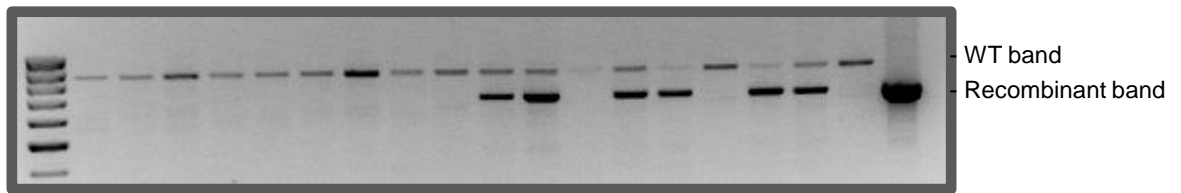
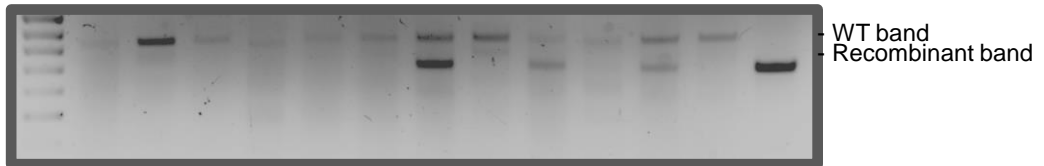
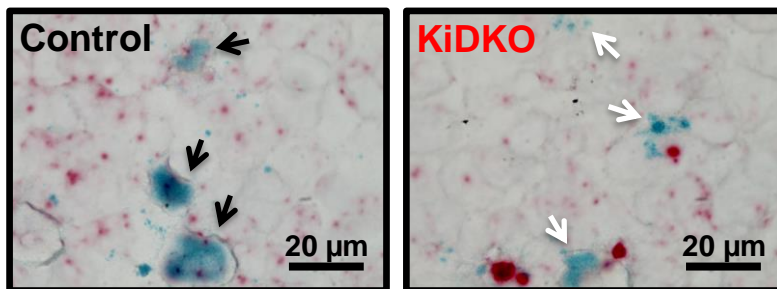
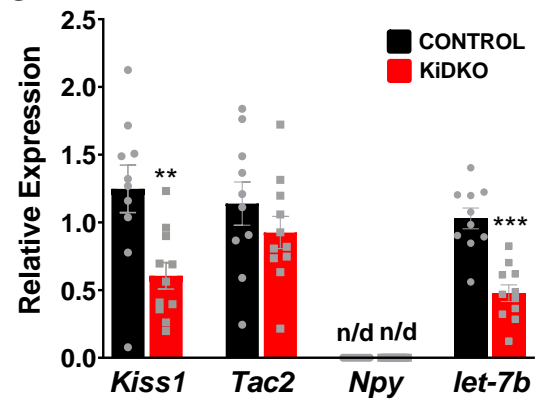
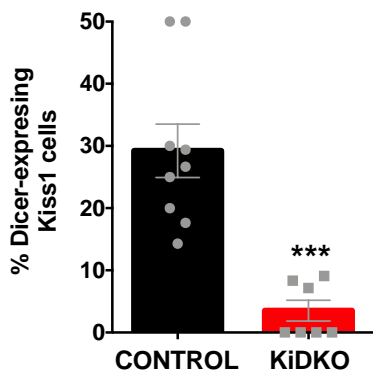
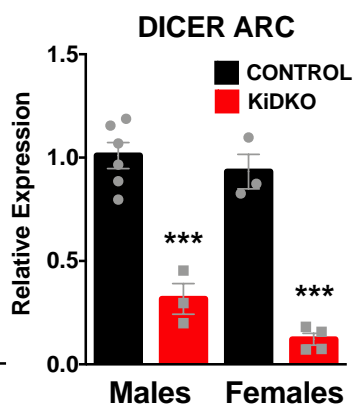
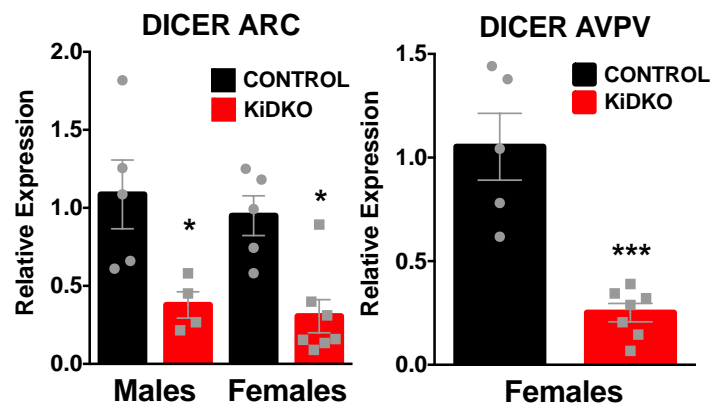


a**PND28**

100-bp POA ARC CX POA ARC CX POA ARC CX POA ARC CX POA ARC CX POA ARC CX C+
ladders
Mouse 1 Mouse 2 Mouse 3 Mouse 1 Mouse 2 Mouse 3

