

1 **Pituitary Gonadotrophic Hormone Synthesis, Secretion, Subunit Gene Expression and Cell**
2 **Structure in Normal and FSH β KO, FSHRKO, LuRKO, *hpg* and Ovariectomised Female**
3 **Mice**

4
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1 **Abstract**

2 To investigate the relationship between gonadotroph function and ultrastructure we have
3 compared in parallel in female mice, the effects of several different mutations which perturb the
4 hypothalamic-pituitary-gonadal axis. Specifically, serum and pituitary gonadotrophin
5 concentrations, gonadotrophin gene expression, gonadotroph structure and number were
6 measured. FSH β knockout (FSH β KO), FSH receptor knockout (FSHRKO), LH receptor
7 knockout (LuRKO), hypogonadal (*hpg*), and ovariectomised mice were compared to control wild
8 type or heterozygote female mice. Serum levels of LH were elevated in FSH β KO and FSHRKO
9 compared to heterozygote females reflecting the likely decreased oestrogen production in KO
10 females as evidenced by the threadlike uteri and acyclicity. As expected there was no detectable
11 FSH in serum or pituitary and absence of expression of FSH β subunit gene in FSH β KO mice.
12 However, there was a significant increase in expression of the FSH β and LH β subunit genes in
13 FSHRKO female mice. The morphology of FSH β KO and FSHRKO gonadotrophs were not
14 significantly different to control except that secretory granules in FSHRKO gonadotrophs were
15 larger in diameter. In LuRKO and ovariectomised mice, stimulation of LH β and FSH β mRNA,
16 and serum protein concentrations were reflected in subcellular changes in gonadotroph
17 morphology, including more dilated rough endoplasmic reticula and fewer, larger secretory
18 granules. In the GnRH deficient *hpg* mouse gonadotrophin mRNA and protein levels were
19 significantly lower than in control mice and gonadotrophs were correspondingly smaller with less
20 abundant endoplasmic reticula and reduced numbers of secretory granules. In summary, major
21 differences in pituitary content and serum concentrations of the gonadotrophins LH and FSH
22 were found between control and mutant female mice. These changes were associated with
23 changes in expression of the gonadotrophin subunit genes and were reflected in the cellular
24 structure and secretory granule appearance within the gonadotroph cells.

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Introduction

The gonadotrophins, LH and FSH, have a key role in the differentiation, maturation and function of the mammalian reproductive system. The main functions of LH and FSH in female physiology are well known: FSH stimulates follicular maturation and granulosa cell oestrogen production whereas LH stimulates theca cell androgen production in the ovary (the substrate for granulosa cell oestrogen production), triggers ovulation and maintains the progesterone production of the corpus luteum. LH and FSH are synthesised within a single cell type in the anterior pituitary, the gonadotroph, and both consist of a common α subunit and a specific β subunit responsible for conferring biological activity on the heterodimer (1). Cyclical changes in gonadotrophin production and release drive the female reproductive cycle. The regulation of the pulsatile secretion of LH and of tonic secretion of FSH is complex and involves integration of GnRH, activin, inhibin, and ovarian-derived steroid signalling (2,3). Gonadotrophs can be identified by their distinctive secretory granules of variable size and electron-density (3). LH secretion occurs in discrete pulses in response to pulsatile GnRH whereas FSH secretion appears to be released independent of pulsatile GnRH (2). As both LH and FSH are produced and released from the same gonadotrophs via constitutive and regulated secretory pathways, differential packaging of LH and FSH with the granins, secretogonin II and chromogranin A respectively, in gonadotroph secretory granules has been proposed to underlie the different molecular pathways of secretion (4,5).

Mutations in gonadotrophin and gonadotrophin receptor genes severely affect fertility and mouse models exist for many human reproductive abnormalities, with genetic alterations in gonadotrophin secretion or action (6,7). Knowledge of reproductive mutant models has corroborated information from human mutations and provides useful tools for understanding

1 infertility and its treatment. In a recent study we compared the effect of mutations, both
2 naturally occurring and genetically engineered, which perturb the hypothalamic-pituitary-gonadal
3 axis in male mice (8). We now report the effects of the same mutations on reproductive function
4 in female mice. Specifically, we have compared FSH β knockout (FSH β KO), FSH receptor
5 knockout (FSHRKO), LH receptor knockout (LuRKO), hypogonadal (*hpg*) and ovariectomised
6 mice.

7
8 The FSH β KO mouse has a genetically engineered deletion of exons 1, 2 and 3 in the gene
9 encoding the FSH β subunit. As a result there is no production of biologically active FSH within
10 the pituitary, but in contrast to male mice carrying the same mutation, female mice are infertile
11 (9). Ovaries are significantly reduced in size compared to normal litter mates, and uteri are thin
12 and atrophic. Follicular development appears normal up to the pre-antral stage but mature
13 follicles and corpora lutea are not seen within the ovaries indicating that normal oestrous cycles
14 do not occur. Ovulation in FSH β KO mice can be induced by the administration of pregnant
15 mare's serum gonadotropin (PMSG) and human Chorionic Gonadotropin (hCG) (9).

