Increased basal and pulsatile secretion of FSH and LH in young men with 47,XXY or 46,XX karyotypes

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Abstract

Objective: The regulation of normal sexual maturation and reproductive function is dependent on a precise hormonal regulation at hypothalamic, pituitary, and gonadal levels. The aim of this study was to investigate the neuroendocrine integrity of the pituitary–gonadal axis in patients with primary testicular failure due to supernumerary X chromosomes.

Design: Cross-sectional study.

Methods: In this study, 7 untreated patients with primary gonadal insufficiency due to SRY-positive 46,XX (n = 4) and 46,XXY karyotypes (n = 3) aged 18.8 years and 25 age-matched healthy controls participated. Reproductive hormones, testicular size, and overnight LH and FSH serum profiles and overnight urine LH and FSH excretion were determined.

Results: Basal LH and FSH secretion was elevated 6.3- and 25.4-fold respectively in the patients and the amount of LH and FSH secreted per burst were 2.0- and 6.6-fold elevated. We found significantly more LH but not FSH peaks per 24 h, as estimated by the Weibull λ analysis. There was no difference between approximate entropy ratios or Weibull γ analyses indicating comparable orderliness and regularity of LH and FSH secretion. Overnight urinary LH and FSH excretion was significantly elevated in patients compared with controls and correlated significantly with calculated total overnight LH and FSH secretion respectively, thus validating deconvolution.

Conclusion: In this group of patients with severe hypergonadotropic hypogonadism due to a supernumerary X chromosome, higher basal, pulsatile, and total LH and FSH secretion were associated with significantly more LH peaks per 24 h in comparison with healthy controls. Thus, our data indicate that in patients with Klinefelter syndrome and XX male karyotypes the entire hypothalamic–pituitary–gonadal axis has undergone functional changes.

Introduction

The regulation of normal sexual maturation and reproductive function is dependent on a precise hormonal regulation at hypothalamic, pituitary, and gonadal levels. Testosterone (T) secretion by Leydig cells is driven by the pulsatile discharge of luteinizing hormone (LH) from the pituitary gland following episodic stimulation by hypothalamic gonadotropin-releasing hormone (GnRH). LH secretion is under tight negative feedback control by T that is primarily exerted at the hypothalamic level. Consequently, low serum T is accompanied by a compensatory increase in LH concentrations (1). Equally, the neuroendocrine regulation of pituitary follicle-stimulating hormone (FSH) release is under the influence of GnRH, T, estradiol (E2), and inhibin B concentrations, but other factors may influence this axis (2, 3).

Klinefelter syndrome (KS) is characterized by tall stature, gynecomastia, small testes, and androgen deficiency. Many of these characteristics can be attributed to the genetically determined primary gonadal defect characterized by the XXY karyotype. In adults with KS, the testes are small due to hyalinization of the seminiferous tubules, serum T levels are in the low normal range, and serum gonadotropin levels are subsequently elevated (4–8). The patients with SRY-positive 46,XX male karyotype resemble those with KS endocrinologically (9, 10).

The aim of this study was to evaluate the pituitary–gonadal axis in young and adolescent men with gonadal insufficiency due to supernumerary X-chromosome material (47,XXY and 46,XX males) in comparison with age-matched controls using the combined strategy of deconvolution analyses and measurement of serial LH and FSH concentrations.
Materials and methods

Subjects

A total of 32 subjects (25 healthy males aged 16.7–36.1 years (median 18.8 years) and 7 patients aged 16.0–39.1 years (median 21.0 years)) were studied. The controls were healthy community-dwelling men recruited from two different studies (11, 12). Twelve subjects (aged 22.5–36.1 years (median 31.2 years)) were part of a study on diurnal rhythm in serum inhibin B levels (11). They were all within 10% of ideal body weight, had no history of genital disorders, infertility or chronic diseases, and did not receive any medication. The remaining 13 participants (aged 16.7–18.8 years (median 17.1 years)) were recruited from a larger study on fetal growth performed on their mothers during 1985–1987 (13). Only young males with normal birth weights and normal fetal growth in the third trimester were included in the present study as described previously (12).

The patients consisted of three non-mosaic 47,XXY males and four SRY-positive 46,XX males.

The patients were referred to our clinic due to prenatal diagnosis, aberrant childhood/adolescent behavior or excessive growth, hypogonadism, or infertility. Chromosome analysis was performed on peripheral blood lymphocytes in each case. Karyotypes were established in 30 metaphases from each patient. All XX males were SRY positive (detected by PCR). The patients substituted with androgen (transdermal T, Testogel (Besins, Paris, France) (n = 4), peroral testosteroneundecanoat, Restandol (Organon, Oss, The Netherlands) (n = 2)) were withdrawn from treatment 14 days prior to and during the study.