Control**KiDKO****PND1**

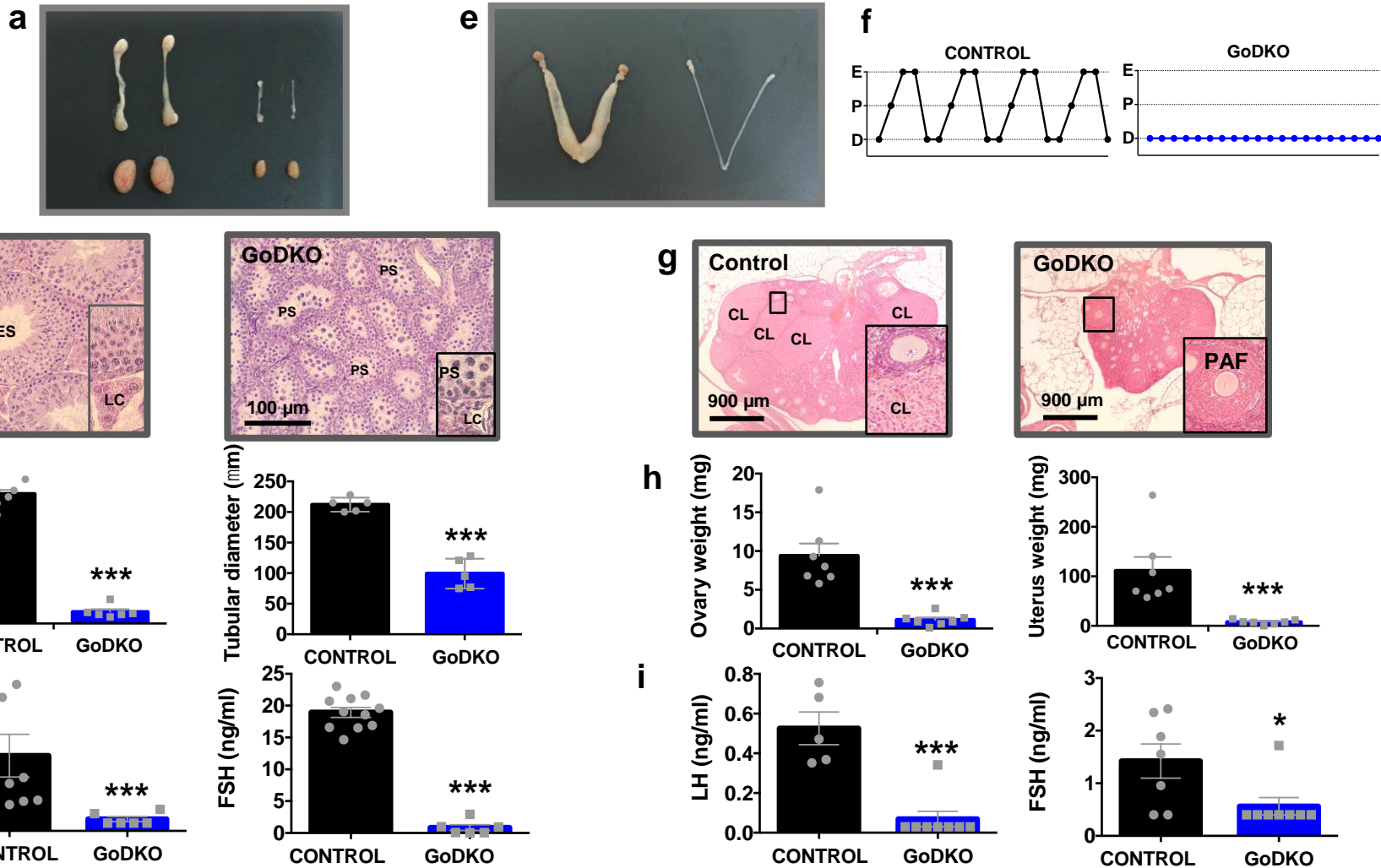
100-bp HTAL CX HTAL CX HTAL CX HTAL CX HTAL CX HTAL CX C+
ladders
Mouse 1 Mouse 2 Mouse 3 Mouse 1 Mouse 2 Mouse 3

Control**KiDKO****b****c****d****e****PND14****PND28**

Suppl. Figure S1. Generation and validation of the *KiDKO* mouse line. Ablation of *Dicer* in *Kiss1* neurons was achieved in vivo using a Cre-LoxP approach, by crossing a *Kiss1*-Cre mouse line with a *Dicer*^{loxP/loxP} mouse line. Panel (a) illustrates PCR detection of the recombinant event leading to *Dicer* allele deletion in the preoptic area (POA) and arcuate nucleus (ARC), areas with prominent *Kiss1* neuronal populations, but not in the cortex (as tissue devoid of *Kiss1* cells), in 28 day-old mice (PND28). In addition, recombination was assessed also in the hypothalamus at PND1. An amplicon of expected size, denoting effective recombination, was detected in the ARC, POA and whole hypothalamus of *KiDKO* mice, whose identity was confirmed by direct sequencing. In contrast, no amplification was observed in the cortex of *KiDKO* mice, neither in the POA or cortex of control mice. Three individuals per genotype were analyzed. In addition, in panel b, results of dual labeling, using BaseScope™ in situ hybridization in OVX females at PND54, are presented. Note that, while in controls, *Kiss1* cells (labelled in green) are shown to co-express *Dicer* exon 23 (labelled in red; see *black arrows*), in *KiDKO* mouse hypothalamus, such co-expression drops to virtually negligible levels, despite detectable kisspeptin content. Scales In panel (c), TaqMan qPCR analyses in FACS isolated *Kiss1* neurons from peripubertal (PND28) *KiDKO*-YFP mice are shown. Despite preserved expression of *Tac2* gene in ARC *Kiss1* neurons, *Kiss1* mRNA levels were partially decreased and expression of an abundant miRNA, namely *let-7b*, was markedly suppressed in *KiDKO* mice, the latter attesting for effective ablation of the miRNA synthesizing enzyme, *Dicer*, in our mouse line. In addition, *Npy* mRNA was not detected in qPCR assays, demonstrating the absence of contamination with other abundant ARC cells in FACS isolated *Kiss1* neuronal samples. Group sizes: n= 10 control; n=11 *KiDKO*. On the other hand, single-cell analysis of FACS isolated *Kiss1* neurons demonstrated a dramatic reduction in the percentage of *Kiss1* neurons expressing *Dicer* in *KiDKO* mice (d). Group sizes: n= 9 control; n=7 *KiDKO* animals. In the same context, *Dicer* expression was significantly reduced in ARC *Kiss1* neurons isolated by FACS from infantile (PND14) and peripubertal (PND28) *KiDKO* male and female mice, and in AVPV *Kiss1* neurons from peripubertal females (e). Group sizes: PND14 (n= 6 control; n=3 *KiDKO* males and n= 3 control; n=4 *KiDKO* females); PND28 (n= 5 control; n=4 *KiDKO* males and n= 5 control; n=7 *KiDKO* females). The values are represented as the mean ± SEM. * P < 0.05; **P < 0.01; ***P < 0.001 vs. corresponding control groups. nd= not detectable.

