SIRT3 protects endometrial receptivity in patients with polycystic ovary syndrome

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Abstract

Background: The sirtuin family is well recognized for its crucial involvement in various cellular processes. Nevertheless, studies on its role in the human endometrium are limited. This study aimed to explore the expression and localization of the sirtuin family in the human endometrium, focusing on sirtuin 3 (SIRT3) and its potential role in the oxidative imbalance of the endometrium in polycystic ovary syndrome (PCOS).

Methods: Endometrial specimens were collected from both patients with PCOS and controls undergoing hysteroscopy at the Center for Reproductive Medicine, Peking University Third Hospital, from July to August 2015 and used for cell culture. The protective effects of SIRT3 were investigated, and the mechanism of SIRT3 in improving endometrial receptivity of patients with PCOS was determined using various techniques, including cellular bioenergetic analysis, small interfering ribonucleic acid (siRNA) silencing, real-time quantitative polymerase chain reaction, Western blot, immunofluorescence, immunohistochemistry, and flow cytometry analysis.

Results: The sirtuin family was widely expressed in the human endometrium, with SIRT3 showing a significant increase in expression in patients with PCOS compared with controls (P < 0.05), as confirmed by protein and gene assays. Concurrently, endometrial antioxidant levels were elevated, while mitochondrial respiratory capacity was reduced, in patients with PCOS (P < 0.05). An endometrial oxidative stress (OS) model revealed that the downregulation of *SIRT3* impaired the growth and proliferation status of endometrial cells and reduced their receptivity to day 4 mouse embryos. The results suggested that SIRT3 might be crucial in maintaining normal cellular state by regulating antioxidants, cell proliferation, and apoptosis, thereby contributing to enhanced endometrial receptivity.

Conclusions: Our findings proposed a significant role of SIRT3 in improving endometrial receptivity in patients with PCOS by alleviating OS and regulating the balance between cell proliferation and apoptosis. Therefore, SIRT3 could be a promising target for predicting and improving endometrial receptivity in this patient population.

Keywords: Apoptosis; Endometrial receptivity; Oxidative stress; Polycystic ovary syndrome; Sirtuin 3

Introduction

Infertility is defined as the inability to achieve a clinical pregnancy after regular unprotected sexual intercourse for more than 1 year, affecting approximately 10% of the population. This complex condition arises from various etiological factors, including female, male, and unexplained factors. Ovulation disorders, accounting for approximately 25–35% of infertility cases, are particularly significant. Notably, polycystic ovary syndrome (PCOS) stands out as the most prevalent ovulation

disorder. Patients with PCOS often seek medical assistance and undergo assisted reproductive technology (ART) to achieve pregnancy. [2] Successful pregnancy outcomes

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Chinese Medical Journal 2025;138(10)

Received: 05-12-2023; Online: 09-05-2024 Edited by: Yanjie Yin



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depend largely on embryo quality and endometrial environment.^[3] The use of oocytes donated by the patients with PCOS did not decrease the overall pregnancy success rate.^[4] This finding suggested that the reduced fertility in patients with PCOS was associated with not only factors such as internal physiological disorders and diminished oocyte quality but also compromised endometrial receptivity.^[5,6] Previous studies focused on the connection between compromised endometrial receptivity in patients with PCOS and disrupted regulatory mechanisms, including imbalances in endometrial oxidative stress (OS), chronic low-grade inflammation, metabolic disorders, endocrine dysregulation, and abnormalities in endometrial hormone receptors. Among these factors, OS imbalance plays a prominent role.^[7]

OS is characterized by the overproduction of reactive oxygen species (ROS), wherein the oxidative burden surpasses the capacity of antioxidant defense systems to neutralize these radicals. This imbalance leads to the chronic accumulation of ROS in various cell types, including vascular endothelial cells, endometrial epithelial cells, and stromal cells. Consequently, this aberrant redox state triggers internal signaling pathways that directly or indirectly induce cellular and tissue damage, affecting components such as deoxyribonucleic acid (DNA), lipid membranes, and protein. [8-10] Within the context of OS, mitochondrial respiration plays a crucial role in ROS generation. Disruptions in mitochondrial function can have serious consequences, such as a significant reduction in the activities of cellular respiratory chain complexes I and IV, impairments in oxidative phosphorylation, reduced adenosine triphosphate (ATP) synthesis, changes in cell membrane permeability, and an abnormal increase in cellular OS levels.^[7] Increasing evidence supports significantly elevated OS levels in patients with PCOS compared with normal individuals, as evaluated by circulating markers including malondialdehyde, catalase (CAT), superoxide dismutase (SOD), and glutathione peroxidase (GPx). [6,11,12] Previous studies identified increased concentrations of OS markers in the follicular fluid from patients with PCOS compared with controls, potentially leading to a decline in oocyte quality and affecting subsequent steps of the reproductive process. [13,14] However, studies on the correlation between endometrial OS and PCOS remain inadequate.

