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INCREASED CLEARANCE OF CORTISOL BY 5 β -REDUCTASE IN A SUBGROUP OF WOMEN WITH ADRENAL HYPERANDROGENISM IN POLYCYSTIC OVARY SYNDROME

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Abstract

Objective—Increased peripheral metabolism of cortisol may explain compensatory ACTH-dependent adrenal steroidogenesis and hence hyperandrogenism in polycystic ovary syndrome (PCOS). Previous studies have described an increased 5 α -reduction of cortisol or impaired regeneration of cortisol by 11 β -HSD1 in PCOS. However, these observations may be confounded by obesity. Moreover, the relationship between alterations in cortisol metabolism and the extent of adrenal androgen hyper-secretion in response to ACTH has not been established. This study aimed to examine the association between cortisol metabolism and ACTH-dependent adrenal hyperandrogenism in PCOS, independently of obesity.

Design—We compared 90 PCOS women (age 18–45 yr) stratified by adrenal androgen responses to ACTH₁₋₂₄ and 45 controls matched for age and body weight.

Methods—PCOS women were stratified as normal responders-NR, intermediate responders-IR, and high responders-HR to 250 μ g ACTH₁₋₂₄: NR (n= 27) had androstenedione and DHEA responses within 2 SD of the mean in controls; IR (n= 43) had DHEA responses >2 SD above controls; HR (n= 20) had both androstenedione and DHEA responses >2 SD above controls.

Results—All groups were similar for age, body weight, and body fat distribution. Basal testosterone, androstenedione, and 5 α -dihydrotestosterone plasma levels were similarly elevated among the three groups of PCOS compared with controls, whereas basal DHEA-S was higher in HR (2.8 \pm 1.2 μ g/mL) and IR (2.4 \pm 1.1 μ g/mL) than in NR (1.8 \pm 0.8 μ g/mL) and controls (1.7 \pm 0.6 μ g/mL). The HR group had the lowest basal plasma cortisol levels (101 \pm 36 ng/mL versus IR 135 \pm 42 ng/mL, NR 144 \pm 48 ng/mL, and controls 165 \pm 48 ng/mL; all P < 0.01), but the greatest

cortisol response to ACTH₁₋₂₄ (60-0) cortisol 173±60 ng/mL versus IR 136±51 ng/mL, NR 114±50 ng/mL, and controls 127±50 ng/mL; all $P < 0.01$), and the highest urinary excretion of total and 5 β -reduced cortisol metabolites (eg 5 β -tetrahydrocortisol/cortisol ratio 25.2±15.3 versus IR 18.8±10.7, NR 19.7±11.4, and controls 17.2±13.7; all $P < 0.05$). There were no differences in urinary excretion of 5 α -reduced cortisol metabolites or in 5 α -dihydrotestosterone/testosterone ratio between groups.

Conclusions—Adrenal androgen excess in PCOS is associated with increased inactivation of cortisol by 5 β -reductase that may lower cortisol blood levels and stimulate ACTH-dependent steroidogenesis.

Keywords

5 α -reductase; 5 β -reductase; adrenal hyperandrogenism; polycystic ovary syndrome; obesity

Introduction

The polycystic ovary syndrome (PCOS) affects between 5-7% of women. Although the ovary is the principal source of androgen excess in most of these patients, between 40-70% also demonstrate elevated circulating levels of adrenal androgens, particularly DHEA-S (1-3). The mechanism of adrenal androgen excess in these patients remains poorly understood. One hypothesis is that adrenal hyperandrogenism in PCOS arises from an exaggerated secretory response of the adrenal cortex to ACTH (4). This theory is supported by the close correlation between the responsiveness of adrenal androgens (i.e. androstenedione and dehydroepiandrosterone; DHEA) to ACTH and circulating DHEA-S levels (4). The mechanism leading to the increased responsiveness to ACTH for androstenedione and DHEA in PCOS remains to be determined. It has been postulated that this condition may be secondary to a compensatory overactivation of the hypothalamic-pituitary-adrenal (HPA) axis in response to increased metabolic clearance of cortisol. Similarly to what has been suggested in simple obesity (5), in PCOS increased cortisol metabolism might increase CRF and ACTH secretion by decreasing the HPA axis negative feedback signal, thus maintaining normal cortisol levels at the expense of increased ACTH-dependent steroidogenesis, and therefore of adrenal hyperandrogenism.

Most metabolism of cortisol in humans is by enzymes in the liver (A-ring reductases) and kidney (11 β -hydroxysteroid dehydrogenase type 2; 11 β -HSD2). The metabolic clearance rate of cortisol is also influenced by the extent of regeneration of cortisol from inactive cortisone, by 11 β -HSD1, mainly in the liver (6).

A-ring reductases catalyse 5 α - or 5 β -reduction of several steroids, converting cortisol to inactive metabolites and testosterone to 5 α - or 5 β -dihydrotestosterone (DHT). 5 β -DHT is inactive but its 5 α -isomer (5 α -DHT) is the most potent natural androgen. In humans, two isoenzymes of 5 α -reductase are encoded by separate genes. 5 α -Reductase type 1 enzyme is found in non-genital skin, liver, brain, muscle and to a lesser extent in stromal cells of adipose tissue. 5 α -Reductase type 2 is predominantly expressed in male accessory reproductive structures, such as the prostate, epididymis and seminal vesicles, and genital skin (7). 5 β -Reductase is expressed mainly in liver and at lower levels in testis and colon (8).