16
17 The FSHRKO mouse carries a genetically engineered deletion in the gene encoding the FSH
18 receptor. In contrast to male mice the resulting inability to respond to circulating FSH results in
19 infertility in female mice (10). As in adult FSH β KO females, ovaries are significantly reduced in
20 size relative to normal mice, with follicular development arrested at the pre-antral stage (11).
21 The uterus is atrophic and the vagina imperforate and the mice do not undergo oestrous cycles.
22 In contrast to FSH β KO females there is no response to exogenous FSH (10).

23
24 The LuRKO mouse has a genetically engineered deletion of exon 11 of the LH receptor. As a
25 consequence mice are unable to respond to circulating LH and both males and females are

1 infertile (12). In adult females, follicular development progresses to the antral stage but no
2 mature follicles or corpora lutea are found and the mice are acyclic. Uteri are thin and atrophic
3 and vaginal opening is delayed (12).

4
5 The *hpg* mouse is a naturally occurring mutant with a deletion in the gene encoding the
6 hypothalamic gonadotrophin releasing hormone (GnRH) (13). Adult female mice homozygous
7 for the deletion present with under-development of the reproductive tract. The ovaries are very
8 small and follicular development does not proceed beyond the pre-antral stage. The uterus is
9 atrophic and the vagina imperforate. *Hpg* mice are deficient in gonadotrophic and ovarian steroid
10 hormones (14,15) and peptides, but ovarian and uterine development can be restored by the
11 administration of exogenous GnRH (16) or gonadotrophins (17).

12
13 The aim of this study was to investigate for the first time pituitary synthesis and regulation of LH
14 and FSH and to relate to gonadotroph ultrastructure in several mutant female mice in parallel in a
15 single study. (8). Throughout the paper we have referred to these mice in the order described
16 above, namely FSH mutants, LH receptor mutant, *hpg*, and ovariectomised mice as this reflects
17 the severity of the consequences of the mutations on female gonadal axis function.

18 19 **Materials and Methods**

20 **Mutant mice**

21 Breeding colonies of FSH receptor deficient, FSHRKO mice, FSH β deficient (10), FSH β KO
22 mice (9) and luteinising hormone receptor deficient, LuRKO mice (12) were established in our
23 laboratory. As both FSHRKO and FSH β KO females are infertile but males are fertile, breeding
24 pairs of KO males and heterozygous females were used to generate heterozygous and KO
25 offspring in a 1:1 ratio. Heterozygous females were used as controls for these two lines. The *hpg*

1 mutation was identified by PCR analysis of tail DNA as described previously (18). FSHR, FSH β
2 and LHR mutations were identified as described in Hirst et al (21). Hypogonadal (*hpg*) mice (4)
3 with a deletion in the gene encoding the GnRH gene (16) from the original colony discovered at
4 the MRC Laboratories, Harwell, Oxford (4) were bred within our department. The *hpg* mice were
5 on a C3H/HeH-101/H genetic background and the knockout mice on a mixed C57B16/129
6 background. All procedures were carried out in accordance with the Animals (Scientific
7 Procedures) Act 1986 and with the approval of a local ethical review committee.

8

9 **Serum and tissue collection**

10 All procedures were carried out under anaesthesia, (Rompun:Ketaset: 0.1 ml/Kg of a 20%:4%
11 [v/v] solution, Veterinary Supplies, University of Oxford) in 8-week old mice. For the analysis of
12 gonadotrophin hormones blood was collected from the jugular sinus, serum separated and frozen
13 at -20°C for assays. For the analysis of gonadotrophin subunit mRNA or gonadotrophin
14 hormone, pituitaries were dissected out, snap frozen in liquid nitrogen and stored at -70°C until
15 assayed. Ovaries and uteri were dissected out and weighed.

16

17 **Ovariectomy**

18 Animals were anaesthetised as described above. An incision of 0.5 cm was made in the dorsal
19 skin and abdominal wall of the right flank, and the ovary, oviduct and distal region of the uterine
20 horn were exposed. The ovarian artery was located and clamped, and the ovary was dissected
21 free from the surrounding fat and oviduct and removed. Abdominal wall and skin were sutured
22 and the procedure repeated on the left side. Mice were killed one month post ovariectomy and
23 comparisons were made with unoperated control mice.

24

25 **Hormone assays**

1 Serum and pituitary levels of FSH and LH were measured using in-house immunofluorimetric
2 assays (Delfia, Wallac OY, Turku, Finland) as described previously (8).

3
4 **RNA Extraction**
5 Total RNA was extracted from individual pituitaries with Trizol (Life technologies, Paisley, UK)
6 and residual genomic DNA was removed by DNase treatment (*DNA-free*, Ambion Inc supplied
7 by AMS Biotechnology, Abingdon, UK). DNase-treated RNA was quantified by
8 spectrophotometric measurement at λ 260nm. In total, 1 μ g of RNA was reverse transcribed using
9 random hexamers (Ambion) and Moloney murine leukaemia virus reverse transcriptase (Life
10 Technologies).