Sirtuins are a family of nicotinamide adenine dinucleotide (NAD+)-dependent protein deacetylases playing a vital role in oxidative metabolism and lifespan extension in lower organisms. [15-19] In mammals, this family consists of seven members (SIRT1-7), each exhibiting distinct subcellular localization. Specifically, SIRT1, SIRT6, and SIRT7 are predominantly localized in the nucleus; SIRT2 in the cytoplasm; and SIRT3, SIRT4, and SIRT5 in the mitochondria. [20,21] These versatile enzymes participate in various biological processes. SIRT1 primarily participates in energy metabolism, gene transcription, and OS; SIRT2 in cell cycle regulation; SIRT3 in energy metabolism and apoptosis; SIRT4 in the tricarboxylic acid cycle and insulin secretion; SIRT5 in the ornithine cycle; SIRT6 in DNA repair and inflammation; and SIRT7 in nucleolus stress and rRNA transcription. [22] Although existing studies

investigating the role of sirtuins in reproductive functions have predominantly focused on SIRT1 and SIRT3 in ovarian function, energy metabolism, and OS,^[23] studies exploring the role of sirtuins in the endometrium are limited. A previous study showed that resveratrol, a SIRT1 activator, played a crucial role in endometrial receptivity by upregulating E-cadherin expression.^[24] Additionally, several studies have suggested distinctive roles for sirtuins in endometrial cancer, with SIRT1 shown to accelerate the proliferation of endometrial carcinoma cell lines.^[25–28] In the context of PCOS, numerous studies have reported the presence of OS in affected patients.^[29] However, the relationship between the imbalanced oxidative endometrium of PCOS and sirtuins remains poorly understood.

This study aimed to investigate the expression and localization of sirtuins in the human endometrium, focusing on SIRT3 and its potential role in the imbalanced oxidative endometrium of PCOS.

Methods

Study participants and tissue collection

In this study, 71 participants were enrolled from the Reproductive Medicine Center of Peking University Third Hospital, including 34 patients with PCOS and 37 controls. All specimens were acquired with the informed consent of patients after approval from the Ethics Committee of Peking University Third Hospital (No. S2018042). The diagnostic criteria for PCOS were based on the presence of at least two of the following three features: (1) oligomenorrhea or amenorrhea, (2) clinical and/or laboratory-confirmed hyperandrogenism, and (3) polycystic ovaries observed via ultrasonography, with the exclusion of related disorders.^[30] The control group consisted of women experiencing infertility due to male factors, excluding those with endometriosis, tubal effusion, endometrial cavity fluid, endometrial polyps, submucosal uterine fibroids, and pelvic tuberculosis. Eligible participants were 35 years of age or younger and had not received any hormonal treatment in the last 3 months. The endometrial samples were obtained during hysteroscopic examination in the hyperplasia phase, as confirmed by histopathological analysis. Fresh specimens were collected, preserved in phosphate-buffered saline (PBS; Gibco, NY, USA) on ice, washed three times with PBS, and subsequently stored in liquid nitrogen for longterm preservation.

Cell culture and induction of OS

The endometrial tissue samples obtained from the hysteroscopic examination were dissociated into single cells using mechanical and enzymatic digestion as described previously. The endometrial cells obtained from both the PCOS and control groups were cultured in Dulbecco's Modified Eagle Medium (DMEM)/F12 medium (Gibco) containing 10% fetal bovine serum (Hyclone, Loga, USA), penicillin (100 U/mL, Gibco), and streptomycin (100 µg/mL, Gibco). The cells were subjected to 250 µmol/L hydrogen peroxide (H₂O₂) (FreeMoreBio, Beijing, China) treatment

for 12 h to induce OS, and then cultured in a CO₂ incubator (Thermo Fisher Scientific, Waltham, MA, USA) under a 5% CO₂ atmosphere at 37°C. The following experiments and measurements were performed in parallel for both the control and OS groups.

Cellular bioenergetic analysis

The Seahorse Bioscience XF analyzer (Billerica, MA, USA) was employed to measure the real-time energy metabolism of the cells to assess mitochondrial function. Endometrial cells derived from groups with either OS or PCOS, as well as the control group, were cultured in the same energy metabolism orifice plate, with a cell density of $4-6 \times 10^4$ (24-well plate, 250 µL per well). Subsequently, the cells in the orifice plate were washed three times with the cell culture medium and subsequently treated with oligomycin (1 µg/mL; Sigma-Aldrich, St. Louis, MO, USA), carbonyl cyanide 4-(trifluoromethoxy) phenylhydrazone (FCCP) (0.7 µmol/L; Sigma-Aldrich), antimycin A (1 µmol/L; Sigma-Aldrich), and rotenone (1 µmol/L; Sigma-Aldrich) in the dosing wells surrounding each cell hole on the second day. The oxygen consumption rate (OCR) of the cells was measured using the Seahorse Bioscience XF analyzer following incubation under a 5% CO₂ atmosphere at 37°C for 20 min.

Small interfering ribonucleic acid (siRNA) silenced the gene expression of SIRT3

The *SIRT3* siRNA lipid complex was co-cultured with the endometrial cells to downregulate the expression of the *SIRT3* gene. Specifically, the cells were cultured with RPMI-1640 culture medium (Invitrogen, Carlsbad, CA, USA) for 24 h. Then, a mixture of *SIRT3* siRNA (600 pmol) and RNA interference (RNAi; GenePharma, Shanghai, China) mate (Lipofectamine 2000, 10 μL; Invitrogen) reagents was added to the cells in a 60-mm Petri dish containing RPMI-1640 (400 μL) on the following day. The RNAi-treated cells were then incubated under a 5% CO₂ atmosphere at 37°C for 24–48 h.

RNA extraction and real-time quantitative polymerase chain reaction (RT-qPCR)

Total RNA was extracted from endometrial tissues using the TRIzol reagent (Ambion, Austin, TX, USA) following the manufacturer's protocol. Subsequently, complementary DNA was synthesized from total RNA using the Thermo Fisher Scientific kit (Thermo Fisher Scientific). RT-qPCR was performed to analyze the expression of sirtuin genes using the SYBR Select Master Mix (Thermo Fisher Scientific). Each 20-µL qPCR reaction included 0.4 µmol/L of forward and reverse primers for sirtuin genes, and the reactions were performed in triplicate. The specific oligonucleotide primers used in this study were represented in Supplementary Table 1, http://links. lww.com/CM9/C6. The PCR cycling conditions were as follows: an initial denaturation step of 2 min at 50°C, followed by another denaturation step of 2 min at 95°C. Subsequently, 40 amplification cycles were performed, consisting of 15 s at 95°C, 15 s at 56°C, and 40 s at 72°C on the ABI 7500 Real-Time PCR System (Invitrogen).