In cohorts of patients with PCOS, lower ratios of cortisol/cortisone metabolites (9) or higher 5 α -reduced cortisol metabolites (10-12) in urine have been reported, suggesting reduced 11 β -HSD1 activity (9), or increased 5 α -reductase activity (10-12), respectively. However, these results may be confounded by coexistent obesity, since obesity in men and women is associated with impaired liver 11 β -HSD1 activity (13, 14) and increased urinary excretion of A-ring reduced cortisol metabolites, particularly 5 α -reduced metabolites (15). Moreover, the relationship between alterations in cortisol metabolism and the extent of adrenal androgen hyper-secretion or increased responsiveness to ACTH has not been established.

The aim of the present study was to characterize whether exaggerated adrenal androgen responses to exogenous ACTH in PCOS are related to altered peripheral cortisol metabolism, independently of obesity. To achieve this we stratified PCOS patients according to androstenedione and DHEA response to ACTH.

Materials and Methods

Subjects

We investigated 90 unmedicated women with PCOS, aged 18-45 yr, and 45 controls recruited from the general population and comparable for age and weight. PCOS women had at least two of: (i) chronic oligoanovulation (luteal serum progesterone below 2 ng/mL (16)); (ii) hirsutism (Ferriman-Gallwey score \geq 8 (17) or elevated serum total and free testosterone levels (18)); (iii) polycystic ovarian morphology at ultrasound, according to the Rotterdam consensus conference criteria (19). Hyperprolactinemia, Cushing's syndrome, congenital adrenal hyperplasia and androgen secreting tumors were excluded by laboratory analysis (20). Controls had normal ovaries by ultrasound, no signs of hyperandrogenism and regular ovulatory menstrual cycles (progesterone levels \geq 8 ng/mL during the luteal phase (16)). Women were classified as normal-weight if body mass index values (BMI) was $<$ 18 kg/m² and $<$ 25 kg/m², overweight if BMI was \geq 25 kg/m² and $<$ 30 kg/m² and obese if BMI was \geq 30 kg/m² (21). Twenty-eight (31%) PCOS were normal-weight, 10 (11%) were overweight and 52 (58%) were obese. Within controls, 15 (33%) were normal-weight, 5 (11%) were overweight and 25 (56%) were obese. None of the subjects included in the study had thyroid dysfunction, abnormal prolactin levels, cardiovascular, renal or liver diseases on clinical examination and routine laboratory findings. The protocol was approved by the local ethics committee and written informed consent was obtained.

Assessment protocol

Anthropometric data (height, weight, and waist circumference), systolic and diastolic blood pressure, and fat mass and fat-free mass by bioimpedance (Akern-BIA, Pontassieve, Florence, Italy) were obtained from each subject, as previously described (22). The number of menses in the previous 6 months was recorded. Studies were performed between d 5 and 10 of the menstrual cycle, or randomly during severe oligomenorrhea or amenorrhea, after excluding pregnancy by appropriate testing. All subjects completed a 24-h urine collection and attended at 0800-0830h after overnight fast. Basal blood samples for hormonal (total testosterone, androstenedione, 17OH-progesterone, DHEA-S, 5 α -DHT, sex hormone binding globulin-SHBG, cortisol) and metabolic (glucose, insulin, total cholesterol, HDL-

cholesterol, triglycerides, ALT, AST) determinations were collected before a 75g oral glucose tolerance test (OGTT) with blood samples taken after 30, 60, 90, 120 and 180 min for glucose determination and after 60, 120 and 180 min for insulin determinations. On a second day, an ACTH₁₋₂₄ stimulation test (Synacthen, 250 µg i.v. at 0800-0830h) was performed, and samples for cortisol, DHEA, androstenedione, and 17OH-progesterone determinations were obtained at baseline and 60 minutes after stimulation. Samples were immediately chilled on ice and centrifuged; 24-h urine and serum were stored at -20 C and plasma at -80 C.

Biochemical assays in plasma

The assays employed for biochemical measurements have been reported elsewhere (23, 24) except 5α-DHT which was measured by an HPLC-RIA method (25).

The free androgen index (FAI) was calculated as the ratio between total testosterone and SHBG (26). Insulin resistance was estimated using the Quantitative Insulin-Sensitivity Check Index (QUICKI) and the Insulin Sensitivity Index during the OGTT (ISI) (27, 28).