11
12 **Quantitative real-time PCR**
13 PCR experiments were carried out in a 25 μ l volume using a 96-well plate format. Primers and
14 probes were designed using Primer Express (Applied Biosystems, Warrington, UK) and probes
15 were synthesised with FAM 5' and TAMRA 3' (Taqman[®]). Primers were used at a final
16 concentration of 300 nM and probes at a concentration of 150 nM in ABI universal master mix
17 (Applied Biosystems, Warrington, UK). Primers and probes were selected from sequences
18 generated using Primer Express (Applied Biosystems, Warrington, UK). Primer and probe
19 sequences are listed in Table 1. To our knowledge alteration of gonadotrophin expression would
20 be unlikely to affect pituitary thyrotrophs and so TSH β subunit gene activity was measured as a
21 negative control for comparison with gonadotrope β subunit activity . Fluorescence was detected
22 on an ABI 7700 system (Applied Biosystems, Warrington, Cheshire). No reverse transcription
23 controls for each sample were screened to check for the presence of residual genomic DNA.

24

1 To measure cDNA levels, a threshold cycle (Ct) was selected within the exponential phase of the
2 amplification for all standards and samples. Arbitrary standards were generated by serial dilutions
3 of a cDNA pool from normal adult male pituitaries. A standard curve was generated by plotting
4 standards against Ct values, sample values were read from this standard curve and mRNA levels
5 were normalised relative to an endogenous control *mGapdh* mRNA (Applied Biosystems,
6 Warrington, UK), to allow comparison of different mRNAs between samples. A comparison
7 between *mGapdh* and *wbscr* (33) found that *mGapdh* showed a more stable pattern of expression
8 across pituitary glands from normal and mutant mice and was therefore selected as the
9 housekeeping gene of choice.

10

11

1 **Tissue processing and electron microscopy**

2 Pituitary glands for electron microscopy analysis were processed and analysed as described
3 previously (8). Briefly, the tissue was contrasted with uranyl acetate (2% (w/v) in distilled
4 water), dehydrated in ethanol and embedded in LR Gold resin. Ultrathin sections (50-80 nm)
5 were prepared using a Reichart-Jung ultracut microtome and mounted on nickel grids (Agar
6 Scientific, Stanstead, Essex, UK). For identification of bi-hormonal gonadotroph secretory
7 granules, sections were labelled for LH, and FSH as previously described (8) Specificity of
8 antibody labelling was confirmed in negative control sections in which the primary antibody was
9 replaced with non-immune serum. Finally, sections were counterstained with lead citrate and
10 uranyl acetate and examined on a JOEL 1010 transmission electron microscope (JOEL USA Inc.,
11 Peabody, MA, USA). For each pituitary (n=4), four randomly orientated sections, each
12 containing ten grid squares of intact tissue were counted for individual gonadotrophs identified
13 by immunogold labelling for LH and FSH. The total secretory cell population was determined by
14 counting the total number of nucleated pituitary (granulated and folliculo-stellate) cells per grid.
15 Gonadotroph number was expressed as a percentage of the total secretory cell population.
16 Immunogold identification of somatotrophs using a rabbit anti-mouse GH primary antibody
17 (1:4000; NHPP) was also carried in order to quantify the percentage of somatotrophs. This was
18 intended as a control for changes in the size of the gonadotroph population as changes in the
19 somatotroph population were not expected in the mutant mice investigated (19). Cell
20 morphology was analysed digitally using Axiovision, version 4.5 (Zeiss, Oberkochen, Germany)
21 (8). The following parameters were measured: cytoplasmic, nuclear and total cell areas; granule
22 area, granule density, granule diameter, percentage of secretory granules located in a 300-nm
23 depth of the plasma membrane and rough endoplasmic reticulum (rough ER). The methodology
24 is described in full in reference (8).

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Statistical analysis

Normal and heterozygous mice from the *hpg* and LuR colonies were initially analysed separately. Where there was no significant difference, data from normal mice and heterozygous mice were combined and expressed as control heterozygous/normal (H/N). Heterozygous females were used as normal controls in FSHR and FSH β colonies. Each mutant line was compared with its own control, comparisons were not made across different lines. Means were compared by one way analysis of variance (ANOVA). Where a significant overall difference was detected between groups, differences between individual means were assessed by the Bonferroni test and unpaired T-test. $P < 0.05$ was considered statistically different between mice and controls in the same line.

Results

Pituitary and serum gonadotrophin concentrations and pituitary gene expression

Figure 1 shows pituitary and serum gonadotrophin concentrations and figure 2 gonadotrophin subunit gene expression for each mutant and control. Table 2 provides a summary of changes measured for each mutant and control.