Western blot analysis

Endometrial tissue protein extraction was performed using a protein lysate (250 mg tissues/mL) (CWBIO, Jiangsu, China), protease inhibitor (1:100; CWBIO), and phenylmethanesulfonyl fluoride (1:100; CWBIO). The concentration of extracted protein was determined using the BCA protein assay kit (Beyotime, Shanghai, China) and a microplate reader. Gels with appropriate concentrations of 10% and 5% for separation and concentration, respectively, were prepared using the Western blot kit (CWBIO). Electrophoresis was performed at 90 V for 30 min, followed by 110 V for 120 min. The resulting protein samples were transferred onto a polyvinylidene fluoride (PVDF) membrane through electrotransfer at 300 mA for 2–3 h. After washing the membrane three times with 0.1% Tris-buffered saline/Tween (TBST; Amresco, Solon, OH, USA), it was blocked with 5% skimmed milk powder in TBST for 1 h. The membrane was then incubated with a SIRT3 antibody (1:1000; Abcam, Cambridge, UK) overnight at 4°C, followed by incubation with peroxidase-conjugated donkey anti-goat IgG secondary antibody (1:2000; Beyotime) at room temperature for 1 h. Finally, the Western blot strips were detected after being washed three times with TBST.

Immunofluorescence

The endometrial cells were fixed with 4% paraformaldehyde (Sigma-Aldrich) for 30 min, followed by treatment with 1% Triton-X-100 (Sigma-Aldrich) for 7 min and incubated with 1% bovine serum albumin (BSA) blocking solution (Sigma-Aldrich) for 1 h at room temperature. The specimens were then incubated with primary antibodies for sirtuins (SIRT1 1:500; SIRT2 1:100; SIRT3 1:100; SIRT4 1:100; SIRT5 1:100; SIRT6 1:100; and SIRT7 1:50; Abcam) in blocking solution at room temperature for 1 h, followed by incubation with secondary antibodies (1:200 dilution, ZF-0314, ZF-0317; ZSGB-BIO, Beijing, China) at room temperature for 1 h. The nuclei were labeled with Hoechst 33342 (1:100 dilution, H3570; Invitrogen) in the last 5 min. Immunofluorescence was observed using a confocal microscope (Zeiss LSM 710; Zeiss, Oberkochen, Germany).

Immunohistochemical analysis

In this study, the endometrial tissues were fixed with 4% paraformaldehyde, embedded in paraffin, sectioned, and dewaxed. Antigen retrieval involved boiling the sections in citrate buffer at pH 6.0 for 25 min, followed by three washes with PBS. To eliminate endogenous peroxidase, we treated the sections with a 3% $\rm H_2O_2$ solution at room temperature for 15 min. Subsequently, the sections were incubated with the primary antibody of SIRT3 (1:200; Abcam), followed by incubation with the horseradish peroxidase–labeled secondary antibody (ZSGB-BIO), to visualize the protein expression. The immunohistochemical images were captured using a light microscope (Nikon, Tokyo, Japan).

Apoptosis detection

The endometrial cells were collected and processed for flow cytometry analysis (BD, Franklin Lakes, NJ, USA).

Briefly, the cells were digested and collected in a 15-mL centrifuge tube. Each sample contained approximately $1-5 \times 10^6$ cells, which were then centrifuged at 1000 rpm (r = 10 cm) for 5 min and resuspended after discarding the medium. Annexin V labeling solution (100 µL; BD) and fluorescein isothiocyanate (FITC) (5 µL; BD) were added to the cells, and the mixture was incubated in the dark at room temperature for 25 min. After centrifugation and one wash with incubation buffer, fluorescent 7-aminoactinomycin D (AAD) (10 µL; BD) was added and the cells were incubated again in the dark at room temperature for 20 min. Following the final centrifugation and wash, the cells were resuspended in 300 µL of incubation buffer. Flow cytometry analysis was performed using a fluorescence wavelength of 488 nm, with FITC fluorescence detected using a bandpass filter with a wavelength of 515 nm and propidium iodide detected using a wavelength greater than 560 nm.

Bioinformatics analysis

To examine the impact of SIRT3 siRNA, we employed principal component analysis (PCA) to assess the differential expression of genes across various sample types. The identification of differentially expressed genes (DEGs) in endometrial samples between individuals treated with SIRT3 siRNA and controls was conducted using the "limma" package of R software (version 3.54.2; R Foundation, Vienna, Austria). The screening criteria for DEGs were set as a *P* value <0.05 and \log_2 fold change (FC)| >1. Subsequently, the DEGs were visually represented through a heatmap and a volcano plot using the "ggplot2" package (version 3.4.2). To gain a comprehensive understanding of the biological pathways associated with the identified DEGs, gene set enrichment analysis (GSEA) was performed. The GSEA software (GSEA_4.1.0, CapitalBio Technology, Beijing, China) was employed to analyze the data, with gene sets exhibiting an adjusted *P*-value (FDR) < 0.05 deemed statistically significant.