MALES

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Suppl. Figure S2. Profound hypogonadotropic hypogonadism in adult GoDKO mice. Representative images of the gonads (testes, ovaries) and sex organs (epididymis, uterus) for adult control and GoDKO mice of both sexes are presented in (a; males) and (e; females). For the latter, individual profiles of ovarian cyclicity, monitored by vaginal cytology, are also shown in (g). In addition, representative histological sections of the testis (b) and ovary (g) are presented for control and GoDKO mice. Control mice showed full spermatogenesis, denoted by the presence abundant elongated spermatids (ES), and prominent Leydig cells (LC) in the interstitium. In contrast, KO mice showed atrophic seminiferous tubules with spermatogenic arrest at the primary spermatocyte (PS) level, as well as poorly differentiated Leydig cells. In the case of the ovary, while control females showed cycling ovaries with abundant growing follicles and corpora lutea (CL), in GoDKO females large antral follicles and corpora lutea were not found in KO mice, while preantral (PAF; see inset) or early antral follicles were the most advanced healthy growing follicles. In addition, testicular weight and tubular diameter in males (c; control n=9, GoDKO n=6 for testis weigh and control n=5, GoDKO n=5 for tubular diameter), and ovarian and uterus weight in females (h; control n=7, GoDKO n=7) are presented for control and GoDKO mice, for which serum LH and FSH levels are also shown (d; control n=11; GoDKO n=6 for both LH and FSH, i; control n=5 for LH and control n=7 for FSH; GoDKO n=8 for both LH and FSH). The values are represented as the mean \pm SEM. * $P < 0.05$; *** $P < 0.001$ vs. corresponding control groups.

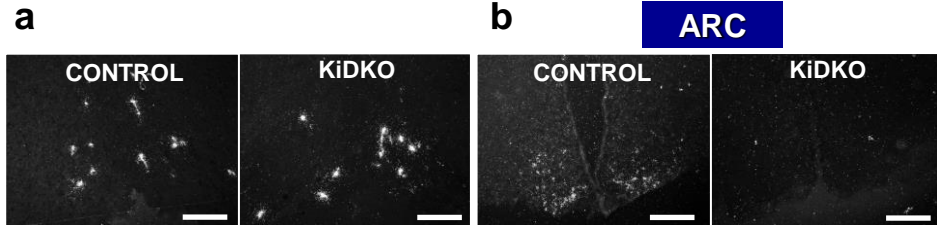
MALES

KiDKO

GnRH mRNA

Kiss1 mRNA

ARC

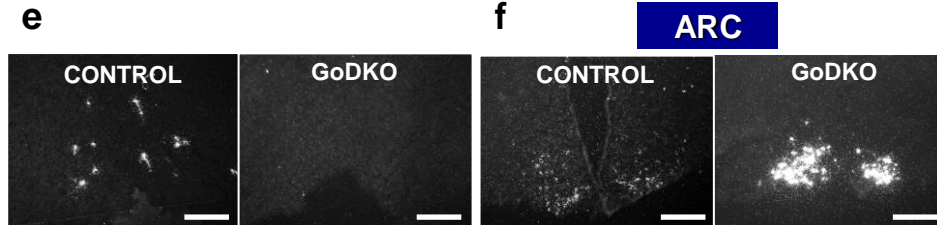


GoDKO

GnRH mRNA

Kiss1 mRNA

ARC

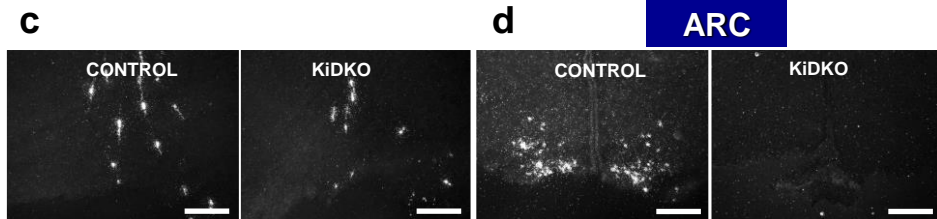


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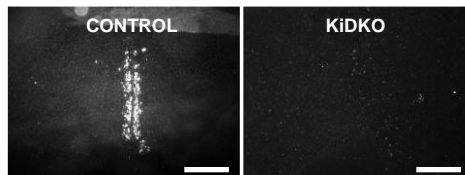
GnRH mRNA

Kiss1 mRNA

ARC



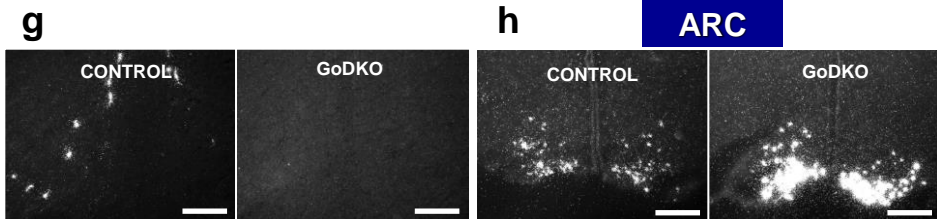
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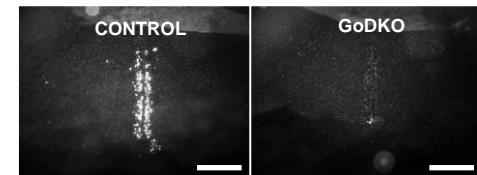
GnRH mRNA