In FSH β KO adult females, pituitary and serum levels of FSH were at, or below the limit of detection of the FSH assay throughout. Furthermore, no expression of the FSH β gene was detected, confirming the absence of synthesis of FSH in this mutant. In control FSH β H female mice, serum FSH was 22% of the pituitary content of FSH. Serum LH levels were 0.3% of pituitary content in control FSH β H females and 0.7% in FSH β KO females. mRNA levels of both common α and TSH β subunit genes were not significantly different in FSH β KO female mice compared to control whereas LH β mRNA was significantly ($P < 0.01$) increased.

1 In FSHRKO females serum and pituitary concentrations of FSH were significantly higher
2 compared with FSHRH females and serum concentrations of LH were significantly higher
3 reflecting higher expression of FSH β and LH β subunit genes in the FSHRKO females relative to
4 heterozygote females. There was no significant difference in expression of common α or TSH β
5 subunit genes between KO and heterozygous females in FSHR females (Fig. 2).

6
7 In contrast, in LuRKO mice, serum concentrations of FSH were significantly ($P<0.01$) higher,
8 approximately doubled, compared with concentrations in control LuRH/N females (Fig. 1) but
9 pituitary content of FSH was not significantly different. A similar pattern was measured for LH;
10 in LuRKO females serum LH was significantly higher, being increased approximately 30-fold
11 ($P<0.01$) compared with control females, but there was no significant difference in LH pituitary
12 content. mRNA levels of common α ($P<0.01$), LH β ($P<0.01$) and FSH β ($P<0.05$) were
13 significantly higher in LuRKO females relative to LuRH/N females, whereas there was no
14 significant difference in amount of TSH β mRNA measured.

15
16 In *hpg* females the pituitary content of FSH and LH was significantly reduced ($P<0.01$) compared
17 to normal/heterozygous females. Serum levels of FSH and LH were also significantly lower than
18 in control mice, ($P<0.01$). mRNA levels of common α , LH β and FSH β genes were significantly
19 lower in *hpg* females compared to controls but the amount of TSH β mRNA was not significantly
20 different. As expected, in ovariectomised mice both pituitary and serum LH and FSH were
21 significantly increased ($P<0.01$) compared to intact controls. mRNA levels of common α , LH β ,
22 FSH β genes were all also significantly increased. The amount of TSH β mRNA was reduced in
23 ovariectomised mice but did not reach significance.

24

1 **Gonadotroph morphology**

2 Figure 3 shows representative electron micrographs of gonadotrophs in control and transgenic
3 mice and Table 3 shows the quantitative analysis of gonadotrophs and their organelles.
4 Gonadotrophs in rodents and sheep can be identified by their distinctive population of secretory
5 granules which have variable size and electron density (3). In the control mice secretory granules
6 were mainly moderately electron-dense of variable size (100-250nm). There was an absence of
7 the distinctive subpopulation of granules that display an electron-dense core and a relatively
8 electron-lucent 'halo'. Cell and cytoplasmic areas showed no significant difference between
9 controls (Fig. 3A) and FSH β KO (Fig. 3B), FSHRKO (Fig. 3C) and LuRKO (Fig. 3D) mice.
10 However, gonadotrophs in *hpg* female mice (Fig. 3E) were significantly ($P<0.01$) smaller than in
11 control mice. Gonadotrophs in ovariectomised mice (Fig. 3F) were similar to control
12 gonadotrophs in size. No significant difference was measured in nuclear area between control
13 and mutant mice, with the exception of significantly ($P<0.01$) smaller nuclei in *hpg* gonadotrophs
14 (Table 3).

15
16 Secretory granule numerical density represents the balance between gonadotrophin
17 synthesis ~~granule formation, storage~~ and release (20). Secretory granule density and distribution
18 were not significantly different in FSH β KO and FSHRKO gonadotrophs compared to control
19 (Table 3). However, granule density was significantly reduced compared to control in *hpg*
20 ($P<0.01$), LuRKO ($P<0.05$), and ovariectomised ($P<0.01$) gonadotrophs. Secretory granule
21 diameter was significantly increased in LuRKO (Fig. 3D; $P<0.01$), FSHRKO (Fig. 3C; $P<0.05$)
22 and ovariectomised (Figs. 3F, 4B; $P<0.01$) gonadotrophs compared to control but was
23 significantly reduced ($P<0.01$) in *hpg* (Fig. 3E) mice. Rough ER expansion reflects increased
24 secretory protein synthesis. Rough ER was significantly ($P<0.01$) reduced in the *hpg* (Fig. 3E)
25 but significantly increased in the LuRKO ($P<0.01$; Fig. 3D) and ovariectomised ($P<0.05$; Fig. 3F)

1 mice. The percentage of granules at the perimeter of the cell (which reflects the readily
2 releasable pool of granules) was significantly ($P<0.05$) increased in the LuRKO gonadotrophs but
3 was not significantly different in the other mutant and ovariectomised mice (Table 3). No
4 changes were seen in the thyrotroph population for any of the morphology parameters measured
5 (data not shown).