Statistical analysis

The statistical analysis of the data was performed using SPSS software (version 26.0; IBM, Armonk, NY, USA). Each experiment was replicated at least thrice, and the results were presented as mean \pm standard deviation (or mean \pm standard error). The independent-samples t-test was employed to evaluate the differential gene expression between the two groups. One-way analysis of variance was employed to compare two or more datasets. The χ^2 test was used to analyze the embryo count and other relevant data. A P value <0.05 indicated a statistically significant difference.

Results

Clinical data

The endometrial specimens were obtained from 37 patients diagnosed with PCOS and 34 patients with male factor infertility as controls. The clinical data, including the mean age, endometrial thickness, and levels of luteinizing

hormone in the PCOS group and control group with no significant difference. The body mass index (BMI) of patients with PCOS was significantly higher than that of the controls $(26.04 \pm 3.47 \text{ kg/m}^2 \text{ vs. } 22.37 \pm 3.62 \text{ kg/m}^2)$, t = -4.353, P < 0.001), indicating that most patients with PCOS in this study were overweight or obese, possibly as a result of metabolic dysfunction. [33] Consistent with the diagnosis of PCOS, the antral follicle count (23.11 \pm 7.72 vs. 11.82 \pm 3.38, t = -7.863, P < 0.001) and the basal testosterone hormone of patients with PCOS were significantly higher than that of the controls. However, despite the abnormal metabolic state of patients with PCOS, no significant difference in the cumulative clinical pregnancy rate was observed between the PCOS and controls $(54.54\% \text{ vs. } 56.25\%, \chi^2 = 0.006, P > 0.05)$ after ART [Supplementary Table 2, http://links.lww.com/CM9/C6], which was consistent with previous findings. [34]

Expression of sirtuin family member SIRT1-7 in the human endometrium

This study reported the expression and subcellular localization of seven sirtuin family members in endometrial cells, with SIRT1 and SIRT6 showing localization to both cytoplasm and nucleus, SIRT2 mainly localizing to cytoplasm, SIRT3-5 mainly localizing to cytoplasm (mitochondria), and SIRT7 showing localization to both nucleus and nuclear membrane [Figure 1A]. Additionally, we performed qRT-PCR to examine the mRNA expression of SIRT1-7 in all human endometrial specimens and found that SIRT3-5 mRNA expression in endometrial specimens of patients with PCOS was higher than that in the control group, with SIRT3 mRNA expression showing a statistically significant increase (P < 0.05) [Figure 1B]. To verify the elevated SIRT3 expression, we conducted immunofluorescence, immunohistochemistry, and Western blot analysis and found significantly higher SIRT3 protein expression in the PCOS group compared with the control group [Figures 1C-E]. These results suggested that SIRT3 was overexpressed in the endometrium of patients with PCOS and localized not only to mitochondria but also to the nucleus in patients with PCOS compared with the control group.

Activation of anti-OS and a decrease in mitochondrial respiratory function in the endometrium of patients with PCOS

Due to the established association between SIRT3 and OS, we conducted further investigations to assess the markers of antioxidants and mitochondrial energy metabolism, which served as the primary source of ROS. We observed an upregulation in the expression of antioxidant genes, including SOD, CAT, and GPx, in the endometrium of patients with PCOS compared with the control group, although these differences were not statistically significant. Considering the upregulation of SIRT3, we inferred that the endometrium of patients with PCOS experienced chronic OS, potentially affecting mitochondrial function. We observed a decrease in the energy metabolism of mitochondria in the endometrium of patients with PCOS compared with the control group

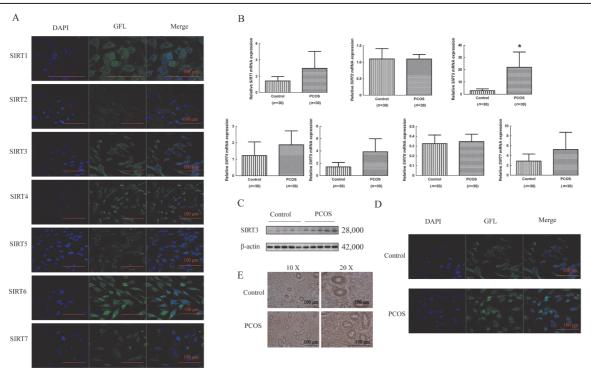


Figure 1: Expression of sirtuin family member SIRT1–7 in the human endometrium. (A) Sirtuin proteins were expressed in endometrial cells. The green fluorescence visualized these proteins. DAPI was used to visualize the nuclei. Original magnifucation \times 63; scale bar = 100 μ m. (B) Expression of mRNA of SIRT1-7. (C) Protein levels of SIRT3 in the PCOS and control groups as assessed using Western blotting. (D) Distribution of SIRT3 in the nucleus of PCOS endometrial cells was significantly higher than that in the control group. Scale bar = 100 μ m. (E) Immunohistochemical analysis in the PCOS and control groups. Scale bar = 100 μ m (*P<0.05 Vs. the control group). DAPI: Diamidino-phenyl-indole; GFL: Green fluorescence; mRNA: Messenger RNA; PCOS: Polycystic ovary syndrome; SIRT: Sirtuin.

using Seahorse Bioscience XF, particularly a significant reduction in basal respiration (P < 0.05; Figure 2). Consistent with previous findings, our results also indicated that the endometrium of patients with PCOS was in a state of chronic OS, resulting in mitochondrial dysfunction.

Modeling chronic OS in endometrial cells

To establish an OS model, we cultured the endometrial cells with a culture medium containing H_2O_2 , based on the hypothesis that it upregulates SIRT3 expression. Previous studies suggested treatment durations of 4–12 h and H_2O_2 concentrations ranging from 125 µmol/L to 500 µmol/L to induce chronic OS in endometrial cells. [35,36] However, in this study, we observed that treatment with H_2O_2 at a concentration of 125 µmol/L for 12 h could not induce obvious chronic OS, whereas a concentration of 500 µmol/L for 4 h resulted in a high number of cell deaths [Figure 3]. Therefore, based on our findings, we identified that treatment with H_2O_2 at a concentration of 250 µmol/L for 12 h was the most suitable condition for inducing chronic OS in endometrial cells.