None of the participants (patients and controls) had any acute or chronic illness, none received medications, or had any other verified endocrinopathy.

Study design

The participants were admitted to the department in the evening. An i.v. catheter was inserted into an antecubital vein and connected to a constant withdrawal pump (Swemed lab pump 3003; International AB, Frölunda, Sweden) by a heparinized catheter. From 2000 to 0800 h, blood samples (2 ml) were collected at 20-min intervals, and consequently 36 continuous blood samples were drawn from each participant. Out of the 25 controls, 12 were examined for 24 h with sampling every 30 min. Only the blood samples collected from 2000 to 0800 h were used for this study in these subjects.

All patients and 13 of the controls (with 20-min blood samples) collected urine overnight (12 h).

Testicular size and morphology were estimated by ultrasonography (14). Height was measured by a stadiometer (Holtain Ltd, Crymych, UK) to the nearest 0.1 cm and weight on a digital scale (SECA delta, model 707) with a precision of 0.1 kg. Body mass index (weight/height$^2$ (kg/m$^2$), BMI) was calculated.

Hormone analyses

Blood samples were clotted and centrifuged, and serum was stored at −20°C until hormone analyses were performed. Male reproductive hormones (T, E2, sex hormone-binding globulin (SHBG), and inhibin B) were determined in the morning blood sample at 0800 h. In all serum samples from each individual, concentrations of LH and FSH were measured by time-resolved immunofluorometric assay (DELFIA; Wallac Inc., Turku, Finland) with detection limits of 0.05 and 0.06 IU/l respectively. Intra- and interassay coefficients of variation (CV values) were both below 5% in the LH and FSH assays.

T, E2, and SHBG were measured by time-resolved fluoroimmunoassay (DELFIA, Wallac) with detection limits of 0.23 nmol/l. 18 pmol/l, and 0.23 nmol/l respectively. The intra- and inter-assay CV values were <6% for both T and E2 and <5.1% for SHBG. Inhibin B was determined using a specific two-sided enzyme immunometric assay from Oxford Bio-Innovation Ltd (Oxford, UK). The sensitivity of the inhibin B assay was 18 pg/ml, and the intra- and interassay CV values were <12 and 17% respectively.

Free T concentration was calculated by the equation of Vermeulen et al. using a fixed albumin concentration of 43 g/l (15).

Urinary LH and FSH

The volume of the 12 h urine collection was measured. To the urine sample (2 ml) from each individual urine collection was added 80 μl BSA (25% BSA stock) and 150 μl glycerol, and the urine samples were stored at −20°C until analysis. Urinary LH (u-LH) and FSH (u-FSH) concentrations were measured by immunofluorometric assay (DELFIA, Wallac), as described previously (16).

Method for analyzing pulsatile LH and FSH secretion

Hormonal secretion profiles were analyzed using deconvolution analysis in both the Klinefelter patients and the control subjects, in order to compare the endogenous LH and FSH secretion and its related secretion and elimination parameters in both study groups. Here we used a recently developed deconvolution method (17, 18) with fixed half-life for LH and fixed fast and slow half-lives for FSH. This fully automated Matlab and mathematically verified program uses the following steps/strategies in order to analyze LH and FSH secretion. First, the program detrends the data independent of the pulse frequency, after which the data are descaled. Thereafter, multiple potential pulse time sets are created by a smoothing process, based
on a mathematical model derived from the non-linear heat diffusion equation. Subsequently, the program calculates candidate pulse time sets. A rational model structure for the secretion and elimination process includes parameters for basal secretion ($\beta_0$), two half-lives ($\alpha_1$ and $\alpha_2$), basal and variable pulse mass ($\eta_0$ and $\eta_1$), variance of pulse mass ($\sigma_\lambda$), assay/measurement error ($\epsilon$), and three-component waveform parameters ($\beta_1$, $\beta_2$, and $\beta_3$). During the next step of the analysis, the program calculates all parameters simultaneously. Once these parameters are defined, the last step of the analysis is to compare the goodness of fit with that using other pulse time sets based on the Akaike’s information criterion (AIC) (19). The secretion and elimination parameters that the program finally calculates include basal secretion (in IU/l vdl per 12 h), pulsatile secretion (in IU/l vdl per 12 h), mass per pulse (in IU/l vdl), the time delay to maximal secretion from pulse onset (mode, in min), the regularity of the distribution of the pulses during the sample time period (Weibull $\gamma$) and the number of pulses per 24 h (Weibull $\lambda$). The $\lambda$ estimation of the Weibull distribution of interpulse intervals is more precise than the number of pulses observed per time interval sampled, since all interpulse intervals without bias introduced by edge artifacts of peaks potentially falling on the leading or trailing edge of the LH or FSH time series are used to calculate the mean 24 h LH pulse frequency as $\lambda$. Hence, the Weibull $\lambda$ estimation is used in the present results.