6
7 Figure 4 shows the distribution of immunogold labelling of LH and FSH on secretory granules in
8 control C57 mice (Fig. 4A) and ovariectomised mice (Fig. 4B). LH immunolabel detected with
9 15 nm immunogold particles and FSH immunolabel, detected with 5 nm immunogold particles
10 was distributed together in the majority of secretory granules and no particular distribution was
11 observed in relation to granules of a particular electron density.

12

13 **Gonadotroph number**

14 No significant difference in the proportion of LH/FSH-immunoreactive gonadotrophs (as a
15 percentage of total secretory cell number) was measured between control, FSH β KO, FSHRKO,
16 LuRKO and ovariectomised mice (Table 4). However, gonadotroph number was significantly
17 reduced ($P<0.05$) in *hpg* females compared to control.

18

19 **Ovary and uterine weights and histology**

20 Ovarian weights in adult FSH β KO, FSHRKO and LuRKO mice were significantly ($P<0.01$)
21 reduced to 30-50% of the corresponding heterozygote control females (Fig. 5A). In *hpg* mice
22 ovarian weights were 14% of control weights (Fig. 5). Uterine weights in all mutant mice were
23 significantly ($P<0.01$) reduced to approximately 7% of the weight of heterozygote controls (Fig.
24 5B). The vagina remained imperforate in all mutant mice at 8 weeks of age.

25

1 **Discussion**

2 This study has compared in female mice the effects of mutations which perturb the hypothalamic-
3 pituitary-gonadal axis in order to advance understanding of the structure and function of
4 gonadotrophs *in vivo* in conditions of altered GnRH, LH, FSH and ovarian steroid and peptide
5 signalling. Pituitary cells produce large amounts of protein hormones, packaged into dense-cored
6 secretory granules so that large amounts of hormone are rapidly available when required (21).
7 Granules must translocate and dock at the plasma membrane and undergo a series of priming
8 events for granule fusion to occur and granules adjacent to the plasma membrane are hence
9 described as the 'readily-releasable' pool (22). Real-time imaging studies in adrenal chromaffin
10 and β -cells have shown that there is preferential secretion of newly-formed secretory granules
11 that move within seconds-minutes to the plasma membrane, whereas older granules if not
12 secreted move away from the plasma membrane to within the cell (23). Therefore, secretory
13 granules located in proximity to the plasma membrane in gonadotrophs may similarly represent
14 release-competent younger granules. Secretory granule size in adrenal chromaffin cells has been
15 shown to correlate with the size of quanta of catecholamine released (24) so it is possible that
16 gonadotroph granule size similarly reflects the amount of gonadotrophin content. Although there
17 have been few dynamic studies of gonadotrophs to understand how granule volume and number
18 are regulated, the combined study of gonadotroph ultrastructure and hormone measurements
19 allows inferences to be made between secretory pathway function and structure.

20

21 At the ultrastructural level gonadotrophs are distinctive for the varied size and electron density of
22 the secretory granules contained compared to other pituitary secretory cell types. In rat and sheep
23 gonadotrophs the different secretory storage granules have been categorised into three
24 populations, distinguished by differential packaging with granin proteins, namely small, electron-
25 dense, LH and secretogranin II (SgII) positive granules; large, electron-lucent, FSH and

1 chromogranin A (CgA) positive granules and intermediate-sized granules containing an electron
2 dense LH and SgII positive core and an electron-lucent FSH and CgA-positive 'halo' outer
3 region (3,25). These granule populations are particularly distinctive in the male rat gonadotroph
4 but less so in the female rat (25). Consistent with these observations, in the present study,
5 gonadotrophs in female control mice showed a less extreme variation in secretory granule
6 appearance compared to male mice, granule diameter was overall smaller, and there was no
7 obvious compartmentalisation of FSH to larger, electron-lucent secretory granules (8).
8 Interestingly, granules were relatively more margined to the plasma membrane in control
9 female mice than in male perhaps due to differences in feedback control mechanisms. Although
10 changes in gonadotroph structure with oestrous cycle have been reported in the rat (26) we did
11 not detect any significant differences in gonadotroph morphology across the mouse estrous cycle
12 (data not shown).

13
14 As seen in male FSH β KO and FSHRKO mice, the female FSH mutants present with the same
15 phenotype, but in contrast to male mice, significantly higher concentrations of FSH (FSHRKO)
16 and LH (both mutants) were measured in both serum and pituitary, indicating a loss of feedback
17 regulation of these hormones at the level of the pituitary and hypothalamus. The high
18 concentrations of these gonadotrophins were associated with significant increases in mRNA
19 levels of the FSH β subunit gene in FSHRKO females and LH β subunit gene in both mutant
20 females. In ~~normal~~ control mice, cyclic changes in ovarian steroid production regulate
21 gonadotrophin secretion and release throughout the oestrus cycle (27,28). It is likely that the high
22 levels of FSH and LH measured are associated with the absence of biologically active oestrogen
23 in these mutant mice as evidenced by the atrophic uteri and imperforate vaginae. Consistent with
24 these observations female oestrogen receptor ER α KO mice are also infertile with increased
25 serum concentrations of gonadotrophins (29,30). The early stages of ovarian follicle

1 development do not require gonadotrophin input (31,32) but FSH is required to progress beyond
2 the pre-antral stage. In the absence of FSH stimulation, follicles do not acquire LH receptors and
3 are unable to convert thecal supplies of androgens to oestrogen (33,34). Ovarian concentrations
4 of inhibins A and B have been found to be significantly lower in FSH β KO and FSHRKO mice
5 compared to normal female mice (35) and in turn feedback regulation of FSH is impaired.