Kiss1 mRNA

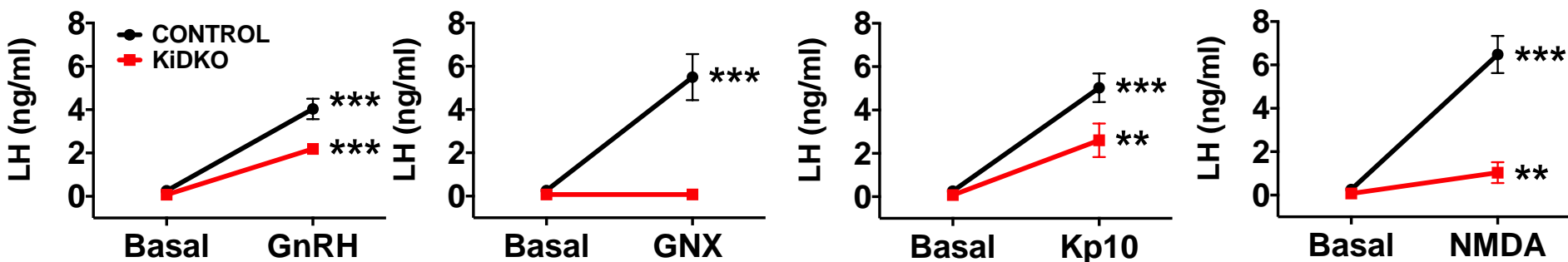
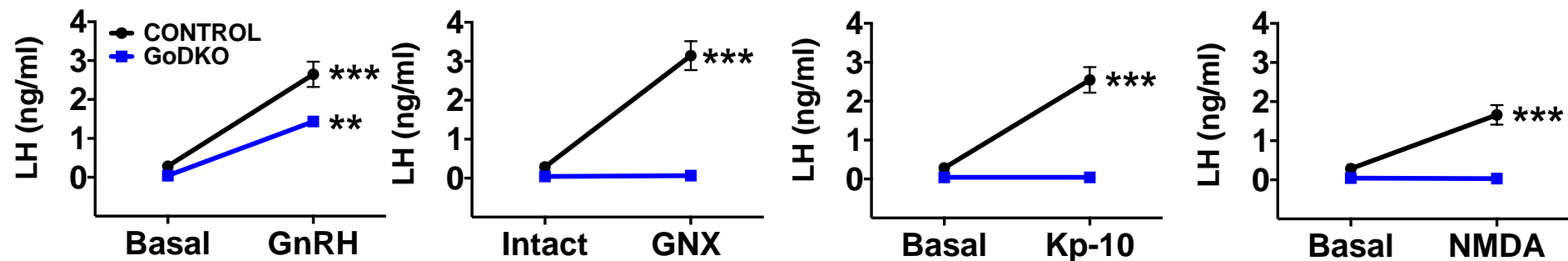
ARC



AVPV



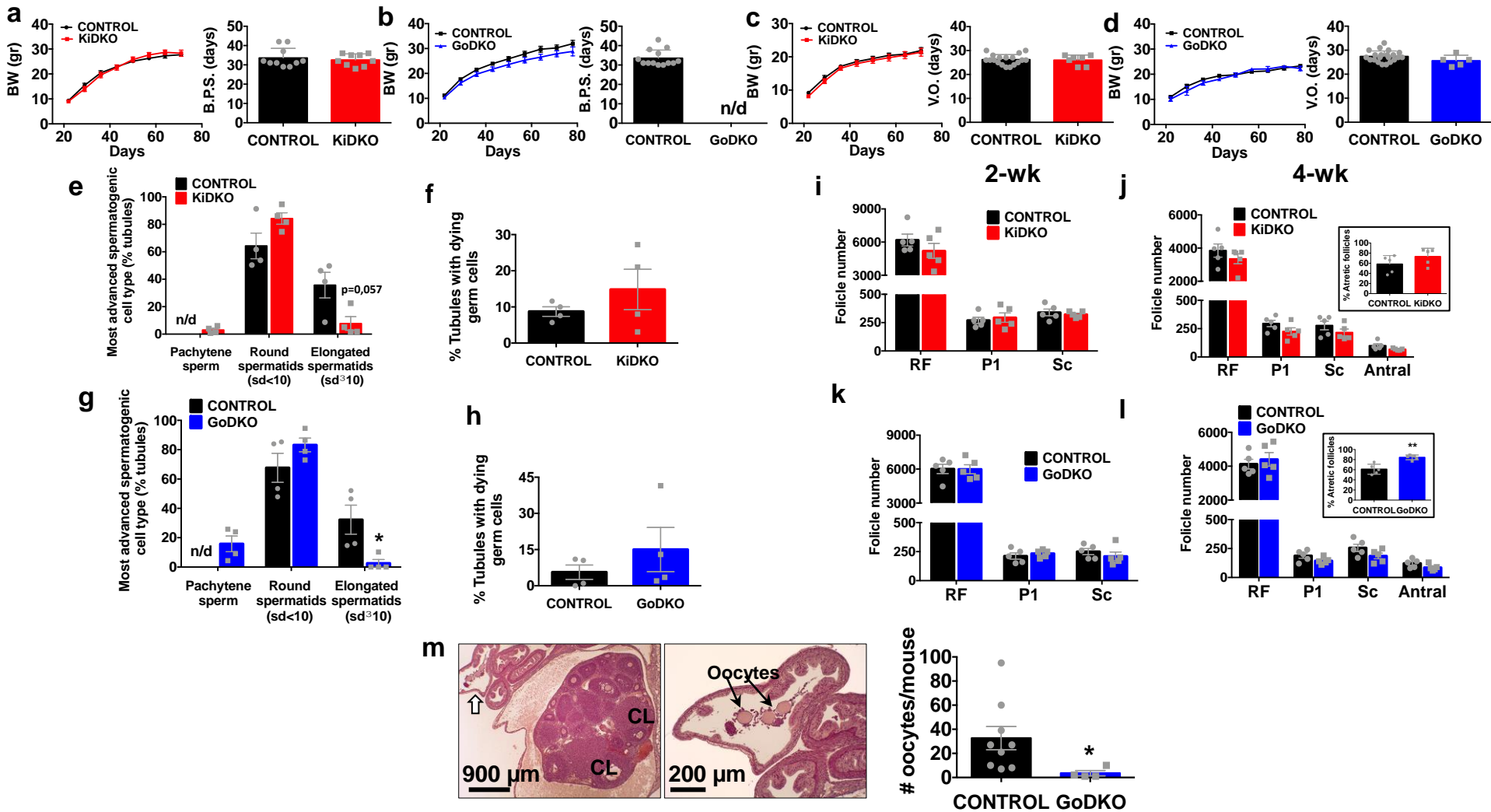
Suppl. Figure S3. Changes in hypothalamic *Kiss1* and *GnRH* expression in *KiDKO* and *GoDKO* mice. Expression of *Kiss1* and *GnRH* mRNAs in specific hypothalamic areas (*Kiss1*: ARC and AVPV; *GnRH*: POA) of the two mouse lines was assessed by in situ hybridization (ISH). POA *GnRH* expression was fully preserved in male (a) and female (c) adult *KiDKO* mice, whereas in adult *GoDKO* animals it was totally absent (e, males; g, females). In turn, *Kiss1* mRNA levels were undetectable in the ARC of male (b) and female (d) *KiDKO* mice, as well as in the AVPV of female *KiDKO* mice (d). In contrast, enhanced *Kiss1* expression was detected in the ARC of male (f) and female (h) *GoDKO* mice, possibly due to the lift of the sex steroid negative feedback caused by hypogonadism in this line. Such decrease in ovarian steroids is likely causative for the substantial suppression of *Kiss1* mRNA levels seen in the AVPV of female *GoDKO* mice (h). Scale bars correspond to 200 μ m.