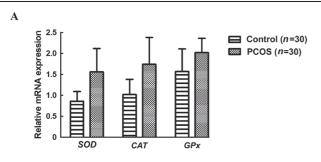
Expression of related factors and functions after the exposure of endometrial cells to 250 µmol/L H₂O₂

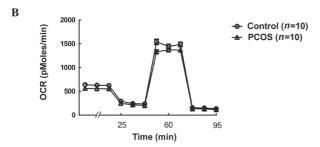
Upon exposure to 250 μmol/L H₂O₂, OS was induced in endometrial cells, leading to the upregulation of antioxidant markers and the expression of *SIRT3*, *SIRT4*, and *SIRT5* [Figures 4A, B, D, E]. It was observed that the expression

of SIRT3 did not show a significant increase after treatment with 250 µmol/L H₂O₂ for 4 h. However, after 12 h of treatment, a noticeable difference in the expression levels of SIRT3 was evident compared to the control group [Figures 4C, 4F]. The assessment of mitochondrial energy and metabolism, conducted using the Seahorse Bioscience XF assay, revealed a noticeable impairment in mitochondrial function, including basal respiration, ATP production, maximal respiration, and spare capacity [Figure 5]. Therefore, our findings suggested that chronic OS could upregulate the expression of SIRT3 and lead to mitochondrial dysfunction, indicating a potential pathogenic mechanism of PCOS and suggesting therapeutic strategies. However, we did not observe a significant decrease in the adhesion rate in the chronic OS group compared with the control group [Figure 5]. This result was consistent with the comparable cumulative pregnancy rate observed among patients with PCOS [Supplementary Table S2, http://links.lww.com/ CM9/C6]. Therefore, we believed that SIRT3 might play a vital role in maintaining endometrial receptivity by providing protection against OS.

SIRT3 protected the endometrial receptivity by anti-OS

To elucidate the role of SIRT3 in endometrial receptivity, we employed siRNA-mediated knockdown of the *SIRT3* gene in endometrial cells, followed by co-culturing with mouse embryos. The adhesion rate was used to evaluate the endometrial receptivity in *SIRT3* knocked-down cells, revealing a significantly lower adhesion rate in these cells compared with the control group (58.89% *vs.* 73.86%,





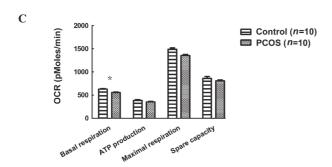


Figure 2: Elevated levels of antioxidants and decreased mitochondrial respiratory capacity in the endometrium of patients with PCOS. (A) Genes of anti-OS were upregulated in PCOS endometrial tissues. (B,C) OCR of PCOS endometrial cells decreased compared with that in the control group (*P <0.05 vs. the control group). ATP: Adenosine triphosphate; CAT: Catalase; GPx: Glutathione peroxidase; mRNA: Messenger RNA; OCR: Oxygen consumption rate; OS: Oxidative stress; PCOS: Polycystic ovary syndrome; SOD: Superoxide dismutase.

P < 0.05) after 48 h of co-culture [Figures 6A–C]. Morphological abnormalities, including cell shrinkage and small bubble-like vacuoles, were observed in SIRT3 knocked-down endometrial cells [Figures 6D-F]. These observations were confirmed by flow cytometry analysis, indicating that the cells progressed to apoptosis [Figure 6G]. We also found significant alterations in the expression of genes associated with apoptosis, including P53 (P = 0.053), P21 (P = 0.052), and BCL2 (P < 0.05),in SIRT3 knocked-down endometrial cells [Figure 6H]. The expression of genes involved in the anti-OS pathway, such as CAT (P < 0.05), SOD (P < 0.01), and GPx, was markedly reduced in SIRT3 knocked-down endometrial cells [Figure 6H]. Collectively, these findings suggested that SIRT3 was a key factor in the anti-OS pathway that influenced the proliferation and apoptosis of endometrial cells, resulting in improved endometrial receptivity.

PCA and differential gene expression analysis

The PCA indicated no notable differences between samples from the two groups [Supplementary Figure 1A, http://links.lww.com/CM9/C6]. Moreover, employing siRNA-mediated knockdown of the SIRT3 gene in endometrial cells resulted in a significant reduction in SIRT3 expression (P = 0.0063; Supplementary Figure 1B, http:// links.lww.com/CM9/C6). Subsequently, different types of samples were screened for DEGs based on the PCA results. The outcome was visualized through a volcano map and a heat map [Supplementary Figures 1C, D, http://links. lww.com/CM9/C6]. Overall, 180 DEGs were identified, with 94 genes upregulated and 86 genes downregulated. The top 10 most significant DEGs were selected and plotted accordingly [Supplementary Figure 1D, http:// links.lww.com/CM9/C6]. Further investigations using GSEA-KEGG pathway enrichment analysis of the DEGs revealed 24 enrichment pathways, particularly including the oxidative phosphorylation and apoptosis pathways