6

7 The ultrastructure of gonadotrophs in FSH β KO female mice however was not significantly
8 different in any of the parameters measured despite the loss of FSH stores, increased pituitary LH
9 content and increased LH serum concentrations. With respect to secretory granule morphology
10 increased LH content was not reflected in an increase in granule diameter but this may have been
11 counteracted by the absence of FSH stores. In male FSH β KO mice a similar profile of
12 morphology was evident except that secretory granule diameter was reduced (8) probably
13 reflecting the consequence of the loss of the significantly larger FSH stores in the male (8). In
14 female FSHRKO mice gonadotrophs the only difference measured was an increase in secretory
15 granule diameter compared to control gonadotrophs which may reflect the approximate doubling
16 of pituitary FSH content.

17

18 As observed in male LuRKO mice, serum levels of both FSH and LH were significantly
19 increased above control levels in LuRKO females but in contrast to male LuRKO mice, the
20 pituitary content of FSH and LH were maintained in the female despite the increased release of
21 both hormones from the pituitary (8). The increased synthesis and release of FSH and LH in
22 LuRKO females was associated with significantly higher mRNA levels for all three
23 gonadotrophin subunit genes. The increase in the production of the normally abundant common α
24 subunit, contrasted with the unchanged level of common α subunit in FSH β KO and FSHRKO
25 females. A further difference between FSH β KO, FSHRKO and LuRKO females relates to

1 inhibin production within the ovary. High levels of ovarian and serum inhibin B in the LuRKO
2 may account for the more moderate rise in FSH β mRNA levels and serum FSH levels compared
3 with the marked rise in serum LH in LuRKO females. Analysis of gonadotroph ultrastructure in
4 female LuRKO mice revealed fewer, but larger diameter secretory granules, possibly maintaining
5 pituitary gonadotrophin stores, which were marginalised to a greater degree towards the plasma
6 membrane compared to control. The greater number of secretory granules distributed towards
7 the plasma membrane and raised concentrations of serum LH and FSH are consistent with
8 increased granule release. In addition, rough ER was more dilated than control, likely indicating
9 increased protein synthesis as protein synthesis inhibitors have been shown to reverse dilated
10 rough ER in other systems (37).

11
12 In adult *hpg* female mice, despite the absence of steroid or peptide feedback to the pituitary there
13 was no increase in either FSH or LH relative to control females and subunit gene mRNA levels
14 remained at or below control levels, emphasising the primary role of hypothalamic GnRH in
15 facilitating gonadotrophin synthesis within the pituitary. In agreement with previous studies
16 (20,38) *hpg* gonadotrophs were smaller than control with less prominent rough ER, reduced
17 secretory granule area and smaller secretory granules. The reduced size of *hpg* gonadotrophs
18 correlated with the profound lack of synthesis and secretion of FSH and LH (although increased
19 gonadotrophin synthesis and secretion, as in the LuRKO female, did not correlate with an
20 increase in cell area).

21
22 In contrast, in ovariectomised female mice, where all ovarian regulatory factors have been
23 physiologically removed, a significant increase was measured in FSH β and LH β subunit gene
24 mRNA levels, in addition to a significant increase in serum and pituitary content of FSH and LH

1 in this study and as previously reported (39,40). At the subcellular level, many correlates of
2 hyper-stimulated gonadotrophin production and secretion were detected in ovariectomised mice.
3 Dilated rough ER correlated with the increased protein synthesis and glycosylation associated
4 with increased production of the gonadotrophins. The reduction in granule area per cytoplasmic
5 profile was consistent with increased release of granules. In ovariectomised female mice,
6 secretory granules increased in size and granules with an electron-dense core and electron-lucent
7 halo, frequently seen in male mice and rats were observed for the first time in this study. Thus,
8 a high pituitary FSH content appears to be associated with the presence of larger granules with
9 electron-lucent 'halos' (8). This suggests that ovariectomy in female mice induces sufficiently
10 high FSH production and storage for formation of granule halos during granule processing.
11 Interestingly the pituitary content of FSH in male mice one month post ovariectomy was reduced
12 to levels seen in hpg mice (8) suggesting that removal of testicular feedback to the pituitary
13 results in depletion of pituitary stores and release of all newly synthesised hormone indicating a
14 fundamental difference in gonadotrophin feedback control between male and female mice.

