

Effects of testosterone on PPAR γ and P450arom expression in polycystic ovary syndrome patients and related mechanisms

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Abstract. – **OBJECTIVE:** Polycystic ovary syndrome (PCOS) causes ovulation disorder and infertility in female patients. This study recruited PCOS patients and adopted testosterone therapy to analyze its effect on expression of nuclear peroxisome proliferation activated receptor γ (PPAR γ), and cytochrome P450 aromatase (P450arom) in ovary granular cells of patients.

PATIENTS AND METHODS: PCOS and non-PCOS patients were treated with testosterone at 1.0 nmol/L, 10 nmol/L. qRT-PCR was used to detect the expressions of PPAR γ and P450arom while Western blot was performed for the evaluation of protein expression.

RESULTS: In the experimental group, the level of PPAR γ mRNA was significantly elevated whilst the expression of P450arom was statistically decreased ($p < 0.05$). The treatment of 10 nmol/l testosterone significantly elevated PPAR γ mRNA and protein expression, and decreased P450arom expression ($p < 0.05$). The level of PPAR γ was positively correlated with testosterone concentration, and was negatively correlated with P450arom expression ($p < 0.05$).

CONCLUSIONS: Testosterone causes hyperandrogenism microenvironment of PCOS patients, with a correlated increase of PPAR γ and reduction of P450arom.

Key Words:

Testosterone, Polycystic ovary syndrome, PPAR γ mRNA, P450arom.

Introduction

Polycystic ovary syndrome (PCOS) manifests as ovulation disorder, and represents the major cause of women menopause or female infertility. At hormonal level, hyperandrogenism and hyperinsulinemia were developed in PCOS patients, with a rising level of follicle stimulating hormone (FSH) or luteinizing hormone (LH). Abundant

small ovary follicles without matured ones were found in PCOS patients due to the dysfunction of the hypothalamus-pituitary-ovary axis, and disorder of follicle selection or arresting. PCOS patients are more susceptible for metabolic disorders including type 2 diabetes and hypertension¹⁻³. It has been demonstrated that, during the pathogenesis of various diseases, hyperandrogenism primarily contributed to the disorder of glucose or lipid metabolism in PCOS women⁴.

Nuclear peroxisome proliferation activator receptor γ (PPAR γ) belongs to nuclear receptor superfamily, and regulates the occurrence and development of chronic metabolic disorders including diabetes, metabolic syndrome or hypertension^{5,6}. In ovary tissues, PPAR γ is mainly expressed in ovary granular cells at the development stage, and regulate estrogen secretion and ovary function under the direction of LH peak and low values⁷. Cytochrome P450 aromatase (P450arom) is highly expressed in granular cells, and can lead to hyperandrogenism once being suppressed for normal transformation⁸. In this study, the effect of testosterone on PPAR γ and P450arom in PCOS patients was determined.

Patients and Methods

Patients Information

A total of 30 PCOS patients (25-40 years old, average age = 29.2 ± 3.5 years) who were admitted to Jinan Maternity and Child Care Hospital from October 2015 to September 2016 were recruited. All patients were for the compliance with PCOS diagnostic criteria: infrequent menstruation; infertility; hairiness; obesity; LH in early phase ovary or after drug retraction/FSH > 3 and/or above normal testosterone level; enlargement of bilateral ovary; More than 10 large follicles

(> 10 mm diameter) in unilateral ovary. Another 30 non-PCOS patients (aging between 25 and 35 years, average age = 28.5 ± 3.1 years) who were characterized by normal menstrual cycle and ovulation were recruited. No significant difference existed regarding age, height or body weight between two groups, which were thus comparable ($p > 0.05$).

Inclusive criteria: No treatment of hormone drugs within 3 months; no malignant lesion occurred in ovary or uterus; no severe liver/renal disease.

Exclusive criteria: Complication of uterus body or cervical inflammation; other gynecology disease or endometrial disorder; long-term user of hormonal drugs; complication with uterus, ovary or oviduct.