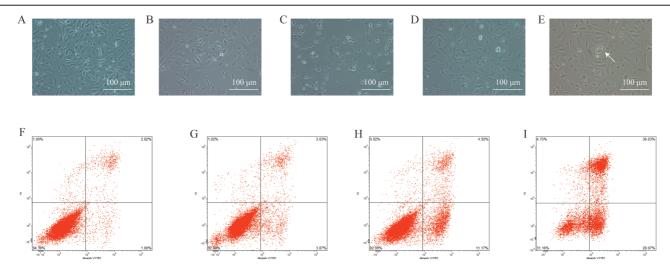


Figure 3: Model of chronic OS in endometrial cells. (A) Control group of endometrial cells. (B) State of the cells after culturing with 125 μ mol/L H_2O_2 for 12 h. No significant abnormality was observed. (C) A large number of cells died after culturing with 500 μ mol/L H_2O_2 for 4 h. (D,E) State of the cells after culturing with 250 μ mol/L H_2O_2 for 4 h and 12 h, respectively. No significant abnormality was observed. Scale bar = 100 μ m. The arrow shows cellular oncosis. (F) Apoptosis in the control group. (G,H) Apoptosis of endometrial cells cultured with 250 μ mol/L H_2O_2 for 4 h and 12 h. The apoptosis percentage was 3.03% and 4.92%, respectively. (I) A higher percentage of cell apoptosis was observed when the cells were cultured with 500 μ mol/L H_2O_2 for 4 h, reaching 36.03%. H_2O_2 : Hydrogen peroxide; OS: Oxidative stress.

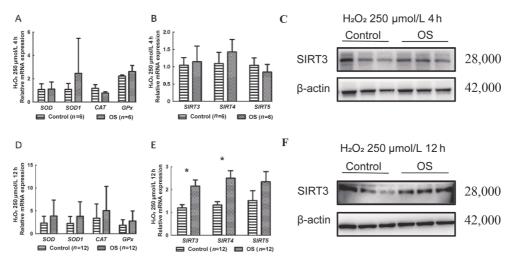


Figure 4: Expression of related factors was evaluated after exposure of endometrial cells to 250 μ mol/L H₂0₂ for 4 h and 12 h. (A–C) Endometrial cells cultured using 250 μ mol/L H₂0₂ for 4 h. The expression of anti-OS markers and *SIRT3*–5 was not significantly different between the OS and control groups. (D–F) Endometrial cells cultured with 250 μ mol/L H₂0₂ for 12 h. The expression of anti-OS markers was upregulated, and the expression of *SIRT3* significantly increased in the OS group compared with the control group (*P<0.01 V8. the control group). CAT: Catalase; GPx: Glutathione peroxidase; H₂0₂: Hydrogen peroxide; OS: Oxidative stress; SOD: Superoxide dismutase.

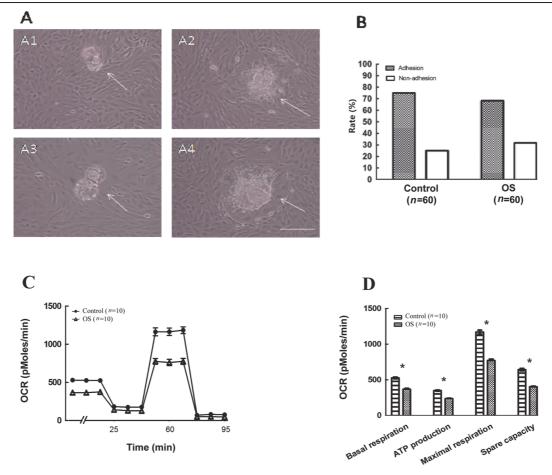


Figure 5: Expression of related functions was evaluated after exposure of endometrial cells to 250 μ mol/L H₂O₂ for 12 h. Embryos from both the non-adhesion (A1) and adhesion groups (A2) interacting with endometrial cells in the control group. Embryos of non-adhesion (A3) and adhesion groups (A4) interacting with OS endometrial cells. (B) Adhesion rate of embryos to OS endometrial cells was not significantly different from that in the control group. (C) OCR curve of endometrial cells in the OS and control groups. (D) Basal respiration, ATP production, maximal respiration, and spare capacity of the OS group significantly decreased compared with that in the control group (* $^{\circ}P$ <0.001 $^{\circ}P$ <0.001 $^{\circ}P$. Hydrogen peroxide; OCR: Oxygen consumption rate; OS: Oxidative stress.

[Supplementary Figure 1E, http://links.lww.com/CM9/C6]. These findings hold the potential for elucidating the

underlying mechanisms by which SIRT3 regulates OS in the endometrium.