15
16 Since oestrogen has been shown to negatively regulate both GnRH at the hypothalamus and to
17 moderate release of LH and FSH from the pituitary, removal of oestrogen will facilitate both
18 increased release of gonadotrophins from the pituitary and allow increased stimulation of
19 gonadotrophin synthesis through increased GnRH activity. This difference in regulation between
20 male and female mice would allow for tight control of LH under normal conditions and for rapid
21 release and replacement of LH following an ovulatory surge in normal female mice. Since FSH
22 is under the joint control of inhibin and oestrogen produced from the follicular granulosa cells at
23 different stages of maturity, this may allow the different pattern of FSH increase seen throughout

1 the follicular phase of the cycle. However, in the absence of oestrogen regulation, the overriding
2 effect is GnRH stimulation of increased synthesis of FSH within the pituitary

3
4 In summary although the genetic mutations carried by the female mice in this study all result in
5 infertility, a different pattern of hormonal changes was found in each mutant which was reflected
6 in gonadotrophin subunit gene expression and gonadotroph ultrastructure. Consistent between
7 the female mutants are the observations that increased gonadotrophin synthesis and release
8 corresponded with increased rough ER density, a decrease in granule numerical density and an
9 increase in granule diameter corresponding to increased FSH production. The present study
10 looked at the single time point of 8 weeks of age. Further studies are investigating changes at
11 earlier developmental stages to determine when differences in gonadotroph structure and function
12 appear.

13

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17

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33

34 **Legends**

1 **Fig. 1.** Pituitary and serum luteinising hormone (LH) and follicle-stimulating hormone (FSH)
2 levels in adult FSH β heterozygote control (11), FSH β KO (10), FSHR heterozygote control (7),
3 FSHRKO (13), LuR heterozygote/normal control (25), LuRKO (10), *hpg* heterozygote/normal
4 control (18), *hpg* (13) and ovariectomised female mice (9). Number of mice in each group is
5 shown in brackets. Results are expressed as the mean \pm SEM. **P < 0.01 versus respective
6 strain control. Open columns, control mice; filled columns, mutant mice, knockout (KO) or *hpg*;
7 striped columns, ovariectomised (Ovx) mice.

8
9 **Fig. 2.** Gonadotroph common α , luteinising hormone (LH β), follicle-stimulating hormone
10 (FSH β) and thyroid-stimulating hormone (TSH β) subunit mRNA levels, normalised to *mGapdh*
11 mRNA in adult FSH β H, FSH β KO, FSHRH, FSHRKO, LuRH/N, LuRKO, *hpg*H/N, *hpg* and
12 ovariectomised female mice. Results are expressed as the mean \pm SEM. **P < 0.01 versus
13 respective strain control, n = 4 mice per group. Open columns, control mice; filled columns,
14 mutant mice, knockout (KO) or *hpg*; striped columns, ovariectomised (Ovx) mice.

15
16 **Fig. 3.** Representative electron micrographs of gonadotrophs in normal and female mutant mice:
17 (A) C57 Normal; (B) FSH β KO; (C) FSHRKO; (D) LuRKO; (E) *hpg*; (F) ovariectomised (Ovx).
18 Scale bar = 2 μ m. Representative organelles are labelled: m, mitochondria; rer, rough
19 endoplasmic reticulum; sg secretory granule; n, nucleus; g, Golgi apparatus. In Fig. 3D
20 arrowheads indicate granules at the periphery of the gonadotroph.

21
22 **Fig. 4.** Representative electron micrographs to show the morphological appearance of
23 gonadotroph secretory granules in female (A) C57 Normal; (B) ovariectomised (Ovx) mice.
24 Scale bar = 200 nm. Secretory granules analogous to the characteristic granules distinctive of
25 male gonadotrophs comprising a dense core and electron 'lucent' halo were observed in

1 ovariectomised gonadotrophs only. The cells were immunolabelled for luteinising hormone (LH)
2 detected with 15 nm immunogold particles (indicated with arrows) and follicle-stimulating
3 hormone (FSH) detected with 5 nm immunogold particles (indicated with filled arrowheads).

4
5 **Fig. 5.** Ovarian and uterine weights in adult FSH β H (18) FSH β KO (15), FSHRH (31), FSHRKO
6 (33), LuRH/N (15), LuRKO (6) and *hpg*H/N (13), *hpg* (9) and ovariectomised (5) female mice.
7 Open columns, control mice; H heterozygote, H/N heterozygote and normal, filled columns; KO,
8 knockout mice, *hpg*, naturally occurring mutant, striped columns; Ovx, ovariectomised one week
9 and 4 weeks. Results are expressed as mean \pm S.E.M, **P<0.01 versus respective strain control.
10 The number of mice in each group is shown in brackets.