a**b**

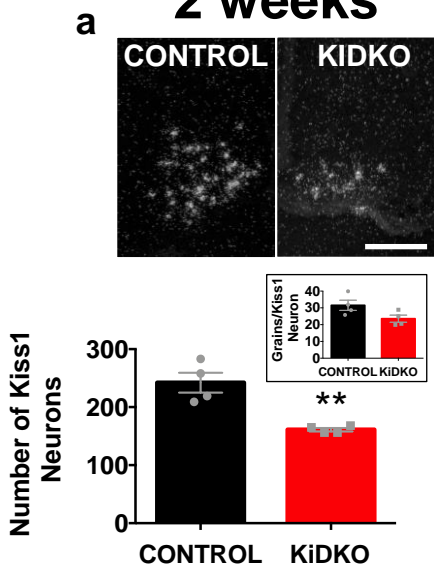
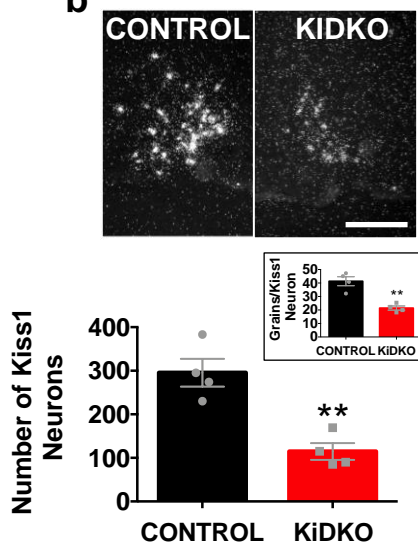
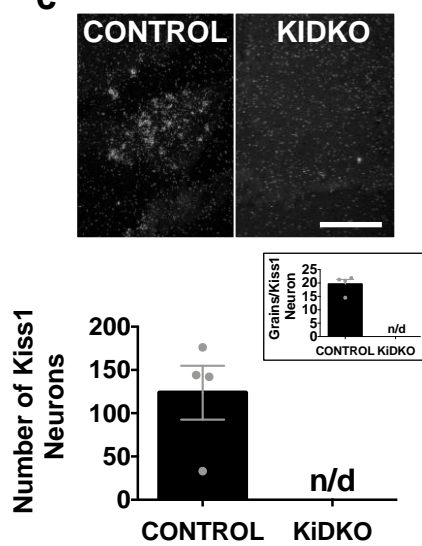
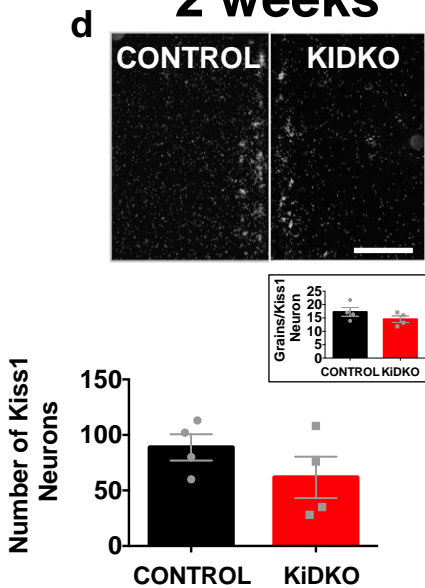
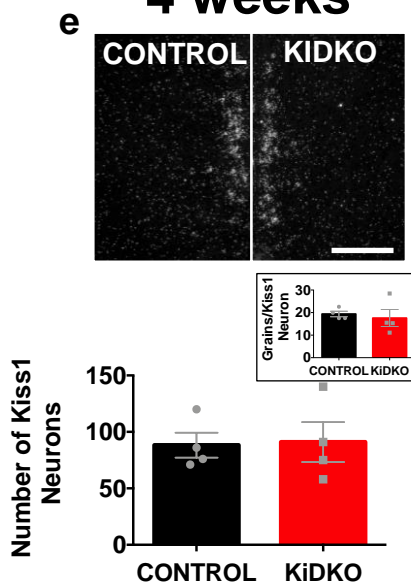
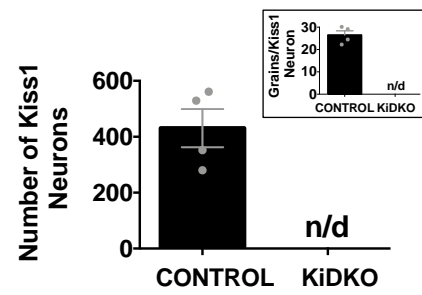
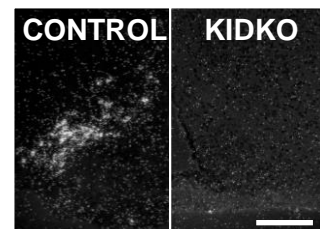
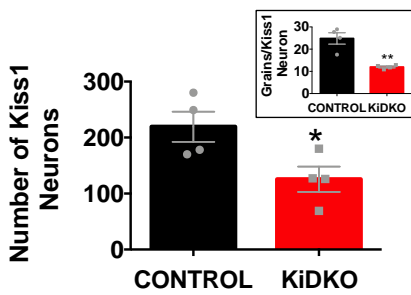
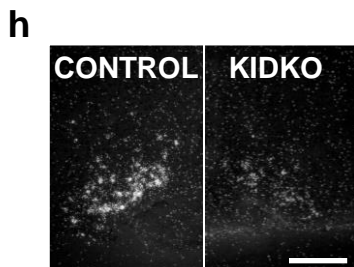
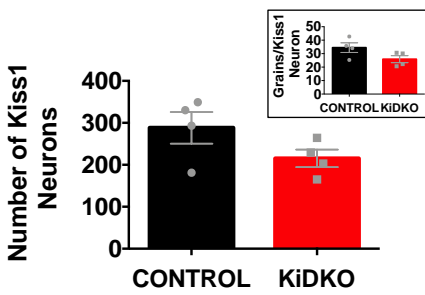
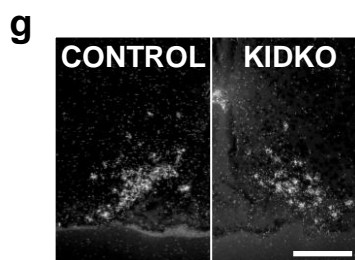
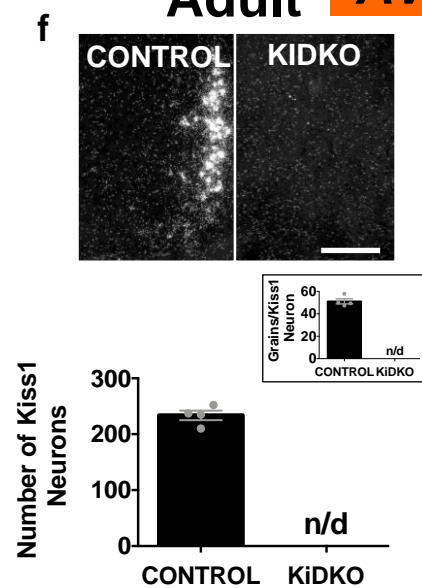
Suppl. Figure S4. LH responses to different neuroendocrine stimuli in adult KiDKO and GoDKO male mice. In upper panels (a), LH levels in adult KiDKO mice, at basal conditions and after administration of a single bolus of GnRH (0.25 μ g, ip), Kp-10 (1 nmol, icv) or NMDA (1 nmol, icv) are presented. In addition, changes in serum LH levels, 2-weeks after gonadectomy (GNX), in KiDKO mice are shown. Similar analyses were conducted in adult GoDKO mice, and data are presented in the lower panels (b). Group sizes, for the KiDKO line: n=6 control in basal conditions and n=7 control after injection; n=5 KiDKO in basal conditions and n=8 KiDKO after injection; for the GoDKO line: n=11 control in basal conditions and n=10 control after injection; n=6 GoDKO in basal conditions and n=5 GoDKO after injection. The values are represented as the mean \pm SEM. **P < 0.01; ***P < 0.001 vs. corresponding basal groups.

MALES

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Suppl. Figure S5. Reproductive indices in *KiDKO* and *GoDKO* mice at puberty and adulthood. Data from the two models, *KiDKO* (*Kiss1*-specific Dicer KO) and *GoDKO* (*GnRH*-specific Dicer KO), are