Urine excretion of cortisol and its metabolites

5β-Tetrahydrocortisol (THF), 5α-THF, 5β-tetrahydrocortisone (THE), cortols, cortolones, and cortisone were measured in 24-h urine by electron impact gas chromatography-mass spectrometry (GC-MS) with minor modifications from a previously described method (29). Briefly, a urine aliquot was equilibrated with Internal Standards (11-epiTHF, 11α-hydrocortisone). The steroids were purified by Sep-Pak C₁₈ extraction, digested with β-glucuronidase, reextracted and converted to HDMS-TSIM derivatives. The samples were injected into a GC-MS (Agilent: GC 6890 – MS 5973) in Selected Ion Mode. Cortisol and its metabolites were quantified by the ratio of metabolite: Internal Standard area against standard curves for each steroid, included in every assay batch. Accuracy and precision were tested in every batch by measuring cortisol in a reference plasma sample (DG KG, Referenzinstitut für Bioanalytische Geschäftsstelle, Im Mühlenbach 52a D-53127 Bonn). Total cortisol excretion was calculated from the sum of 5β-THF, 5α-THF, 5β-THE, cortols and cortolones (30). Relative 5α- and 5β-reduction of cortisol was assessed by Ulick's A-ring reduction quotients, 5α-THF/cortisol, 5β-THF/cortisol, and 5β-THE/cortisone (31). The balance of 5β- and 5α-reductases was also assessed by the ratio 5β-THF/5α-THF (15). The balance between 11β-HSD1 and 11β-HSD2 activities in all tissues was assessed as the ratio of (5α-THF+5β-THF)/5β-THE (29, 32). Renal 11β-HSD2 activity was assessed as urinary cortisol/cortisone ratio (29).

Statistical analysis

Data are shown as means ± standard deviation (SD). Glucose and insulin response to the OGTT was expressed as area under the curve (AUC), which was calculated by the trapezoidal method. Hormone response to ACTH₁₋₂₄ stimulation test was calculated by the difference between 60min and basal concentrations (₍₆₀₋₀₎). Arbitrary cut-off values used to stratify responses of androstenedione and DHEA to ACTH₁₋₂₄ in PCOS women was calculated from the mean+2 SD of ₍₆₀₋₀₎ of androstenedione and DHEA in controls. The data were compared among the groups by analysis of variance (ANOVA). Simple

correlation analyses were used to relate adrenal androgens, cortisol metabolism and insulin sensitivity within PCOS. The impact of obesity (as BMI and % fat mass) on these correlations was tested by multiple correlation analyses.

Statistical analyses were performed on SPSS/PC+ version 8 (Chicago, IL, USA). Two-tailed *P* values < 0.05 were considered statistically significant.

Results

Stratification of PCOS women according to the response of adrenal androgens to ACTH

PCOS women were classified into three groups (normal responders-NR, intermediate responders-IR, high responders-HR) according to the response of androstenedione and DHEA to ACTH₁₋₂₄ compared with the group of controls. NR (n= 27) had androstenedione and DHEA responses to ACTH₁₋₂₄ within 2 SD of the mean in controls; IR (n= 43) had DHEA response >2 SD of controls but androstenedione response within 2 SD of controls; HR (n= 20) had both androstenedione and DHEA responses >2 SD of controls (Figure 1A, 1B). None of the PCOS women evaluated had an isolated androstenedione response >2 SD of controls.

The response of 17OH-progesterone to ACTH₁₋₂₄ was also higher in IR and HR compared with NR and controls (Figure 1C).

Anthropometric parameters (including prevalence of normal-weight, overweight and obesity) and systolic and diastolic blood pressure did not differ among NR, IR, HR and controls (Table 1). Hirsutism score was similarly higher, and number of menses in the previous 6 months similarly lower, in all the PCOS groups compared with controls (Table 1). There were no differences in plasma glucose or triglycerides, but the HR group had the highest total cholesterol and insulin_{AUC} during the OGTT, and the lowest insulin sensitivity scores (QUICKI and ISI) and HDL-cholesterol (Table 1).

Basal plasma levels of testosterone (total and free), androstenedione and 5 α -DHT were similarly higher, and SHBG similarly lower in all three groups of PCOS compared with controls (Table 1). However, basal 17OH-progesterone and 5 α -DHT/total testosterone ratio did not differ between any groups (Table 1).

Basal DHEA-S was higher in IR and HR groups compared with NR and controls (Figure 2A).

Cortisol and its metabolism

The HR group had the lowest fasting plasma cortisol levels (Figure 2B), but the greatest response of cortisol to ACTH₁₋₂₄ (Figure 1D). Both the HR and IR groups had higher total cortisol metabolite excretion (5 β -THF + 5 α -THF + 5 β -THE + cortols + cortolones) than controls (Table 2), which was attributable to increased excretion of 5 β -THF and 5 β -THE. Both the HR and IR groups had higher absolute excretion rates of 5 β -THF and 5 β -THE levels than controls, although only the HR group had 5 β -THF and 5 β -THE excretion higher

than the NR group. In contrast, 5 α -THF excretion did not differ between any groups (Table 2).