**Statistical analysis**

Clinical and biochemical characteristics of the whole group are provided as median and interquartile ranges. Differences in anthropometric measurements and hormone concentration between patients and controls were compared using the Mann–Whitney U test. Associations between urinary excretion and serum gonadotropin secretion were tested with Spearman’s rank correlation coefficient. All statistical analyses were carried out using the statistical package SPSS (version 13; SPSS Inc., Chicago, IL, USA).

**Ethics**

The study was approved by the local ethics committee (KF 01 270862, KF 01-243/97, KF 01-229/02, and KF 01-065/03) and conducted in accordance with the Second Helsinki Declaration. Signed informed consent was obtained from all subjects and also from the parents of the participants under 18 years of age.

**Results**

The overnight LH and FSH serum profiles in patients and controls are illustrated in Fig. 1A–D. The deconvolution analysis revealed a significantly higher basal, pulsatile, and total secretion of LH and FSH in patients and the amount of FSH and LH secreted per burst was significantly higher in patients (Table 1). We found significantly more LH peaks per 24 h in the patients, as estimated by the Weibull $\lambda$ analysis, but no difference in the number of FSH peaks. There was no difference between the Weibull $\gamma$ analyses when comparing patients and accordingly there was no difference between approximate entropy (ApEn) FSH and LH ratios (Table 1).

Our patients had significantly lower basal serum levels of total and free T than controls, whereas inhibin B was below the detection limit in all but one patient (who had a very low concentration (25 pg/ml)). We found no difference between E$_2$ and SHBG between groups (Table 1).

The patients had significantly smaller testicular volumes than controls, and significantly more patients had gynecomastia and a history of cryptorchidism than controls (Table 1).

Overnight urinary concentrations of LH and FSH were significantly higher in patients (Table 1; Fig. 2). There were no overlapping values between patients and controls for FSH. The total urinary excretion of LH and FSH correlated significantly with total overnight LH ($r=0.77$, $P<0.0001$) and FSH secretion ($r=0.67$, $P=0.006$) respectively (Fig. 2A and B).

**Discussion**

In this detailed study of FSH and LH pulsatility in Klinefelter patients, we found significantly higher basal, pulsatile, and total LH and FSH secretion. The number of LH but not FSH pulses in Klinefelter patients was significantly higher than that in the controls. The regularity of pulsing and the mode (shape) of secretion bursts were similar in the two groups.

Basal (non-pulsatile) and total LH secretion were elevated 6.3- and 3.8-fold respectively, whereas the amount of LH secreted per pulse was elevated 2.0-fold in the patients compared with controls. An amplification of the basal and total FSH secretion of 25.4- and 13.1-fold respectively was quantitated. The amount of FSH secreted per burst was 6.6-fold higher in the patients compared with controls. To our knowledge, the hypothalamic–pituitary–gonadal axis in patients with Klinefelter (47.XXY and SRY-positive 46.XX male) syndrome has not been studied in detail previously.

Our findings are in accordance with previously reported data on males with primary testicular deficiency (LH (20, 21) and FSH (3)). By contrast, another study found significantly more FSH peaks in males with primary testicular failure when compared with healthy controls (20). One reason for this discrepancy may be that the three previous studies on males with primary testicular failure included a total of nine patients with 47.XXY karyotype ($n = 1$ (21), $n = 5$ (20), and $n = 3$ (3)), whereas the remaining patients...
(n = 9) had primary testicular failure due to other reasons. Since these studies included patients with gonadal diseases other than KS, they may not be directly comparable with our study. KS is a complex entity in which numerous factors may influence the pituitary–gonadal axis.

