

Supplemental Fig 1.- Hypothalamic gene expression of orexigenic (*Npy* and *Agrp*) and anorexigenic (*Cart* and *Pomc*) neuropeptides of *ob/ob* and C57BL6/J mice subcutaneously treated with oxytocin (50nmol/day) or vehicle (saline) during 14 days, considering the levels in the C57BL6/J saline-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; † $p < 0.05$, †† $p < 0.01$, ††† $p < 0.001$ vs. C57 Oxt.

Supplemental Fig 2.- Food intake-dependent and –independent effects of oxytocin, as determined by using a pair-fed control group. *Ob/ob* mice were treated with vehicle (saline), oxytocin (50nmol/day) or vehicle (saline), but receiving the same amount of food as oxytocin-treated mice (PF, pair-fed group) during 14 days. (A) Delta body composition between the beginning and the end of the treatment measured by magnetic resonance imaging, N=6-20. (B) Epididymal white adipose tissue (eWAT) weight, N=6-7. (C) eWAT gene expression of lipolytic (*Hsl*), lipogenic (*Fasn*) lipid uptake (*Lpl*) and glyceroneogenic (*Pepck*) enzymes, as well as of the oxytocin receptor (*Oxtr*), considering the levels in the *ob/ob* Sal group as 100%, N=6-7. (D) FASN activity in eWAT, N=6-7. (E) Expression of the macrophage marker *Emr1* in eWAT, considering the levels in the *ob/ob* Sal group as 100%, N=6. (F) Representative merged immunofluorescence images of the macrophage marker MAC-2 in the red channel and the nuclear marker, Hoechst 33258 in the blue channel and quantification of the immunofluorescence in percent of MAC-2 positive cells over all cells present on the slice, scale white bar: 100 μ m, N=6-7. (G) Representative hepatic hematoxylin and eosin staining from animals with the different treatments, scale black bar: 100 μ m. (H) Quantification of hepatic triglyceride content, N=6-7. (I) Representative hepatic oil red O staining from animals with the different treatments, scale black bar: 100 μ m. (J) Hepatic gene expression of gluconeogenic (*G6pc*, *Fbp1* and *Pepck*) and glycolytic (*Gck*, *Pfkfb1* and *Pklr*) enzymes, N=6-7. (K-L) Glycemia (K, N=6-17) and insulinemia (L, N=6-7) during a glucose tolerance test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ vs. *ob/ob* Sal; † $p < 0.05$, †† $p < 0.01$ vs. *ob/ob* Oxt.

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

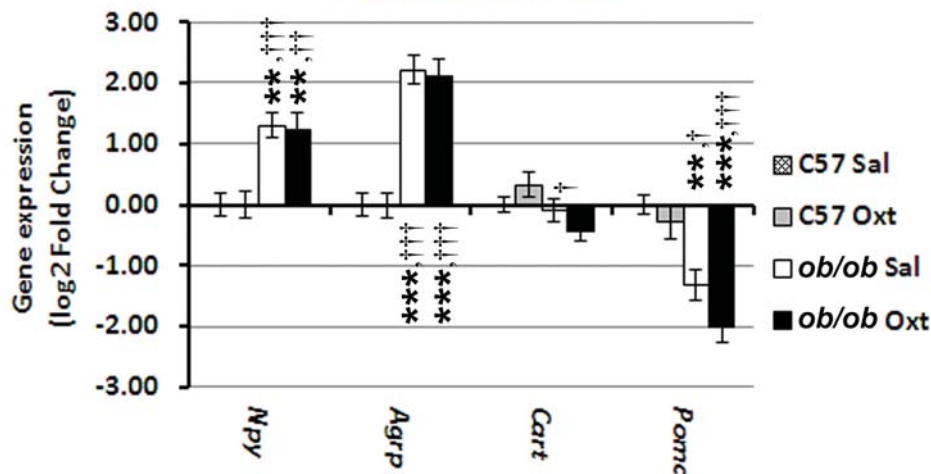
Supplemental Fig 4 *Oxtr* mRNA expression and plasma oxytocin levels in C57BL6/J and *ob/ob* mice. (A) *Oxtr* levels in different mouse tissues, according to the BioGPS database (24). (B) eWAT gene expression of *Oxtr* in saline-treated C57BL6/J and *ob/ob* mice, considering the levels in the C57BL6/J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