presented. Postnatal body weight (BW) gain and balano-preputial separation (BPS, as external marker of puberty) are presented for male *KiDKO* (**a**; control n=10; *KiDKO* n=10) and *GoDKO* (**b**; control n=11, *GoDKO* n=8) mice, and their corresponding controls. In addition, quantitative parameters of spermatogenesis, including the most advanced spermatogenic cell type and % of tubules with dying cells, are presented for *KiDKO* (**e**, **f**; control n=4; *KiDKO* n=4) and *GoDKO* (**g**, **h**; control n=4; *GoDKO* n=4) males. In the case of females, postnatal BW gain and vaginal opening (VO, as external marker of puberty) are presented for *KiDKO* (**c**; BW: control n=5, *KiDKO* n=7; and VO: control n=18, *KiDKO* n=7) and *GoDKO* (**d**; control n=21; *GoDKO* n=5) mice, and their corresponding controls. In addition, quantitative parameters of folliculogenesis, including follicle numbers at 2-wk and 4-wk of postnatal life, as well as the % of atretic follicles (see *insets*), are shown for *KiDKO* (**i**, **j**; control n=5; *KiDKO* n=5) and *GoDKO* (**k**, **l**; control n=5; *GoDKO* n=5) females. Finally, in panel (**m**), representative histological images of ovaries of *GoDKO* female mice after ovulation induction by standard gonadotropin priming are shown, with appearance of newly formed corpora lutea (CL) and released oocytes in the uterine tubes (denoted by *arrows*). Quantification of the #oocytes observed in control and *GoDKO* females after gonadotropin priming is also displayed (control n=9; *GoDKO* n=4). The values are represented as the mean \pm SEM. * P < 0.05; **P < 0.01 vs. corresponding control groups. nd= not detectable.