To our knowledge, we have for the first time carried out the Weibull $\gamma$ and $\lambda$ analyses on patients with KS. $\gamma$ reflects the regularity of distribution of pulses during the sample period and $\lambda$ number of pulses per 24 h (calculated from the actual sample time period) respectively. We found a 1.5-fold increase in $\lambda$ for LH ($P = 0.02$) in the patients, with no difference in mean interpulse interval regularity ($\gamma$) or the time required to achieve maximum secretion (mode of 13 min in both cases).
ApEn analysis was carried out in order to compare the orderliness of secretion patterns of LH and FSH. Although the patients secreted significantly more LH and FSH per 12 h than controls, ApEn ratios of LH and FSH were not different between the two groups, indicating equal orderliness. This contrasts with previous findings in hypogonadal males in which a greater irregularity in FSH secretion was found (3). The reason for this could be the inclusion of males with primary gonadal deficiency due to other reasons than KS in the latter study as discussed above.

Our patients presented with the classical endocrine hallmarks of Klinefelter and 46,XX male syndromes, i.e., hypergonadotropic hypogonadism including highly elevated serum LH and FSH and low levels of serum total and free T. Additionally, inhibin B was undetectable in all but one patient (who had a very low concentration (25 pg/ml)), whereas E₂ and SHBG were within the normal ranges. Urinary FSH and LH levels were clearly elevated in the patients compared with controls.

Gonadotropin synthesis and secretion is dependent upon pulsatile stimulation by the episodic secretion of hypotalamic GnRH into the pituitary portal circulation. Alterations in pulse frequency and amplitude change the relative levels of gonadotropin synthesis and release through changes in gonadotroph gene expression in the anterior pituitary (22, 23).

The underlying mechanism of the GnRH pulse generator is complicated and appears to be regulated at many levels, and by a variety of neural and endocrine factors. The episodic neurosecretion by the GnRH neuron is partly generated by an inhibitory autocrine feedback action of GnRH on its own secretion but is also modulated by gonadal steroid hormones and peptide hormones (23–27). In males, T and E₂ exert independent effects on the regulation of LH secretion, whereas FSH secretion is under negative feedback control by inhibin B, and by E₂ resulting from aromatization of T.

We found no difference in the serum levels of E₂ or SHBG between patients and controls, but free T values were reduced in the patients. Therefore, the elevation in

### Table 1 Clinical characteristics of the patients with supernumerary X chromosomes and controls

<table>
<thead>
<tr>
<th>Clinical data</th>
<th>Patients (n=7)</th>
<th>Controls (n=25)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Height (cm)</strong></td>
<td>181.1 (162.4–186.2)</td>
<td>183.6 (177.0–186.0)</td>
<td>NS</td>
</tr>
<tr>
<td><strong>Weight (kg)</strong></td>
<td>71.2 (67.6–77.7)</td>
<td>75.0 (68.6–82.9)</td>
<td>NS</td>
</tr>
<tr>
<td>BMI (kg/m²)</td>
<td>22.4 (19.5–29.6)</td>
<td>21.3 (20.3–24.9)</td>
<td>NS</td>
</tr>
<tr>
<td><strong>Mean testicular volume by ultrasound (ml)</strong></td>
<td>0.93 (0.79–1.51)</td>
<td>10.75 (8.1–12.2)</td>
<td>0.001</td>
</tr>
<tr>
<td><strong>Gynecomastia (%)</strong></td>
<td>60</td>
<td>10</td>
<td></td>
</tr>
<tr>
<td><strong>Cryptorchidism (%)</strong></td>
<td>30</td>
<td>0</td>
<td></td>
</tr>
</tbody>
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**Basal reproductive hormones**

| Testosterone (nmol/l)                      | 8.0 (7.2–9.9) | 20.3 (17.1–25.5) | <0.001  |
| SHBG (nmol/l)                             | 41 (31–54) | 29 (20–42) | NS      |
| Estradiol (pmol/l)                        | 71 (61–73) | 91 (66–114) | NS      |
| Inhibin B (pg/ml)                         | Detection limit | 199 (140–237) | <0.001  |
| Free testosterone (pmol/l)                | 133 (122–205) | 461 (408–565) | <0.001  |