The A-ring reduction quotients reflecting 5 β -reductase activity (5 β -THF/cortisol and 5 β -THE/cortisone) were higher in HR compared with IR, NR and controls, but they did not differ between the IR and NR groups (Figure 3A, 3C). In contrast, the quotient reflecting 5 α -reductase activity (5 α -THF/cortisol) did not differ between groups (Figure 3B). Similarly, the 5 β -THF/5 α -THF ratio, that reflects the balance of 5 β - and 5 α -reductases, was higher in HR than in IR, NR and controls, but it was similar between the IR and NR groups (Figure 3D). The ratio of cortisol/cortisone, reflecting renal 11 β -HSD2 activity did not differ between groups (Table 2) but the (5 α -THF+5 β -THF)/5 β -THE ratio was similarly lower in all PCOS groups compared with controls, consistent with a lower 11 β -HSD1 activity in PCOS (Table 2).

Correlations between the response of adrenal androgens to ACTH, cortisol metabolism, DHEA-S and insulin sensitivity

Given the variability of hyperandrogenism even within the IR and HR groups (Figure 1), and the inevitably arbitrary cut-offs used to stratify these groups, simple correlation analyses were undertaken in all the PCOS women considered together. The response of androstenedione and DHEA to ACTH₁₋₂₄ were negatively ($P < 0.001$) correlated with fasting cortisol levels (Figure 4A, B) but positively correlated with cortisol response to ACTH₁₋₂₄ ($r = 0.33$, $P = 0.002$ and $r = 0.22$, $P = 0.042$, respectively). The androgen response to ACTH₁₋₂₄ was also positively correlated with DHEA-S fasting levels ($r = 0.22$, $P = 0.037$ for androstenedione; $r = 0.27$, $P = 0.016$ for DHEA), whereas it was not correlated with insulin_{AUC} during the OGTT ($r = 0.19$, $P = 0.113$ for androstenedione; $r = 0.02$, $P = 0.857$ for DHEA), nor with QUICKI ($r = -0.11$, $P = 0.312$ for androstenedione; $r = 0.03$, $P = 0.773$ for DHEA) or ISI ($r = -0.06$, $P = 0.604$ for androstenedione; $r = -0.03$, $P = 0.772$ for DHEA). Both androstenedione and DHEA responses to ACTH₁₋₂₄ were positively correlated with 5 β THE/cortisone ($r = 0.30$, $P = 0.003$ and $r = 0.24$, $P = 0.023$, respectively), whereas only the response of DHEA was significantly positively correlated with 5 β THF/5 α THF ($r = 0.24$, $P = 0.023$). The androgen response to ACTH₁₋₂₄ was not correlated with 5 β THF/cortisol ($r = 0.16$, $P = 0.138$ for androstenedione; $r = 0.20$, $P = 0.063$ for DHEA), nor with 5 α THF/cortisol ($r = 0.03$, $P = 0.819$ for androstenedione; $r = 0.05$, $P = 0.676$ for DHEA), or with the (5 α -THF + 5 β -THF)/5 β -THE ratio ($r = -0.16$, $P = 0.142$ for androstenedione; $r = -0.11$, $P = 0.327$ for DHEA).

Both 5 β THE/cortisone and 5 β THF/cortisol were negatively correlated with fasting cortisol levels ($r = -0.28$, $P = 0.035$ and $r = -0.29$, $P = 0.031$, respectively), but were not correlated with DHEA-S ($r = -0.01$, $P = 0.905$ and $r = -0.04$, $P = 0.745$, respectively). 5 β THE/cortisone, but not 5 β THF/cortisol, was significantly positively correlated with insulin_{AUC} during the OGTT ($r = 0.32$, $P = 0.007$), and negatively with QUICKI ($r = 0.31$, $P = 0.004$) and ISI ($r = 0.26$, $P = 0.015$). 5 β THF/5 α THF ratio also correlated positively with insulin_{AUC} ($r = 0.26$, $P = 0.032$).

These relationships were maintained when BMI or % fat mass were included in multiple regression analyses (data not shown).

Discussion

In this study we used the adrenal androgen (androstenedione and DHEA) responses to exogenous ACTH to stratify PCOS patients in order to examine predictors of adrenal androgen excess. In the 'high responder' group, where both androstenedione and DHEA responses were >2 SD of controls, we found abnormalities of adrenal steroid secretion and metabolism which were distinct both from healthy controls and from PCOS patients with normal responses to ACTH. Similar, but less striking, differences were observed in a group of 'intermediate responders', where DHEA response was >2 SD of controls, whereas androstenedione response was within 2 SD of controls. Specifically, the 'high responders' had evidence of chronic activation of the HPA axis, with an exaggerated cortisol and 17OH-progesterone response to ACTH. However, morning plasma cortisol values were decreased in this group. In addition, this group was characterized by an elevated 24h urinary cortisol metabolite excretion attributable to a selective increase in urinary excretion of 5β -reduced cortisol metabolites, which suggests enhanced 5β -reductase activity. Based on these findings we infer that the activation of the HPA axis in 'high responder' group may represent a compensatory response to enhanced peripheral metabolism of cortisol by 5β -reductase, particularly in the liver, thereby explaining higher adrenal androgen and cortisol responses to ACTH as well as lower than normal fasting cortisol concentrations.

