL6/J group as 100%, N=7-8. (C) Plasma oxytocin levels of saline-treated C57BL6/J and *ob/ob*  
34 mice, N=5-6. \*\*\* $p < 0.001$ .

35 **Supplemental Table 1.**- Sequence of primers used for the Real Time PCR.

36

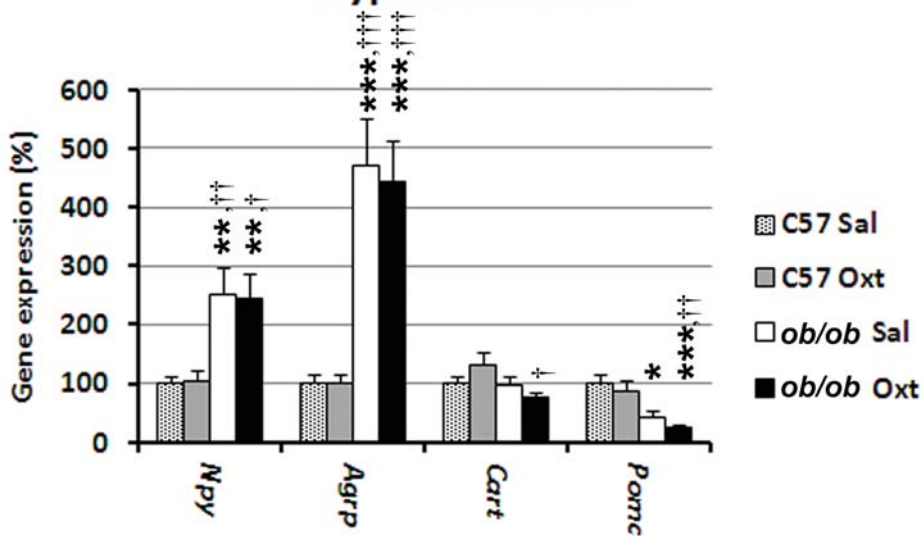
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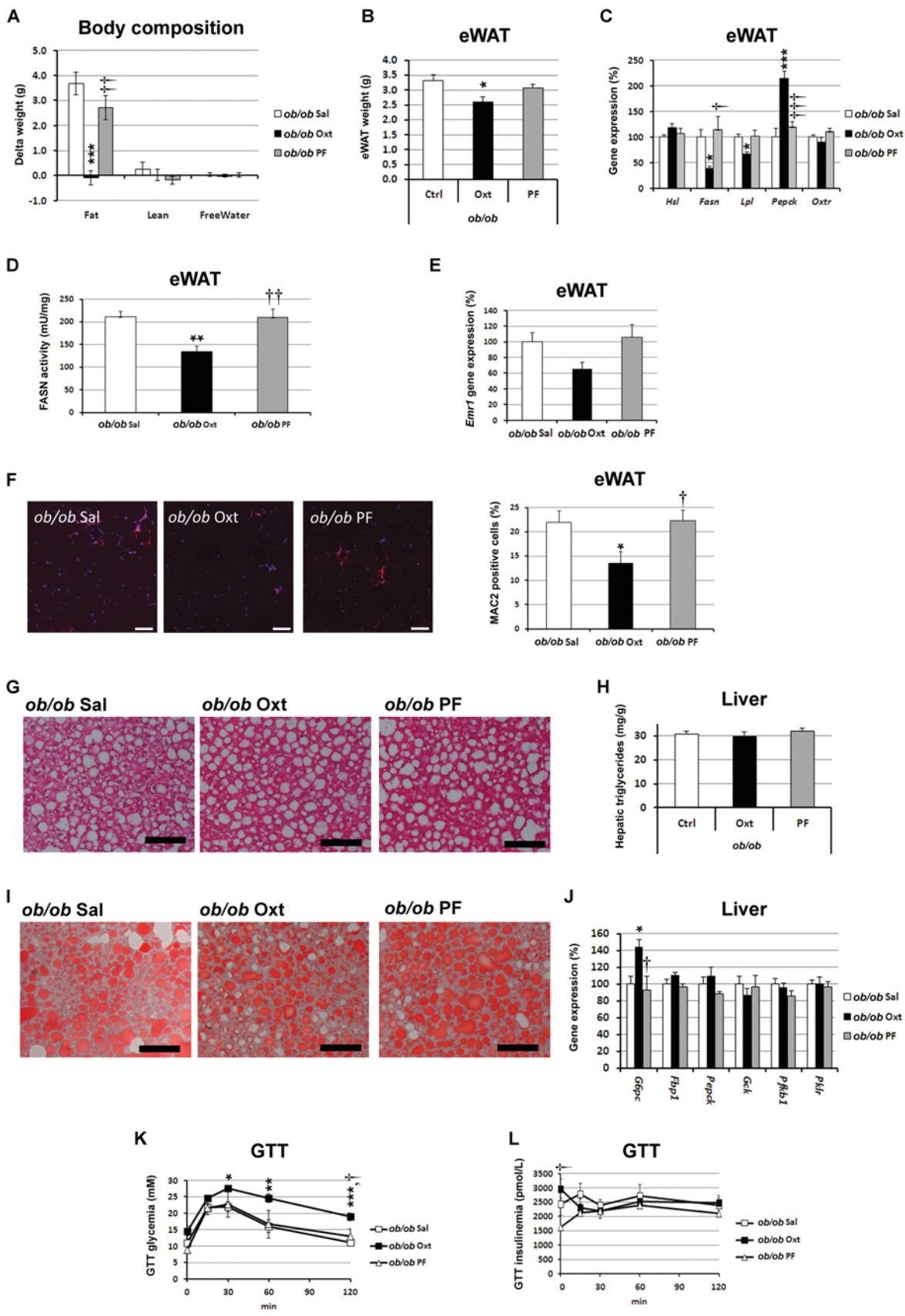
## Hypothalamus