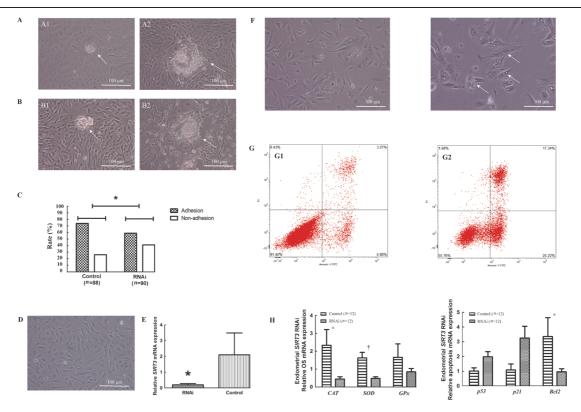


Figure 6: SIRT3 protected the endometrial receptivity by anti-OS. (A) Control group with non-implanted (A1) and implanted embryos (A2). (B) SIRT3 RNAi endometrial cells with non-implanted (B1) and implanted embryos (B2). (C) Embryonic adhesion rate of endometrial cells in the SIRT3 RNAi group was 58.89%, which was significantly lower than the adhesion rate of 73.86% in the control group. (D) Endometrial cells in the control group. (E) Expression of SIRT3 in endometrial cells was significantly inhibited by siRNA. (F) Endometrial cells of SIRT3 RNAi (G) Apoptotic rate of SIRT3 RNAi endometrial cells significantly increased compared with that in the control group. (G1) Control group. (G2) SIRT3 RNAi group. (H) Expression of markers of anti-OS, such as CAT, SOD, and GPx, significantly decreased after treatment with SIRT3 RNAi. The apoptosis pathway was significantly activated after treatment with SIRT3 RNAi. P53 and p21 were the factors promoting apoptosis, whereas Bcl2 was an anti-apoptotic factor (scale bar = 100 µm, "P < 0.05, "P < 0.01 vs. the control group). Bcl2: B-Cell lymphoma 2; CAT: Catalase; GPx: Glutathione peroxidase; OS: Oxidative stress; RNAi: RNA interference; SIRT3: Sirtuin 3; siRNA: Small interfering ribonucleic acid; SOD: Superoxide dismutase.

Discussion

PCOS is a prevalent gynecological endocrine disorder that manifests with clinical or biochemical indications of hyperandrogenism, sporadic ovulation or anovulation, and polycystic ovarian changes. It commonly co-occurs with obesity and dysregulation of glucose and lipid metabolism. Although the precise etiology and pathogenesis of PCOS remain incompletely understood, current research suggests a multifactorial origin encompassing both genetic and environmental factors. Importantly, patients with PCOS face an elevated risk of developing endometrial hyperplasia and endometrial cancer. Therefore, long-term and standardized management strategies are imperative to optimize patient care and mitigate these risks. The administration of medications to regulate menstruation, enhance fertility, and protect the endometrium of patients with PCOS can be further augmented through synergistic approaches, including the supplementation of antioxidants such as resveratrol, melatonin, quercetin, and coenzyme Q, to improve therapeutic efficacy. [37]

OS is a pathological state implicated in various diseases including PCOS.^[38] Sirtuins, a group of deacetylase proteins, play a crucial role in anti-OS, metabolic regulation, and epigenetic modification.^[22] In this study, we investigated the expression of sirtuin genes and proteins in human endometrium and observed a significant increase

in the expression of SIRT3 in patients with PCOS compared with the control group, which could be attributed to the chronic OS state. Using *SIRT3* knockdown and H₂O₂-induced OS cell models, we further demonstrated the protective role of SIRT3 in endometrial receptivity by regulating anti-OS responses and mediating the apoptosis pathway. Our findings emphasized the significance of SIRT3 in maintaining endometrial function and suggested its potential as a therapeutic target for managing PCOS.

Liu et al^[39] showed that although the total antioxidant capacity (TAC) did not exhibit a significant alteration compared with the control group, a slight increase was observed in individual antioxidant markers such as TAC, glutathione (GSH), and SOD. This increase could potentially be attributed to a compensatory response aimed at maintaining redox homeostasis. Consistent with this finding, our study also demonstrated elevated expression levels of oxidation markers, such as CAT, GPx, and SOD, in the endometrium in the PCOS group compared with the control group. Moreover, we observed impaired basal respiration in the mitochondria in the PCOS group. Mitochondria serve as both the primary source of ROS production and a highly vulnerable organelle prone to ROS-induced damage. This vulnerability arises from several factors. First, mitochondrial DNA is located close to its own oxidative phosphorylation machinery, rendering it

particularly susceptible to ROS attack and subsequent mutagenesis. Unlike nuclear DNA, mitochondrial DNA lacks histone protection, further increasing its vulnerability. Additionally, mitochondrial DNA regulates the expression of 13 subunits within the respiratory chain complex and the expression of various transfer RNAs and ribosomal RNAs within the mitochondria. Consequently, mutations affecting these genes can potentially disrupt mitochondrial function. [40] In addition, our RT-qPCR results demonstrated a significant upregulation of SIRT3, an antioxidant, in patients with PCOS compared with controls. This led us to further investigate the expression characteristics of SIRT3 protein using immunohistochemical and Western blotting techniques. Our results confirmed a significant increase in SIRT3 protein expression in the endometrial cells of patients with PCOS and suggested that SIRT3 was located in not only the mitochondria but also the nucleus. Interestingly, previous studies indicated that full-length SIRT3 was translocated to the nucleus under cellular stress conditions, [41] which was consistent with our findings. Moreover, previous studies reported that human SIRT3 was a soluble mitochondrial matrix protein that primarily regulated energy metabolism processes, influenced the flow of mitochondrial oxidative pathways, and ultimately controlled the rate of ROS production. [42–44]

In this study, we aimed to investigate the potential causes of differences in SIRT3 expression between the PCOS and control groups. The retrospective analysis of the clinical data of the two groups revealed no significant differences in basic endocrine levels. In contrast, BMI was significantly higher in patients with PCOS than in the control group. Previous studies have established a close association between obesity or overweight and OS, leading to several diseases such as metabolic syndrome. [45] Numerous studies have emphasized the role of OS in the pathogenesis of obesity and its associated risk factors, underscoring the importance of regulating body weight for better health benefits. [46] In this study, we observed a higher expression of SIRT3 in the endometrial cells of women with PCOS, suggesting a chronic imbalance in the oxidation state that impacted pregnancy outcomes and contributed to a higher clinical infertility rate. To validate this finding, we induced an endometrial oxidative state with H₂O₂ and found significantly upregulated expression of SIRT3 and classical anti-OS enzymes, similar to the PCOS endometrial specimen test results. These preliminary findings suggested that OS was one of the main pathological sources affecting SIRT3 expression. This aligned with previous studies demonstrating increased SIRT3 expression and activation of downstream targets associated with increased cell survival and decreased apoptosis, whether induced by oxidants or arising under pathological conditions.[47,48]

