

Suppl Fig. 1 Assessment of *Prl* mRNA expression in guinea pig decidual tissues. RT-PCR analysis for *Prl* mRNA in the pituitary gland and uterus of a cycling female guinea pig (NP) and in uterine decidual tissue on gestation d15 and d25. Note that *Prl* transcripts were not detected in d15 and d25 guinea pig decidua.

Suppl Fig. 2 Immunocytochemical localization of PRL and GH in the guinea pig anterior pituitary. **A)** Identification of PRL immunopositive cells within the anterior pituitary of the guinea pig using antibodies to porcine PRL. **B)** Demonstration of the specificity of the PRL immunoreactivity within the guinea pig anterior pituitary by competition with recombinant guinea pig PRL. **C)** Identification of GH immunopositive cells within the anterior pituitary of the guinea pig using antibodies to porcine GH. Immunoreactive cells were identified by incubation with TRITC conjugated secondary antibody. Tissues were counterstained with DAPI.

NP-Pituitary

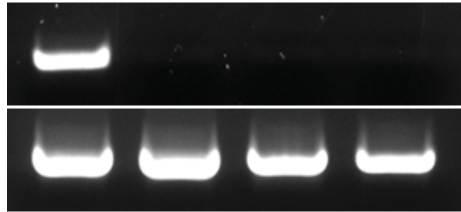
NP-Uterus

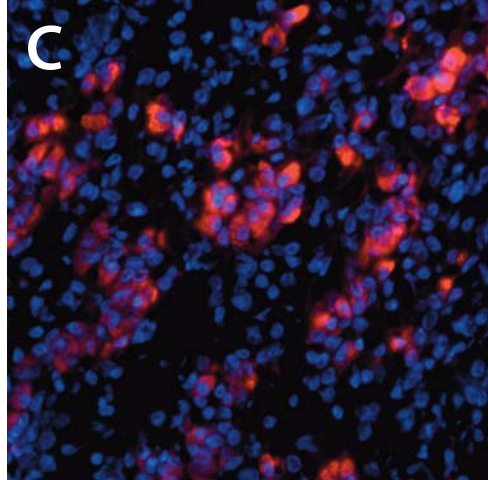
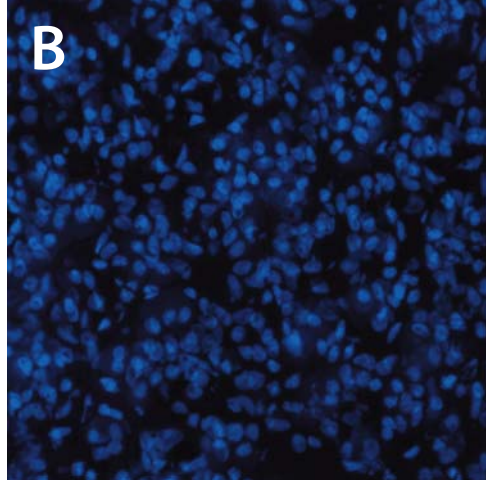
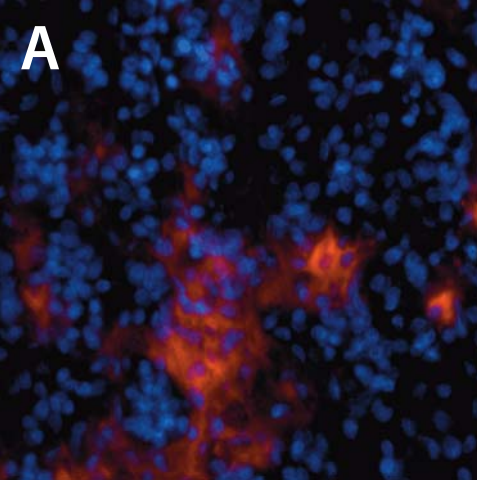
d15 Decidua

d25 Decidua

Prl

18S





Suppl Table 1. Guinea pig primer sequences.

GENE	FORWARD PRIMER	REVERSE PRIMER
<i>Prl</i>	F1: 5'-CCATTACATCCACAGCCTCT-3' F2: 5'-CTACTCTGTCTGGCTGGGAC-3' F3: 5'-CATTTAAAGAATTCACAGCCCATGTGTCCCAGT-3'	R1: 5'-GGAACCTTGAGATAATTGTCGA-3' R2: 5'-CATGGCCTGGGTAAAAAACT-3' R3: 5'-GGAGAGGCTGTGGATGTAAT-3' R4: 5'-GGATCGCTTTTATACCCTTACA-3' R5: 5'-GTTATATGTGGATCCTTAGCAGCTGCCATCATA 3'
<i>Prirp1</i>	F1: 5'-GCTTCCCAGTCCAAATGTATT3' F2: 5'-GGCCTTCAAGACACAGACTAT-3' F3: 5'-GATTTCAATTGATACTTTTCACCA-3' F4: 5'-CATTTAAAGAATTCACAGTCCAAATGTATTG-3'	R1: 5'-GGTAAAAAGTATCAATGAAAT-3' R2: 5'-GGAGGCTGTATGACAACCTT-3' R3: 5'-GTGAGGACGTGGATATTGTCA-3' R4: 5'-CATTTGGACTGGGAAGCTACA-3' R5: 5'-GTTATATGTGGATCCTTAGTTGTGTCTATCA-3'
<i>Prirp2</i>	F1: 5'-CCTGATACTGCTCATGTCAA-3' F2: 5'-GTGGAAAATATTAGGAGAAGAA-3' F3: 5'-GACTTCCATTCCTGAGATCAT-3'	R1: 5'-CAGATGATCTCAGGAATGGAA-3' R2: 5'-CCTTGAACAGCTCTGAAGAGA-3' R3: 5'-GCTCGATCAAAAAGGTAAGGAA-3' R4: 5'-CTTGACATGAGCAGTATCAGGA-3'
<i>Gh</i>	5'-CAGTCCCGAAACCAGTCAG-3'	5'-TATGATGCAGCTCAGTTTTA-3'
18S	5'-ATGCCAGAGTCTCGTTCGTT-3'	5'-CGCGTTCTATTTTGTGGT-3'

Suppl Table 2. GH sequence analysis.

Genus species	Common Name	Accession No.	Amino acids	Percent identity with <i>Cavia porcellus</i>
<i>Cavia porcellus</i>	Guinea pig	AAF36409	216	100
<i>Canis familiaris</i>	Dog	NP_001003168	216	89
<i>Felis catus</i>	Cat	NP_001009337	216	89
<i>Sus scrofa</i>	Pig	NP_999034	217	87
<i>Equus caballus</i>	Horse	NP_001075417	216	87
<i>Mesocricetus auratus</i>	Golden hamster	AAB20368	216	87
<i>Bos taurus</i>	Cow	NP_851339	217	86
<i>Rattus norvegicus</i>	Rat	NP_001030020	216	85
<i>Ovis aries</i>	Sheep	NP_001009315	217	84
<i>Mus musculus</i>	Mouse	NP_032143	216	84
<i>Monodelphis domestica</i>	Opossum	NP_001028165	215	79
<i>Gallus gallus</i>	Chicken	NP_989690	216	71
<i>Macaca mullata</i>	Rhesus monkey	NP_000506	217	66
<i>Homo sapiens</i>	Human	NP_001036203	217	65
<i>Xenopus laevis</i>	African clawed frog	NP_001083848	214	59