MALES**2 weeks****4 weeks****Adult****ARC****FEMALES****2 weeks****4 weeks****Adult****AVPV****ARC**

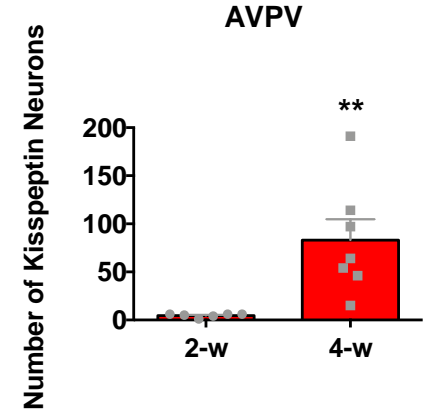
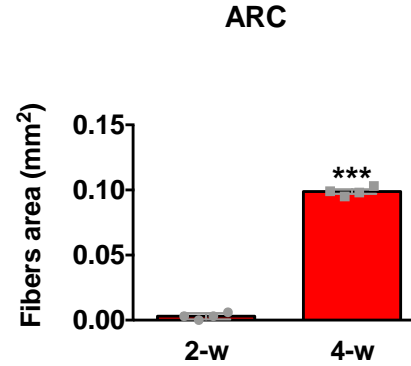
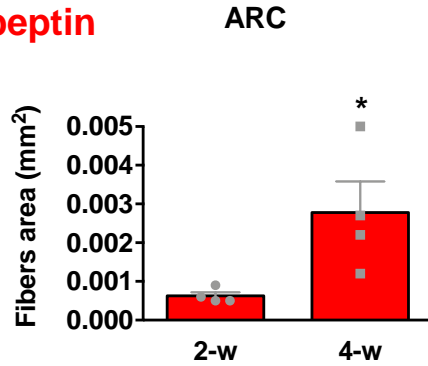
Suppl. Figure S6. *Hypothalamic Kiss1 expression in KiDKO mice along postnatal maturation.* Expression of *Kiss1* mRNA at the ARC and AVPV was assessed by in situ hybridization (ISH). This procedure allowed counting of numbers of *Kiss1*-expressing cells, as well as the grain density per cell (as proxy marker of individual expression). Representative images and quantitative data on #*Kiss1*-expressing neurons and *Kiss1* grain density per cell, are shown in the ARC of males (**a-c**), and the AVPV (**d-f**) and ARC (**g-i**) of females, of control and KiDKO genotypes. Data were collected at three postnatal ages: 2-weeks (corresponding to mini-puberty); 4-weeks (corresponding to early pubertal transition); and adulthood (4-months). Group sizes: n=4 control males; n=4 KiDKO males; n=4 control females; n=4 KiDKO females. The values are represented as the mean \pm SEM. * P < 0.05; **P < 0.01 vs. corresponding control groups. Scale bars correspond to 200 μ m.