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

Sup Fig 1

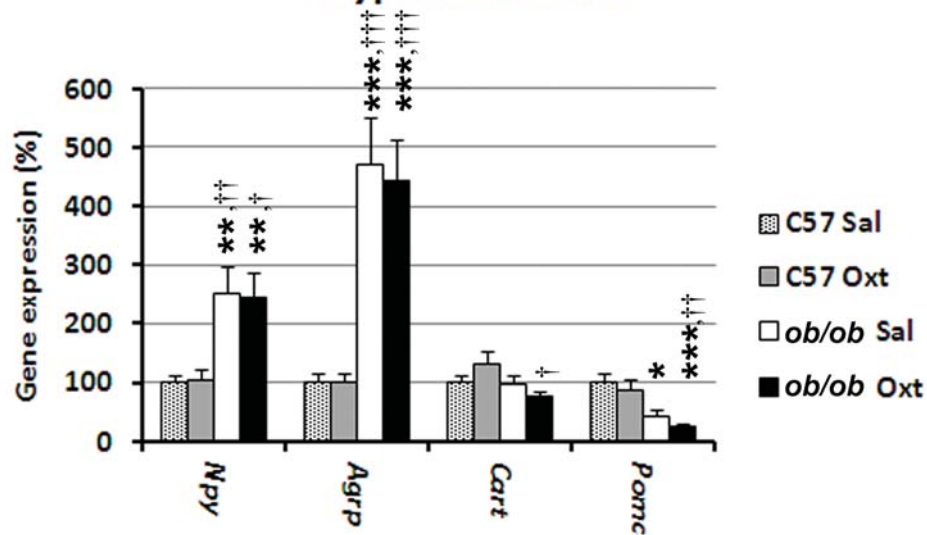
A

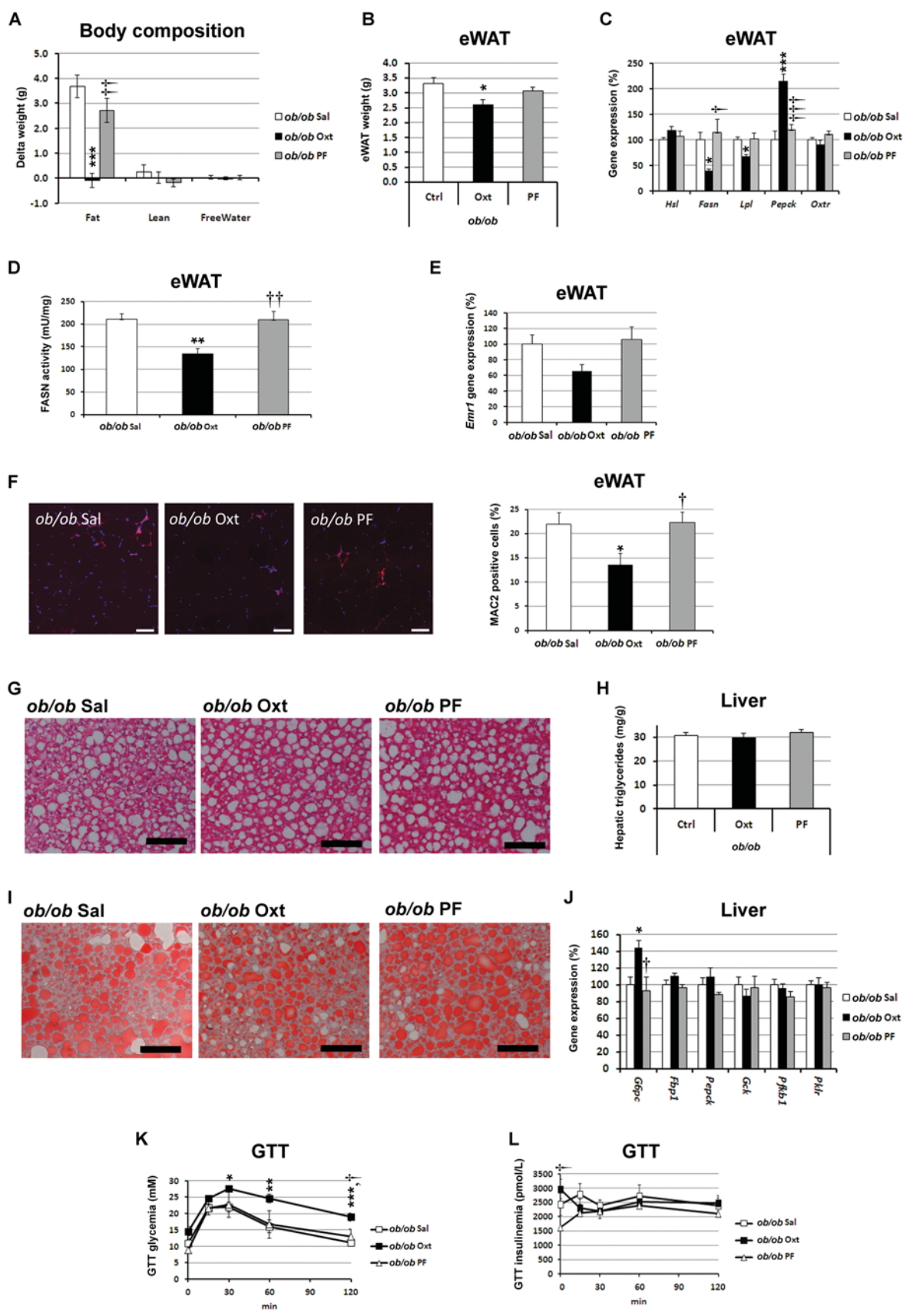
Hypothalamus



B

Hypothalamus