The study protocol was approved by the Research Ethics Committee of Jinan Maternity and Child Care Hospital, and all patients signed the informed consent before the start of investigation.

Reagents

DMEM medium, streptomycin/penicillin and fetal bovine serum (Thermo Fisher Scientific, Waltham, MA, USA); q-PCR test kit, reverse transcription cDNA test kit (Invitrogen, Waltham, MA, USA); Biometra T1 PCR cycler Bio-Rad (Hercules, CA, USA).

Granular Cell Separation

Follicles were collected via vaginal route for extracting follicular fluid and were cultured *in vitro*. Follicular fluid was centrifuged and placed in PBS-Percoll mixture. Upper layer cells were removed in NH_4Cl for further centrifugation, and suspensions of ovary granular cells were filtered and obtained. Cells were re-suspended in DMEM containing 10% FBS and streptomycin/penicillin for 48 h culture. Cells were, then, fixed in paraformaldehyde and stored at -80°C .

Granular Cell Culture

Experimental group A and B were treated with 1 nmol/L and 10 nmol/L testosterone (N=10). Group C was maintained by using normal cell culture medium (N=10). Cells were repeatedly rinsed and mixed with TRIzol for -80°C storage.

qRT-PCR for PPAR γ and P450arom Expression in Ovary Granular Cells

According to the instruction of the PCR kit, PCR mixture was prepared including 0.1 ml cDNA, and 1 ml PCR buffer in 1 ml pre-mix. The reaction plate was sealed by the membrane, 20 μl reaction buffer was added for centrifugation. PCR conditions were: 95°C 5 min, followed by 40 cycles each of 95°C 5 s, and 60°C 30 s (Table I).

Western Blot for PPAR γ and P450arom Protein Expression in Ovary Granular Cells

Cells were rinsed and lysed for further mixture with 0°C RIPA lysis buffer. The mixture was centrifuged, and the supernatant was collected and stored at -80°C . Proteins were boiled at 100°C . Proteins were separated by electrophoresis and were transferred to the membrane, which was incubated with primary antibody (1:100 dilution) for 2 h. After rinsing, goat anti-rabbit secondary antibody was added for 1 h incubation. After development and exposure, the bands in the membrane were scanned and analyzed.

Data Processing

SPSS17.0 statistical software (SPSS Inc., Chicago, IL, USA) was used for data processing and data were presented as mean \pm standard deviation (SD). Enumeration data were presented by chi-square analysis, whilst measurement data were analyzed by Student *t*-test. The correlation between PPAR γ mRNA or P450arom and testosterone was analyzed by Spearman approach. A statistical significance was defined when $p < 0.05$.

Table I. Primer synthesis.

Gene		Temp ($^\circ\text{C}$)	Length (bp)
PPAR γ mRNA	5'-CAGGAGCAGAGCAAAGA-3'	60	274
	5'-GGAATCAGGGTGGTTCA-3'		
P450arom	5'-CAGCGGTCTCCCTTGATA-3'	60	314
	5'-TCTTCTCGGCATTTCTCC-3'		
GAPDH	5'-GCACCGTCAAGGCTGAGAAC-3'	60	580
	5'-TGGTGAAGACGCCAGGGA-3'		

Results

PPAR γ and P450arom Expression in Ovary Granular Cells Before Treatment

Expression of PPAR γ mRNA and P450arom in patients with ovary granular cells was measured. Compared to control group, the expression of PPAR γ was significantly increased in experimental groups, with statistical reduction of P450arom expression ($p < 0.05$, Figure 1).

PPAR γ and P450arom Gene Expression in Ovary Granular Cells After Treatment

We further measured the contents of PPAR γ mRNA and P450arom in PCOS patients after testosterone treatment. The elevation of PPAR γ and the decrease of P450arom levels were presented in a dose of testosterone-dependent manner ($p < 0.05$, Figure 2).