11
12
13
14

1	Subunit gene	Primer	Sequence 5'-3'	ID Number
2	common α	Forward	CTGTTGCTTCTCCAGGGCATA	NM009889
3		Reverse	TTCTTTGGAACCAGCATTGTCTT	
4		Probe	CCCACTCCCGCCAGGTCCAA	
5	LH β	Forward	TGGCCGCAGAGAATGAGTTC	MM25145
6		Reverse	CTCGGACCATGCTAGGACAGTAG	
7		Probe	CCCAGTCTGCATCACCTTCACCACC	
8	FSH β	Forward	GGAGAGCAATCTGCTGCCATA	MM12932
9		Reverse	GCAGAAACGGCACTCTTCCT	
10		Probe	CTGTGAATTGACCAACATCACCATCTCAGTAGA	
11	TSH β	Forward	ACTTCATCTACAGAACGGTGGAAAT	MMTSHB1
12		Reverse	GCGACAGGGAAGGAGAAATAAG	
13		Probe	CCAGGATGCCCGCACCATGTTACT	
14				

15 **Table 1.** Sequences of mouse primer and probe sets for gonadotrophin subunit genes. Probes are
16 dual labelled FAM 5' and TAMRA 3' (Taqman[®]).

17

1 **Table 2.** Summary of the Changes in Pituitary and Serum Follicle-stimulating Hormone (FSH)
 2 and Luteinising Hormone (LH) concentration and mRNA levels in mouse mutant lines.

3

		FSHβKO	FSHRKO	LuRKO	<i>hpg</i>	Ovx
Pituitary	FSH	absent	↑	↔	↓	↑
Serum	FSH	absent	↑	↑	↓	↑
Pituitary mRNA	FSHβ	absent	↑	↔	↓	↑
Pituitary mRNA	common α	↔	↔	↑	↓	↑
Pituitary	LH	↑	↔	↔	↓	↑
Serum	LH	↑	↑	↑	↔	↑
Pituitary mRNA	LHβ	↑	↑	↑	↓	↑
Pituitary mRNA	TSHβ	↔	↔	↔	↔	↓

4

5

6

7 Direction of arrows indicate direction of change, horizontal arrows indicate no overall change.

8 LuR, LH receptor; KO knockout; Ovx, ovariectomised; *hpg* hypogonadal; TSH, thyroid-

9 stimulating hormone.

	Control	FSH β KO	FSHRKO	LuRKO	<i>hpg</i>	Ovariectomised
Cell Area (μm^2)	98 \pm 10	98 \pm 9	110 \pm 10	100 \pm 9	60 \pm 4 ^a	101 \pm 8
Cytoplasm Area (μm^2)	70 \pm 5	77 \pm 6	85 \pm 6	79 \pm 5	45 \pm 2 ^a	80 \pm 7
Nuclear Area (μm^2)	28 \pm 2	21 \pm 3	25 \pm 2	21 \pm 2	15 \pm 1 ^a	21 \pm 2
Granule Diameter (nm)	140 \pm 10	148 \pm 8	175 \pm 9 ^b	170 \pm 9 ^b	112 \pm 6 ^a	175 \pm 6 ^b
Granule Density (/ μm^2)	0.1 \pm 0.007	0.082 \pm 0.006	0.09 \pm 0.005	0.05 \pm 0.008 ^c	0.06 \pm 0.006 ^a	0.055 \pm 0.006 ^b
Rough ER (units)	2.1 \pm 0.1	2.0 \pm 0.2	2.2 \pm 0.2	3.5 \pm 0.3 ^b	1.1 \pm 0.1 ^a	3.0 \pm 0.2 ^c
% Granules at perimeter	51 \pm 5	49 \pm 5	46 \pm 8	65 \pm 5 ^c	45 \pm 3	52 \pm 5

Table 3: Subcellular morphology of gonadotrophs in female 8 week old female normal and mutant mice. ^a p<0.01 vs Normal C3H; ^bP<0.01, ^c P<0.05 vs control. Values expressed as mean \pm SEM (n=4 animals). Three groups of controls were independently assessed C57BL6/129 (transgenic mouse background), C3H (*hpg* background) and LuRH (transgenic heterozygote). No significant differences were found for each of the parameters measured and for simplicity the control data shown are from the C57BL6/129 mice.

Table 4 Percentage of Luteinising Hormone (LH) and Follicle-Stimulating Hormone (FSH)-positive cells (Gonadotrophs) and GH-positive cells (Somatotrophs) in Wild-Type and Mutant Female Mice as a Percentage of Total Anterior Pituitary Secretory Cell number.

	Control	FSH β KO	FSHRKO	LuRKO	<i>hpg</i>	Ovariectomised
% LH and FSH positive cells	14 \pm 2	16 \pm 3	18 \pm 2	18 \pm 2	9 \pm 1*	12 \pm 1
% GH positive cells	55 \pm 3	55 \pm 2	57 \pm 2	50 \pm 2	53 \pm 4	51 \pm 2

Values expressed as the mean \pm SEM (n=4 animals). *hpg*, hypogonadal; FSH, follicle-stimulating hormone; LuR, LH receptor; KO, knockout.