KiDKO

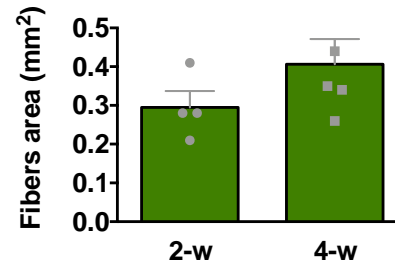
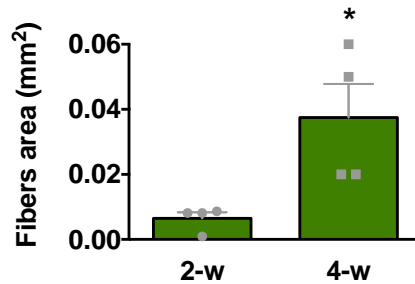
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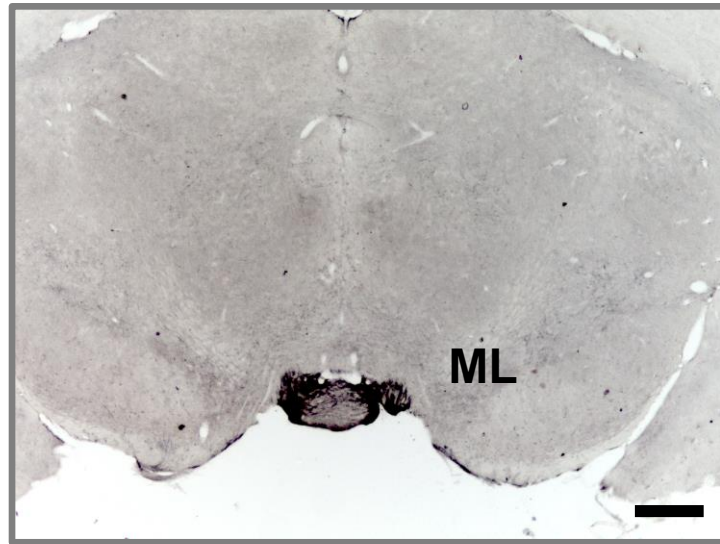
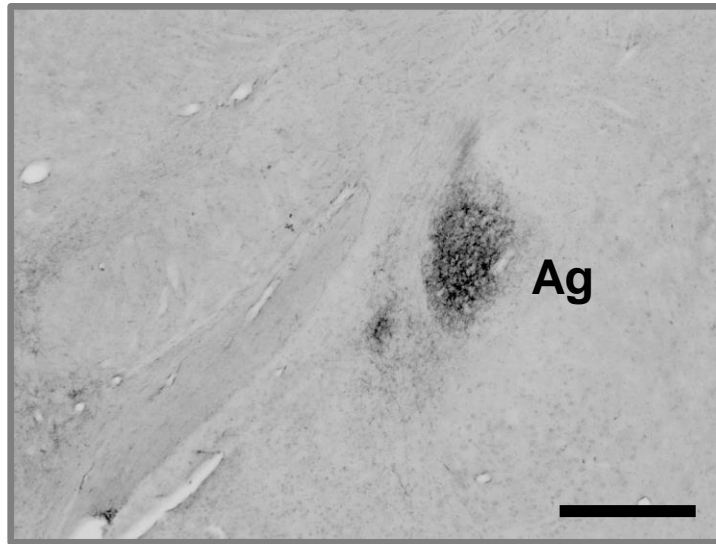
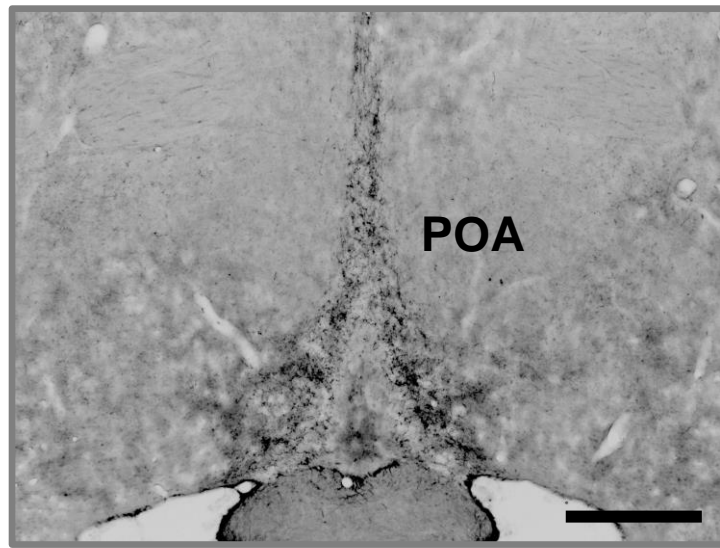
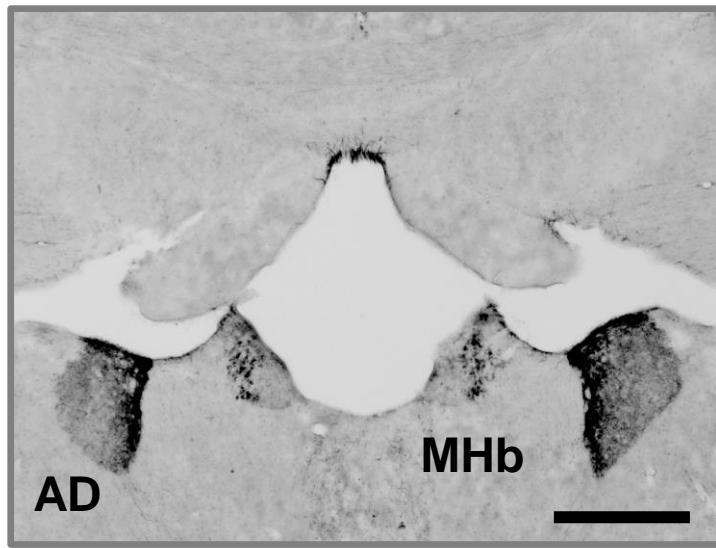
Kisspeptin



NKB



Suppl. Figure S7. Changes in kisspeptin and NKB levels in KiDKO mice during the pubertal transition. Quantification of changes in kisspeptin- (in red) and NKB- (in green) immunoreactivity (IR) in the ARC of male and female KiDKO mice, and of kisspeptin-IR in the AVPV of female KiDKO mice, between 2-wk and 4-wk of postnatal age. Group sizes for ARC densitometry: n = 4 control males; n = 4 KiDKO males; n = 4 control females; n = 4 KiDKO females. Of note, for quantification of kisspeptin cell number in the AVPV, a higher number of animals (n=6 for 2- and n=7 for 4-weeks), was included due to the greater variability. The values are represented as the mean \pm SEM. * P < 0.05; **P < 0.01; ***P < 0.001 vs. corresponding 2-wk group.



MHb: Medial habenular nucleus; **AD:** Anterodorsal nucleus; **POA:** Preoptic area;
Ag: Amygdala; **ML:** Medial Mammillary nucleus

Suppl. Figure S8. *Persistent NKB expression in other brain areas of KiDKO mice.* Representative images of NKB immunoreactivity in various brain areas of KiDKO mice, where NKB expression has been previously reported, including the medial habenular nucleus (MHb), the antero-dorsal nucleus (AD), the preoptic area (POA), and the medial mammillary nucleus (ML). Scale bars correspond to 200 μ m.

Suppl. Table S1. Sequences of primers for in situ hybridization (ISH)

| Riboprobe | GenBank Accession number | Primers | Sequence | Spaning Region |
|------------------|-------------------------------------|----------------|--|---------------------------|
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| | | | Kiss1 reverse 5'-TCA GCC CCG TGC TGC CCG CGC-3' | |
| | | Internals | T3-Kiss1 forward 5'-CAG AGA TGC AAT TAA CCC TCA CTA AAG GGA GAG TGA AGC CTG GAT CCA CAG GC-3' | 111...397 |
| | | | T7-Kiss1 reverse 5'-CCA AGC CTT CTA ATA CGA CTC ACT ATA GGG AGA GCC TGC CTC CTG CCG TAG CGC-3' | |
| GnRH | NM_008145.2 | Externals | GnRH forward 5'-GTT GAC TGT GTG TTT GGA AGG C-3' | 137...371 |
| | | | GnRH reverse 5'-CTT CTT CTG CCC AGC TTC CTC-3' | |
| | | Internals | T3-GnRH forward 5'-CAG AGA TGC AAT TAA CCC TCA CTA AAG GGA GAA GCA CTG GTC CTA TGG GTT GCG-3' | 170...344 |
| | | | T7-GnRH reverse 5'-CCA AGC CTT CTA ATA CGA CTC ACT ATA GGG AGA CAG ACG TTC CAG AGC TCC TCG-3' | |