PPAR γ and P450arom Protein Expression in Ovary Granular Cells

We also tested the expressions of PPAR γ and P450arom in ovary granular cells after testosterone treatment. In the group of 10 nmol/l testosterone, PPAR γ protein expression was significantly up-regulated and P450arom level declined compared to the group of control or low dose of testosterone ($p < 0.05$, Table II, Figure 3).

Correlation Between PPAR γ and P450arom Expression in Ovary Granular Cells As Well As Testosterone

In ovary granular cells, PPAR γ mRNA expression was positively correlated with testosterone

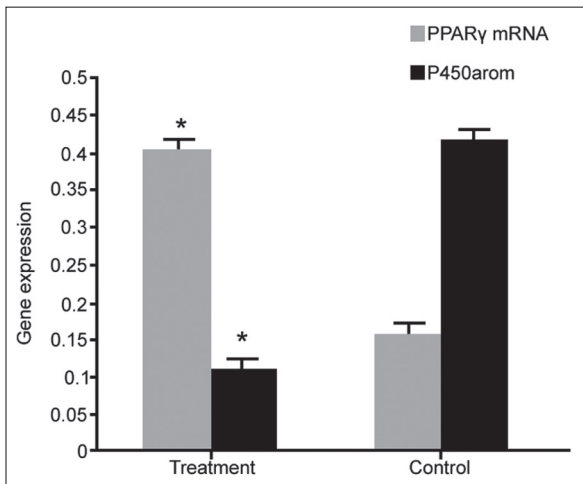


Figure 1. Expression of PPAR γ mRNA and P450arom in ovary granular cells. *, $p < 0.05$ compared to control group.

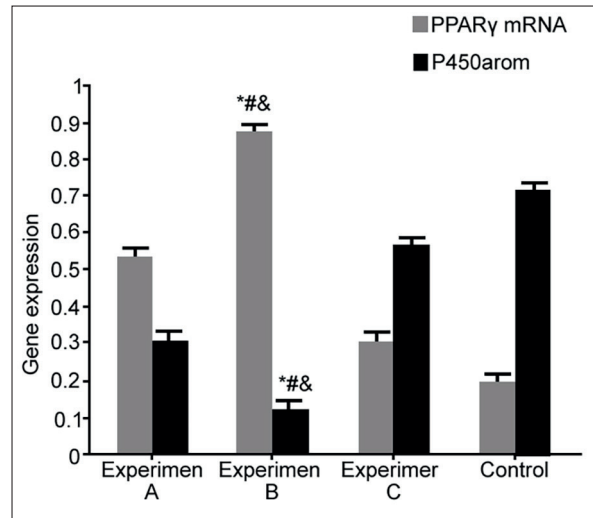


Figure 2. PPAR γ mRNA and P450arom gene expression in ovary granular cells after testosterone intervention. *, $p < 0.05$ compared to control group; #, $p < 0.05$ compared to experimental group A; &, $p < 0.05$ compared to experimental group C.

expression ($r = 0.640$, $p < 0.01$) and was negatively related to P450arom expression ($r = -0.738$, $p < 0.01$).

Discussion

PCOS represents a type of common endocrine disorder in clinics. The fertile period is largely determined by total depository of oocytes and their apoptotic rate⁹⁻¹¹. Hyperandrogenism belongs to a kind of unique endocrine manifestation of PCOS and severely affects normal ovulation, leading to infertility of PCOS women¹². P450arom activity was closely correlated with androgen/estrogen transition. As the enzyme activity was suppressed, androgen cannot be effectively

Table II. Protein expression of PPAR γ and P450arom in ovary granular cells after testosterone.

Group	N	PPAR γ	P450arom
Experiment	30		
A	10	1.932 \pm 0.004	1.376 \pm 0.003
B	10	2.549 \pm 0.008*#&	1.034 \pm 0.002*#&
C	10	1.378 \pm 0.004	1.668 \pm 0.004
Control	30	1.253 \pm 0.002	2.476 \pm 0.011

Note: *, $p < 0.05$ compared to control group; #, $p < 0.05$ compared to experimental group A; &, $p < 0.05$ compared to experimental group C.