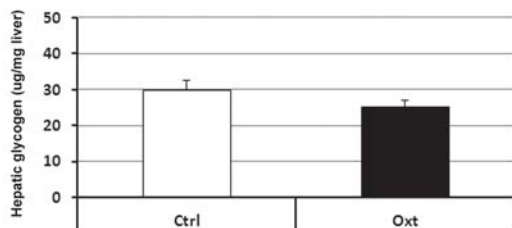




Sup Fig 3

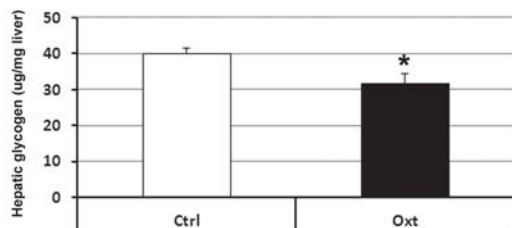
A

ob/ob



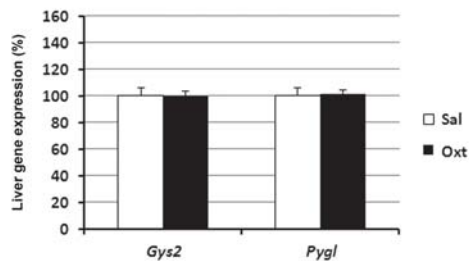
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C57BL6/J