We also found that 'high responder' as well as 'intermediate responder' PCOS groups had higher fasting DHEA-S levels than 'normal responder' and controls. Since increased levels of DHEA-S are commonly referred as an index of adrenal androgen overproduction in PCOS (33), we can infer that an exaggerated secretory response of adrenal androgens to ACTH may be at the basis of adrenal hyperandrogenism in PCOS. This theory, in agreement with previous reports (4), is supported by the close positive correlation between the responsiveness of androstenedione and DHEA to ACTH and circulating DHEA-S levels, observed within the PCOS cohort. However, we can also infer that only when there is an overall hyper-responsiveness of the adrenal cortex to ACTH (i.e. increased response of androstenedione, DHEA, 17OH-progesterone, and cortisol), as observed in the 'high responders', are circulating levels of DHEA-S increased; this is consistent with the adrenal hyperandrogenism, being sustained by an increased inactivation of cortisol by 5β -reductase. Low fasting cortisol levels may be considered an additional marker reflecting increased cortisol metabolism.

Surprisingly, at variance of previous studies, we did not find evidence of increased excretion of 5α -reduced cortisol metabolites in PCOS when compared with controls (10-12, 34). The finding of normal 5α -reductase activity in PCOS is further emphasized by the lack of any difference in 5α -DHT/total testosterone ratio in blood between PCOS and controls. The discrepancy with previous reports may be partially explained by the relatively low number of subjects enrolled in the other studies (10-12, 34), since there is considerable interindividual variation in urinary steroid excretion rates. Indeed, no significant difference in 5α -reductase activity between PCOS and controls was found in another study in which a larger population was screened (9). Another potential confounding factor, not fully explored in the other studies, is the impact of obesity, since we suspect that enhanced peripheral clearance of cortisol by 5α -reductase only occurs in PCOS as a manifestation of co-existent

obesity (13). In our study the prevalence of normal-weight, overweight and obesity was similar between PCOS and controls, so that any confounding effect of body weight on urinary metabolites reflecting 5 α -reductase activity was avoided. Additionally, no relationship of the secretory response of adrenal androgens to ACTH with 5 α -reduced cortisol metabolite excretion was observed within the PCOS group. This result, together with the finding of similar values of both 5 α -reduced cortisol metabolites and 5 α -DHT/total testosterone ratio among the three groups of PCOS, suggests that 5 α -reductase activity is not a determinant of the responsiveness of adrenal androgens to ACTH.

The urine (5 α -THF+5 β -THF)/5 β -THE ratio was decreased in all the PCOS groups compared with controls, which is consistent with some (9), but not all (10, 11), previous reports. Since the urinary free cortisol/cortisone ratio (an index of renal 11 β -HSD2 activity) was similar to controls, we can attribute the lower THFs/THE ratio of PCOS to an impaired 11 β -HSD1 activity, which appears to occur in PCOS irrespective of associated obesity. However, the lack of a significant relationships between adrenal hyperandrogenism and the urine (5 α -THF +5 β -THF)/5 β -THE ratio suggests that it is unlikely that variations in 11 β -HSD1 are an important determinant of ACTH-dependent adrenal hyperandrogenism in PCOS.

However, these data are probably not conclusive, since there is evidence that the urinary THFs/THE ratio is an inadequate indicator of 11 β -HSD1 activity in the presence of altered A-ring reductase activities. For example, one study in men showed that high liver fat content was associated with increased 5 β -reductase activity and with reduced urinary THFs/THE ratios, despite there being no change in conversion of cortisone to cortisol on first pass metabolism by 11 β -HSD1 (35). More specific measurement of regeneration of cortisol from cortisone will be required to understand the role of 11 β -HSD1, and its functional variability, in PCOS.

The mechanisms of up-regulation of 5 β -reductase in the 'high responder' PCOS group is unclear, since published information about the regulation of 5 β -reductase is rather sketchy. It may be that hyperinsulinaemia is involved since insulin sensitising agents ameliorate the increase in 5 β -reductase mRNA and activity which occurs in the liver of obese rats (36). Indeed, we found a positive correlation between 5 β -reductase activity and insulin levels in our cohort of women with PCOS, independently of body composition. Previous studies in rats described programmed regulation of 5 β -reductase activity by androgens, since testosterone treatment of female rats at birth increased the activity of 5 β -reductase after ovariectomy in adulthood (37). However, in our studies the elevation of testosterone was similar in the three PCOS groups. Finally, a selective increase in urinary excretion of 5 β -reduced cortisol and cortisone metabolites has been demonstrated previously in otherwise healthy men with high liver fat content (35). Unfortunately, we did not assess fat accumulation in the liver in the present study, but in PCOS women we did not find any correlation between serum transaminases (ALT and AST) and 5 β -reductase indices.