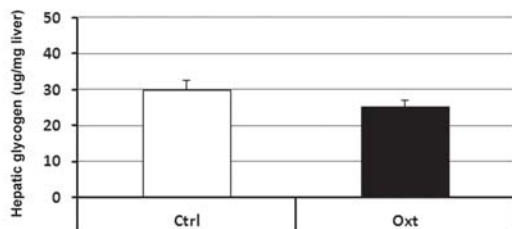
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## Hypothalamus



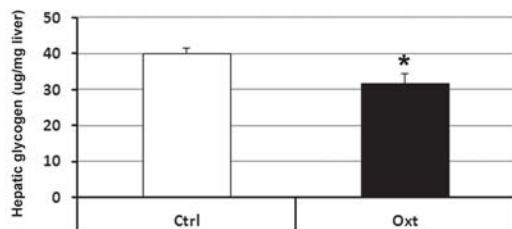


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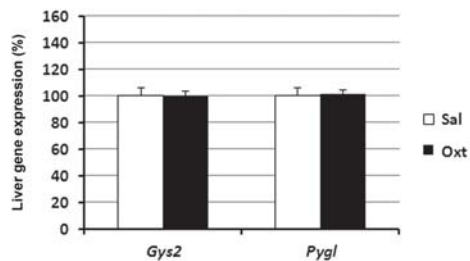
*ob/ob*

B

C57BL6/J

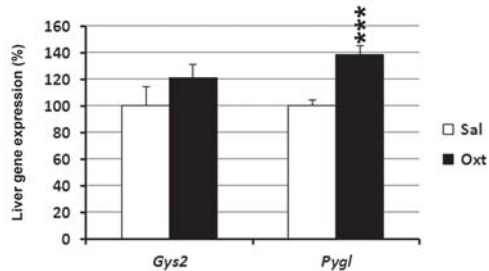


C

*ob/ob*

D

C57BL6/J





Sup Table-Primers

	Genes	Forward	Reverse
<b>White adipose tissue genes</b>	<i>Hsl</i> (official name <i>Lipe</i> )	GGAGCACTACAAACGCAACGA	CCACCGGTAAAGAGGGAACTG
	<i>Fasn</i>	GCCAACCGGCTCTCTTTCTT	GGCTGTGTCCAGGGCAAT
	<i>Lpl</i>	TTCCAGCCAGGATGCAACA	CCACGTCTCCGAGTCCTCTCT
	<i>Oxtr</i>	CATCACCTCCGCTTCTACGG	ATGCCACCACCTGCAAGTA
	<i>Emr1</i> (also known as F4/80)	CAGATACAGCAATGCCAAGCA	GATTGTGAAGGTAGCATTACAAGTG
<b>Liver genes</b>	<i>G6pc</i>	GGAGTCTTGTCAAGCATTGCT	CGGAGGCTGGCATTGTAGAT
	<i>Fbp1</i>	GCACTCTGGTATATGGAGGGATCT	AGCAGCCGCAGCTTTCC
	<i>Pepck1</i> (official name <i>Pck1</i> )	CCACAGCTGCTGCAGAACAC	GAAGGGTCGCATGGCAA
	<i>Pfkb1</i>	TGATCTGTCACCAGGCTGTCA	AGGGCAGCTCATCTGAACTTTT
	<i>Pklr</i>	GAACCATGAAGGCGTGAAGAA	CCCCGAGCCACCATGAT
	<i>Gck</i>	TGGATGGCTCCGTGTACAAG	GATTCGCAGTTGGGTGTCA
	<i>Gys2</i>	GAGTCCTTATCCAGGCTTAATTTCC	GGCAGGCATGATGAAAAACA
	<i>Pygl</i>	CGGTGAACGGTGTAGCAAAGA	CTAGCTCGCTGAAGTCCTGAAT
<b>Hypothalamic genes</b>	<i>Cart</i>	CTGCAATTCTTTCTTGAAGTG	GGGAATATGGGAACCGAAGGT
	<i>Agrp</i>	CCGCTTCTTCAATGCCTTTT	AGGTGCGACTACAGAGGTTTCGT
	<i>Pomc</i>	GCAGAGGCAAACAAGATTGGA	CAGAGAGCTGCCTTTCCGCGACAG
	<i>Npy</i>	AAAACGCCCCCAGAACAAG	CGGGAGAACAAGTTTCATTTCC
<b>Housekeeping gene</b>	<i>Rps29</i>	GCAAATACGGGCTGAACATGT	TCCAACCTTAATGAAGCCTATGTCCTT

Sequence of primers used for the Real Time PCR.