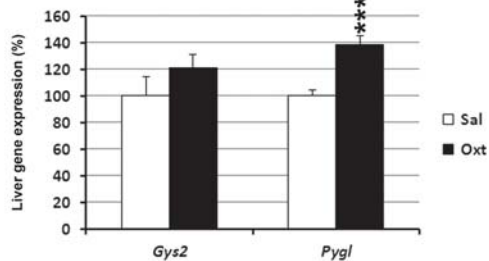
C

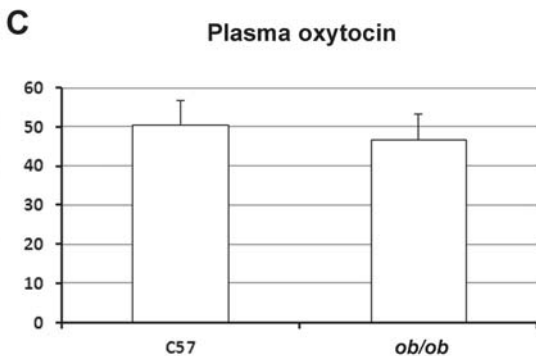
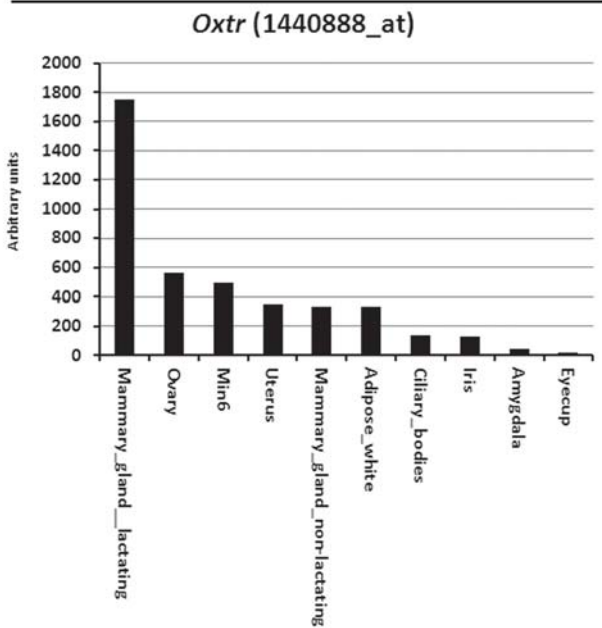
ob/ob



D

C57BL6/J





Sup Table-Primers

| | Genes | Forward | Reverse |
|----------------------------|--|---------------------------|-----------------------------|
| White adipose tissue genes | <i>Hsl</i> (official name <i>Lipe</i>) | GGAGCACTACAAACGCAACGA | CCACCGGTAAAGAGGGAACTG |
| | <i>Fasn</i> | GCCAACCGGCTCTCTTTCTT | GGCTGTGTCCAGGGCAAT |
| | <i>Lpl</i> | TTCCAGCCAGGATGCAACA | CCACGTCTCCGAGTCTCTCT |
| | <i>Oxtr</i> | CATCACCTTCCGCTTCTACGG | ATGCCACCACCTGCAAGTA |
| | <i>Emr1</i> (also known as F4/80) | CAGATACAGCAATGCCAAGCA | GATTGTGAAGGTAGCATTACAAGTG |
| Liver genes | <i>G6pc</i> | GGAGTCTTGTCAAGGATTGCT | CGGAGGCTGGCATTGTAGAT |
| | <i>Fbp1</i> | GCACTCTGGTATATGGAGGGATCT | AGCAGCCGCAGCTTTCC |
| | <i>Pepck1</i> (official name <i>Pck1</i>) | CCACAGCTGCTGCAGAACAC | GAAGGGTCGCATGGCAAA |
| | <i>Pfkfb1</i> | TGATCTGTCACCAGGCTGTCA | AGGGCAGCTCATCTGAACTTTT |
| | <i>Pklr</i> | GAACCATGAAGGCGTGAAGAA | CCCCGAGCCACCATGAT |
| | <i>Gck</i> | TGGATGGCTCCGTGTACAAG | GATTTCGCAGTTGGGTGTCA |
| | <i>Gys2</i> | GAGTCCTTATCCAGGCTTAATTTCC | GGCAGGCATGATGAAAAACA |
| | <i>Pygl</i> | CGGTGAACGGTGTAGCAAAGA | CTAGCTCGCTGAAGTCCTTGAAT |
| Hypothalamic genes | <i>Cart</i> | CTGCAATTCTTTCCTCTTGAAGTG | GGAATATGGGAACCGAAGGT |
| | <i>Agrp</i> | CCGCTTCTTCAATGCCTTTT | AGGTGCGACTACAGAGGTTTCGT |
| | <i>Pomc</i> | GCAGAGGCAAACAAGATTGGA | CAGAGAGCTGCCTTCCGCGACAG |
| | <i>Npy</i> | AAAACGCCCCCAGAACAAG | CGGGAGAACAAGTTTCATTTC |
| Housekeeping gene | <i>Rps29</i> | GCAAATACGGGCTGAACATGT | TCCAACCTTAATGAAGCCTATGTCCTT |

Sequence of primers used for the Real Time PCR.