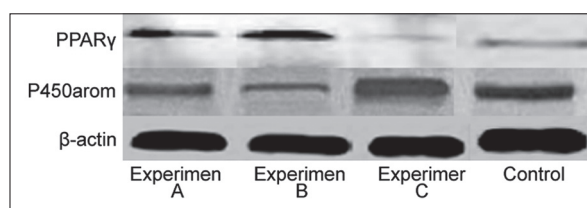


Figure 3. PPAR γ and P450arom protein expression in ovary granular cells after testosterone treatment.

transformed into estrogen, which resulted in focal hyperandrogenism^{9,10}. PPAR γ expression is positively correlated with follicular testosterone concentration and free androgen index of PCOS patients¹³. Liang et al¹⁴ generated PCOS mice and indicated that P450arom was the downstream factor of TGF- β /Smads signal transduction pathway. In human liver/renal tissues, agonist of PPAR can decrease Smad2 phosphorylation level and exert certain inhibitory effects on TGF- β /Smads signal transduction pathway. In ovary granular cells, however, the interaction between PPAR and P450arom is still unclear.

In this work, we found that PPAR γ mRNA was up-regulated and P450arom was down-regulated in PCOS patients and ovary granular cells. Consistently, a previous study¹⁵ showed the statistical correlation between P450arom gene polymorphism and PCOS pathogenesis. The hyperandrogenism occurred in PCOS women, mainly due to the alternation of P450arom substrate and inactivity. Furthermore, the up-regulation of PPAR γ was exhibited in ovary granular cells and PCOS patients, as in agreement with our study.

Faut et al¹⁶ investigated the correlation between antral follicle apoptosis in PCOS patients and PPAR γ expression, and validated that PPAR γ affected pregnant outcome via mediating follicular growth. Additionally, abnormal aromatization of P450arom affects follicular development. In this study, the concentrations of 1 nmol/l and 10 nmol/l testosterone were used for the evaluation of the effect of testosterone on PPAR γ and P450arom from ovary granular cells of recruited patients. Of note, high dose of 10 nmol/l testosterone remarkably increased the level of PPAR γ , with down-regulation of P450arom expression, compared with that of control or low dose of testosterone. The genomic study showed up-regulation of PPAR γ mRNA in ovary tissues of PCOS patients¹⁷. Evidence on PCOS mouse model showed that the hyperandrogenism microenvironment

caused by exogenous interference can change PPAR γ mRNA expression level in ovary granular cells of PCOS mice, leading to altered methylation level¹⁸, which was in line with our finding on positive correlation between testosterone and PPAR γ . Hyperandrogenism can induce phenotypic changes of PPAR γ in ovary granular cells to certain extents and is frequently accompanied by ovary dysfunction¹⁹. The clustered analysis also indicated the correlation between PPAR γ gene polymorphism and PCOS onset and occurrence, which was closely associated with hyperandrogenism²⁰. Mallqueo et al²¹ utilized P450arom inhibitor as a continuously exogenous stimulus to study its role on PCOS-induced hyperandrogenism model of mice. Assays showed that once P450arom gene activity was suppressed, it can directly participate in the onset and progression of PCOS-related hyperandrogenism, which combined with our data on negative correlation between P450arom and hyperandrogenism, suggesting the inhibitory effect of P450arom on hyperandrogenism²¹, which confirmed that the decreased activity of P450arom was associated with the androgen-estrogen transition. As multiple factors were identified in the regulation of PCOS²², and in the future, precise assay for the study of PPAR γ and P450arom, combined with treatment such as metformin, may provide novel idea and strategy for the therapy of PCOS.

Conclusions

Our data described the behavior of PPAR mRNA and of cytochrome P450arom before and after different doses of testosterone in ovary granular cells and PCOS women. Testosterone significantly elevated the PPAR γ expression while down-regulated the level of P450arom in both ovary granular cells and PCOS patients. PPAR γ was positively correlated with testosterone but negatively related to P450arom expression.

Conflict of Interest

The Authors declare that they have no conflict of interests.

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