The consequences of increased 5 β -reductase activity may include not only increased peripheral metabolism of cortisol and compensatory activation of the HPA axis, but also metabolic disturbances, because 5 β -reductase is involved in cholesterol and bile acid metabolism (6). Accordingly, the 'high responder' PCOS women appear to have a more

severe dysmetabolic pattern, with lower plasma HDL-cholesterol levels and hyperinsulinaemia. It has been proposed that insulin may be responsible for both adrenal and ovarian hyperandrogenism in PCOS (38). During a euglycemic hyperinsulinemic clamp ACTH-stimulated steroidogenesis was potentiated in hyperandrogenic women (39), with a disproportionate rise in 17OH-pregnenolone, 17OH-progesterone and DHEA, rather than cortisol, progesterone and androstenedione, suggestive of inhibition of 17,20-lyase activity (39). Similar results were obtained in men (40). However, in our data exaggerated secretory response of androstenedione and DHEA to ACTH was not associated with a disproportionate 17OH-progesterone response to ACTH, suggesting that 17,20-lyase activity is not altered. Moreover, no association of androstenedione and DHEA response to ACTH and insulin levels was found in our cohort of women with PCOS, which does not suggest that insulin is a key driver for adrenal androgen production in PCOS. However, whether very high insulin concentrations, as are obtained during a clamp study, may dysregulate 17,20-lyase activity in PCOS, as previously suggested (39, 40) cannot be excluded by our data.

In conclusion, by stratifying patients according to adrenal androgen responses to exogenous ACTH, and matching subjects for body composition, we have clarified the alterations in cortisol metabolism which occur in PCOS. These data suggest a role for increased inactivation of cortisol by 5 β -reductase in the pathogenesis of adrenal hyperandrogenism in a subgroup of PCOS women, characterised by increased response of androstenedione, DHEA, 17OH-progesterone and cortisol to ACTH with low circulating basal cortisol levels. The ability to define subgroups in PCOS is important in improving therapy for this heterogeneous condition. Most variables are continuously distributed and any cut-off values are inevitably arbitrary and subject to further investigation, however our comparison between groups stratified by adrenal androgen responses to ACTH was supported by continuous correlation analyses. We can speculate that the PCOS group with adrenal hyperandrogenism sustained by an overall hyperresponsiveness of the adrenal cortex to ACTH stimulation may be the most susceptible to existing strategies which enhance negative feedback suppression of the HPA axis, for example with low dose glucocorticoid suppression, or to novel strategies which normalise peripheral cortisol metabolism.

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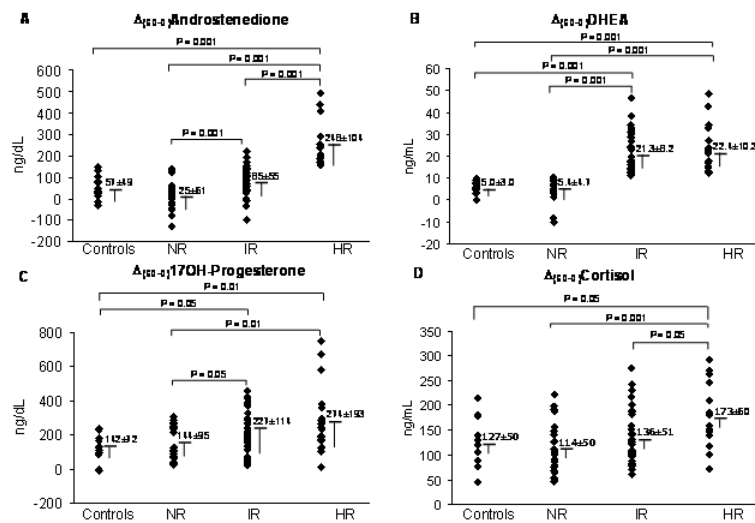


Figure 1.

Response of androgens and cortisol to ACTH₁₋₂₄ ($\alpha_{(60-0)}$) in PCOS normal responders-NR, intermediate responders-IR and high responders-HR, and in controls shown as individual values (scatter plot) and means \pm SD.

$P < 0.001$ for comparison in androstenedione and DHEA, $P = 0.025$ for comparison in 17OH-Progesterone, and $P = 0.001$ for comparison in cortisol among the three groups of PCOS (NR, IR, HR) by one-way ANOVA.

To convert to SI units, multiply DHEA by 3.467 (result in nmol/L) and cortisol by 27.59 (result in nmol/L).

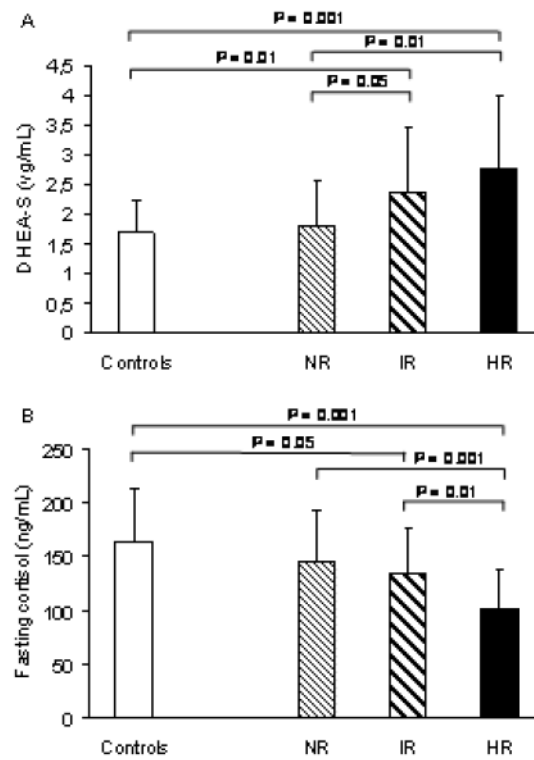


Figure 2.