**Overnight gonadotropin profiles**

| Basal overnight LH secretion (IU/l per 12 h) | 161.4 (136.2–248.2) | 25.7 (14.8–49.7) | <0.001  |
| Total secretion LH (IU/l per 12 h)          | 225.3 (208.4–316.8) | 59.5 (35.5–83.3) | <0.001  |
| LH mass per pulse (IU/l)                    | 8.9 (4.4–11.6) | 4.5 (3.3–5.8) | 0.016   |
| LH pulsatile secretion (IU/l per 12 h)      | 68.6 (47.1–89.2) | 28.8 (17.6–36.6) | <0.001  |
| LH pulse frequency per 12 h                 | 10 (6–11) | 6 (5–7) | NS      |
| LH slow half-life (min)                     | 40.0 (40.0–77.4) | 71.5 (52.6–82.9) | NS      |
| Modeα                                    | 14.3 (13.2–16.3) | 14.6 (13.8–15.7) | NS      |
| Weibull γα                               | 4.3 (3.0–5.3) | 4.2 (3.3–9.5) | NS      |
| Weibull λα                               | 17.8 (13.7–19.8) | 11.6 (10.0–13.8) | 0.023   |
| ApEn LH ratio                            | 0.72 (0.70–0.92) | 0.71 (0.61–0.84) | NS      |
| Basal overnight FSH secretion (IU/l per 12 h) | 25.4 (4.0–46.5) | 1.0 (0.04–2.2) | 0.012   |
| Total secretion FSH (IU/l per 12 h)        | 50.9 (31.6–64.1) | 3.9 (2.7–5.0) | <0.001  |
| FSH mass per pulse (IU/l)                  | 3.3 (2.9–4.6) | 0.5 (0.3–0.7) | <0.001  |
| FSH pulsatile secretion (IU/l per 12 h)    | 28.5 (10.0–36.7) | 2.8 (1.2–4.2) | <0.001  |
| FSH pulse frequency per 12 h               | 7 (3–8) | 5 (3.5–6) | NS      |
| Modeα                                    | 8.3 (7.7–11.4) | 10.7 (8.0–14.0) | NS      |
| Weibull γα                               | 4.8 (2.7–5.7) | 3.2 (2.8–7.4) | NS      |
| Weibull λα                               | 13.5 (4.9–13.9) | 9.6 (7.9–11.1) | NS      |
| ApEn FSH ratio                           | 0.80 (0.68–0.86) | 0.85 (0.76–0.92) | NS      |

**Urinary gonadotropin secretion**

| Urinary LH (IU/l)                        | 20.1 (11.1–33.6) | 3.8 (2.8–5.7) | 0.017   |
| Urinary FSH (IU/l)                       | 32.2 (23.9–49.4) | 1.7 (1.5–2.1) | 0.001   |

Mann–Whitney U-test was used for the comparison of clinical data and basal hormones. Deconvolution analyses were compared using an unpaired t-test (on logarithmic data for LH). LH fast half-life and FSH fast and slow half-lives were fixed at 18, 110, and 620 min respectively.

*Comparison only between patients (n=7) and controls (n=13) with 20-min samples (see the Materials and methods section).
basal and pulsatile LH secretion could be attributable to the reduction in testosterone feedback. Likewise, the increase in basal FSH secretion might be due to the undetectable inhibin concentrations, possibly additionally associated with low free T feedback. The negative feedback from sex steroids and inhibin B exerts its effect on both hypothalamic and pituitary levels, and it may be speculated that an altered hypothalamic–pituitary–gonadal set point at this time may contribute to the testicular destruction resulting in a vicious circle.

Due to the differences in sampling interval, the deconvolution analyses of the patients were compared with a subgroup of controls (aged 17.1 (median) years) who were sampled in an identical way (i.e., 20-min intervals). The patients were, however, slightly but not significantly older (median 21.0 years) than controls. It is well established that the number of both the LH and FSH peaks increase with increasing age (38–40). However, when excluding the two oldest individuals (aged 34.4 and 39.1 years) in the Klinefelter group leaving the patients between 16.0 and 21.6 years of age and comparing these with the controls the number of peaks did not change for either FSH or LH.

In conclusion, we found significantly elevated basal, pulsatile, and total LH and FSH secretion when compared with healthy controls. The number of LH pulses in the Klinefelter patients was significantly higher than that in the controls. Our patients had hypergonadotropic hypogonadism due to the existence of supernumerary X chromosomes and the resulting primary testicular failure and none of the patients had any known pituitary defects. Thus, our data indicate that in the patients with KS and XX male karyotype, the entire hypothalamic–pituitary–gonadal axis has undergone functional changes resulting in an altered set point.

Acknowledgements

This study was in part supported by a research grant from the Kirsten & Freddy Johansen Foundation (A Juul), the Netherlands Organization for Scientific Research (NWO, The Hague, The Netherlands) (P Kok), and the National Institutes of Health Grant (AG023133, R21023777) and National Science Foundation Grant (DMS0107680) (J Veldhuis and D Keenan).
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Received 24 January 2008
Accepted 27 February 2008