Over the past two decades, sirtuins have been extensively studied in mammals. These ubiquitously expressed deacetylase enzymes play a vital role in various processes associated with antioxidants and OS.^[49] SIRT3–5 are primarily localized to the mitochondria, with SIRT3 displaying robust deacetylase activity.^[44] To investigate the role of SIRT3 in anti-OS and endometrial receptivity, we

knocked down SIRT3 in endometrial cells using siRNA. Our results revealed significantly reduced antioxidant levels in SIRT3 RNAi endometrial cells, indicating a critical role of SIRT3 in the anti-OS pathway. Additionally, SIRT3 knockdown resulted in cellular abnormalities, including the accumulation of vacuoles in the cytoplasm and an increased rate of cell apoptosis. The embryo adhesion experiment using SIRT3 RNAi endometrial cells demonstrated significant damage to endometrial receptivity. These findings provided direct evidence supporting the protective role of SIRT3 in endometrial receptivity through its involvement in the anti-OS pathway. Furthermore, our study revealed significant differences in the expression pattern of nuclear SIRT3 between endometrial cells in the PCOS and control groups, indicating the upregulation of nuclear SIRT3 in endometrial cells of patients with PCOS experiencing chronic OS. Previous studies indicated that SIRT3 underwent rapid degradation under cellular stress, including OS and ultraviolet irradiation, thereby influencing the expression of stress-related mitochondrial genes encoded in the nucleus.^[41] Therefore, the rapid removal of SIRT3 from chromatin could induce the expression of genes required for the stress response. [50] Earlier findings indicated that SIRT3 localized not only to the mitochondria but also to the nucleus, deacetylating acetyl-Lys 9 and acetyl-Lys 16 of histones H3 and H4, respectively.^[51] This finding was significant because histone H3 acetylation was implicated in OS related to alcoholism, and SIRT1 activation resulted in H3 deacetylation, attenuating metabolic disease OS and complications. [52,53] Recent studies have also elucidated the close association between epigenetic modifications of H3 and the activation of the ERK pathway, which is involved in OS-mediated DNA damage and cell death.^[54] Consistent with previous findings, our results also confirmed the significance of SIRT3 in anti-OS and the increased expression of SIRT3 in the endometrial cells of patients with PCOS, particularly in the nuclear SIRT3 of endometrial cells with chronic OS, emphasizing its critical role as a key protective factor for maintaining endometrial receptivity.

Endometrial receptivity, which involves complex physiological processes including OS, plays a vital role in successful blastocyst attachment, penetration, and localized stromal changes. In this study, we demonstrated that patients with PCOS exhibited a persistent OS, with upregulated antioxidants, such as SIRT3, resulting in mitochondrial respiratory damage and impaired endometrial receptivity. Previous studies have shown that ROS levels can stimulate normal physiological functions (e.g., cell growth and proliferation), but excessive ROS can cause cellular injury (e.g., damage to DNA, lipid membranes, and proteins), which negatively affects female fertility and in vitro fertilization outcomes. [6] Gupta et al[29] extensively reviewed the role of OS in altering protein pathways, thereby contributing to the pathophysiology of female infertility. Different from the classic pathways reported in previous studies, we identified SIRT3 as an endometrial receptivity protective factor that mediated anti-OS and apoptosis pathways. The knockdown of SIRT3 using siRNA weakened anti-OS pathways while activating apoptosis pathways, leading to a significant increase in proapoptotic factors p53 and p21 and a notable decrease

in the antiapoptotic factor Bcl2 (*P* <0.05). Previous studies have established the involvement of leukemia inhibitory factor (LIF) and vascular endothelial growth factor (VEGF) in facilitating the entry of endometrial cells into the window of implantation. [55] Recent research has identified p53 as a key regulator of maternal reproduction and blastocyst implantation through regulating LIF and VEGF expression, [56,57] inducing endometrial cells to prepare for embryo implantation. [58] Therefore, we believe that the SIRT3-SOD-p53 pathway is crucial for endometrial receptivity during the pathophysiological process of PCOS, but further investigations are needed to verify this observation.

The limitation of this study should be acknowledged. This study aimed to investigate the potential enhancement of endometrial receptivity in patients with PCOS through the involvement of SIRT3, which mitigated OS while regulating cellular proliferation and apoptosis. However, due to the limited time and funding, we did not further explore the precise mechanism by which SIRT3 modulated endometrial OS.

In conclusion, this study proposed a significant role of SIRT3 in enhancing endometrial receptivity in the adverse uterine environment of patients with PCOS and also elucidated the underlying mechanisms. Specifically, SIRT3 alleviated OS damage and regulated the balance between cell proliferation and apoptosis in patients with PCOS. Therefore, SIRT3 could be a promising target for predicting and improving endometrial receptivity in this patient population.

Acknowledgments

We would like to thank all the patients and staff of the Reproductive Medicine Center, Department of Obstetrics and Gynecology, Peking University Third Hospital, for their valuable contributions to this study.

Funding

This study was supported in part by grants from the National Key Research and Development Project of China (Nos. 2021YFC2700303 and 2022YFC2702500) and the National Natural Science Foundation of China (Nos. 81925013, 82288102, 82101714, and 82201806).

Conflicts of interest

None.

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How to cite this article: Zeng ZH, Shan HY, Lin MM, Bao SY, Mo D, Deng F, Yu Y, Yang YH, Zhou P, Li R. SIRT3 protects endometrial receptivity in patients with polycystic ovary syndrome. Chin Med J 2025;138:1225–1235. doi: 10.1097/CM9.0000000000003127