Fasting plasma DHEA-S and cortisol levels in PCOS normal responders-NR, intermediate responders-IR and high responders-HR, and in controls. Data are means \pm SD.

$P = 0.004$ for comparison in DHEA-S and $P = 0.003$ for comparison in cortisol among the three groups of PCOS (NR, IR, HR) by one-way ANOVA.

To convert to SI units, multiply DHEA-S by 2.741 (result in $\mu\text{mol/L}$).

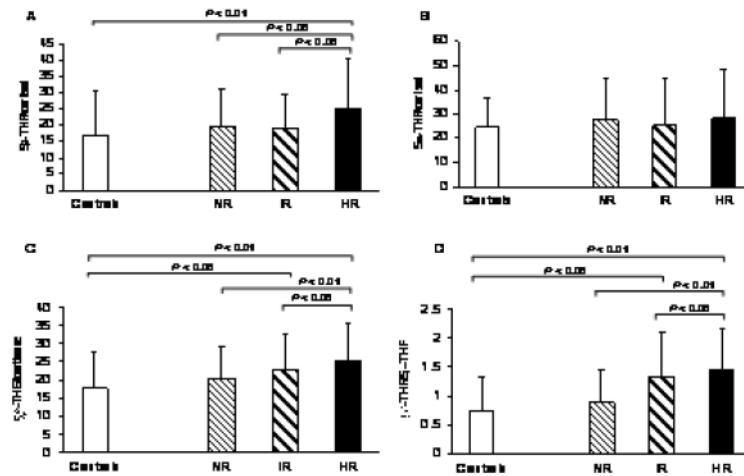


Figure 3.

Relative 5β- (5βTHF/cortisol, 5βTHE/cortisone) and 5α- (5αTHF/cortisol) reduction of cortisol and the balance of 5β- and 5α-reductases (5βTHF/5αTHF) in PCOS normal responders-NR, intermediate responders-IR and high responders-HR, and in controls. Data are means \pm SD.

$P = 0.049$, $P = 0.053$, $P = 0.825$, and $P = 0.044$ for comparison in 5βTHF/cortisol, 5βTHE/cortisone, 5αTHF/cortisol, and 5βTHF/5αTHF levels, respectively, among the three groups of PCOS (NR, IR, HR) by one-way ANOVA.

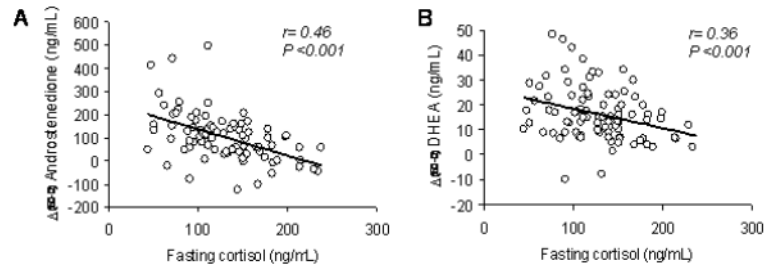


Figure 4. Relationship between androstenedione or DHEA response to ACTH₁₋₂₄ and fasting plasma cortisol levels.

Table 1

Clinical, hormonal and metabolic characteristics in women with PCOS, stratified according to the response of adrenal androgens to ACTH in normal responders-NR, intermediate responders-IR and high responders-HR, and in healthy controls

Variables	PCOS			P value ^a	Controls (n=45)
	NR (n=27)	IR (n=43)	HR (n=20)		
Age (yrs)	26.3±6.0	23.8±4.6	26.8±6.5	0.09	23.7±10.3
Body weight (kg)	84.7±25.4	78.1±18.1	87.2±20.8	0.231	79.5±22.6
Body mass index (kg/m ²)	31.7±8.7	30.0±7.0	32.6±7.6	0.447	30.0±8.3
25 kg/m ²	8 (31%)	14 (33%)	6 (30%)	0.436 ^b	
>25- 30 kg/m ²	2 (7%)	5 (11%)	3 (15%)		
>30 kg/m ²	17 (62%)	24 (56%)	11 (55%)		
Waist circumference (cm)	93.1±18.3	88.7±14.9	96.4±18.3	0.239	91.4±20.4
Fatty mass (%)	63.6±10.0	67.8±11.6	60.4±10.3	0.323	60.2±9.9
Free fatty mass (%)	39.9±11.1	32.2±11.6	36.8±10.3	0.308	39.8±9.9
Systolic blood pressure (mmHg)	123±17	128±11	123±14	0.216	128±14
Diastolic blood pressure (mmHg)	77±13	78±10	79±9	0.799	79±7
Hirsutism (Ferriman-Gallwey score)	10.0±6.9 ^{***}	12.1±6.9 ^{***}	11.4±7.7 ^{***}	0.576	2.2±1.0
Menses (n° of in the previous 6 months)	3.7±2.3 ^{***}	3.8±2.3 ^{***}	3.5±2.1 ^{***}	0.876	6.0±0.2
Fasting 0800-0830h plasma:					
Total testosterone (ng/mL)	0.66±0.26 ^{***}	0.67±0.18 ^{***}	0.70±0.26 ^{***}	0.818	0.48±0.16
FAI	2.44±1.09 ^{***}	2.99±1.56 ^{***}	2.95±1.50 ^{***}	0.282	1.51±0.87
Androstenedione (ng/dL)	382±163 ^{***}	373±138 ^{***}	401±119 ^{***}	0.799	201±57
17OH-progesterone (ng/dL)	157±77	156±107	158±138	0.480	127±85
DHT (ng/mL)	0.22±0.13 ^{***}	0.21±0.09 ^{***}	0.22±0.14 ^{***}	0.983	0.16±0.07
SHBG (nmol/L)	31.4±15.6 [*]	28.6±17.9 [*]	27.9±13.3 [*]	0.706	39.9±18.4
5α-DHT/Total testosterone	0.39±0.47	0.34±0.16	0.36±0.26	0.775	0.34±0.14
Oral glucose tolerance test:					
Glucose _{AUC} (mg/dL.min ⁻¹)	19,865±4,545	19,953±3,663	19,562±3,109	0.944	21,913±3,817
Insulin _{AUC} (μU/mL.min ⁻¹)	12,348±8,328	12,126±11,650	16,875±12,606 [*]	0.354	9,318±5,988
QUICKI	0.341±0.050	0.350±0.052	0.332±0.030 [*]	0.320	0.361±0.042
ISI	5.90±5.37	7.11±6.30	5.34±4.39 [*]	0.477	8.47±5.69
Total-cholesterol (mg/dL)	179.4±30.0	174.2±35.9	191.0±43.0 [*]	0.525	169.0±19.9
HDL-cholesterol (mg/dL)	55.3±16.6	53.7±14.3	46.4±13.6 ^{#, \$, *}	0.061	56.4±15.2
Triglycerides (mg/dL)	102.8±50.3	86.6±46.8	88.7±36.7	0.449	97.4±51.5
ALT (U/L)	20.7±3.8	22.1±7.8	24.0±8.4	0.292	19.1±5.2
AST (U/L)	22.4±8.2	25.6±17.1	30.8±23.3	0.243	21.2±10.5

Data are means ± SD.

^a Comparison among the three groups of PCOS (NR, IR, HR) by one-way ANOVA;

b comparison by two-tailed Fisher's exact test.

* $P < 0.05$,

*** $P < 0.001$ for post-hoc comparison between NR, IR or HR and controls.

$P < 0.05$ for post-hoc comparison between HR and NR.

§ $P < 0.05$ for post-hoc comparison between HR and IR.

FAI, free androgen index; DHT, dihydrotestosterone; SHBG, sex hormone binding globulin; AUC, area under the curve of the oral glucose tolerance test, calculated by the trapezoidal rule.

To convert to SI units, multiply total testosterone by 0.0347 (result in nmol/L), androstenedione by 0.0349 (result in nmol/L), 17-hydroxyprogesterone by 0.0303 (result in nmol/L), DHT by 3.467 (result in nmol/L), glucose by 0.0555 (result in mmol/L), insulin by 6.945 (result in pmol/L), Total- and HDL-cholesterol by 0.0259 (result in mmol/L) and triglycerides by 0.0113 (result in mmol/L)

Table 2

24-h Urine cortisol metabolites in women with PCOS, stratified according to the response of adrenal androgens to ACTH in normal responders-NR, intermediate responders-IR and high responders-HR, and in healthy controls

Variables	PCOS			P value ^a	Controls (n=45)
	NR (n=27)	IR (n=43)	HR (n=20)		
5 β -THF (μ g/d)	1592 \pm 904	1830 \pm 1026*	2013 \pm 803 #, *	0.046	1593 \pm 637
5 α -THF (μ g/d)	2536 \pm 1921	2971 \pm 2543	2653 \pm 2340	0.732	2344 \pm 1407
5 β -THE (μ g/d)	3292 \pm 1631	4136 \pm 3012*	4224 \pm 1703 #, *	0.050	3197 \pm 1783
Total (μ g/d) ^b	11329 \pm 5866	13459 \pm 8250*	13236 \pm 5708*	0.453	10418 \pm 6155
Cortisol/cortisone	0.57 \pm 0.22	0.65 \pm 0.29	0.53 \pm 0.21	0.138	0.57 \pm 0.26
(5 α -THF + 5 β -THF)/ 5 β -THE	1.24 \pm 0.46**	1.35 \pm 0.75*	1.06 \pm 0.36***	0.233	1.61 \pm 0.53

Data are means \pm SD.

^a Comparison among the three groups of PCOS (NR, IR, HR) by one-way ANOVA.

^b Total cortisol metabolites = 5 β -THF + 5 α -THF + 5 β -THE + cortols + cortolones.

Data are means \pm SD.

* $P < 0.05$,

** $P < 0.01$,

*** $P < 0.001$ for post-hoc comparison between NR, IR, or HR and controls.

$P < 0.05$ for post-hoc comparison between HR and NR.