

Original Article

Rehmannioside D mitigates disease progression in rats with experimental-induced diminished ovarian reserve via Forkhead Box O1/KLOTHO axis

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ABSTRACT This study aims to explore the impact of Rehmannioside D (RD) on ovarian functions of rats with diminished ovarian reserve (DOR) and its underlying mechanisms of action. A single injection of cyclophosphamide was performed to establish a DOR rat model, and fourteen days after the injection, the rats were intragastrically administrated with RD for two weeks. Rat estrus cycles were tested using vaginal smears. Ovarian tissues were histologically evaluated, the number of primordial, mature, and atretic follicles was calculated, and the apoptotic rate of granulosa cells. Follicle-stimulating hormone (FSH), luteinizing hormone (LH), and estradiol (E_2) levels were determined by ELISA assays. Protein levels of Forkhead Box O1 (FOXO1), KLOTHO, Bcl-2, and Bax were investigated in ovarian tissues of DOR rats. The binding between FOXO1 and KLOTHO was verified by ChIP assay. High-dose administration of RD into DOR rats improved their estrus cycles, increased ovarian index, enhanced the number of primordial and mature follicles, reduced the number of atretic follicle number, and ovarian granulosa cell apoptosis in addition to inhibiting FSH and LH levels and upregulating E_2 expression. FOXO1 and KLOTHO were significantly suppressed in DOR rats. FOXO1 knockdown partially suppressed the protective effects of RD on DOR rats, and KLOTHO overexpression could restore RD-induced blockade of DOR development despite knocking down FOXO1. FOXO1 antibody enriched KLOTHO promoter, and the binding between them was reduced in DOR group compared to that in sham group. RD improved ovarian functions in DOR rats and diminished granulosa cell apoptosis via the FOXO1/KLOTHO axis.

INTRODUCTION

Diminished ovarian reserve (DOR) refers to the progressive loss of the number or quality of oocytes with advanced age [1], which essentially reflects female reproductive aging [2]. Pathologic DOR (DOR occurs early or is accelerated) is a threat for female reproductive and general health [3]. In clinical practice, diagnosis of DOR is based upon increased follicle-stimulating hormone (FSH) and luteinizing hormone (LH), low levels of estradiol (E_2) and anti-Mullerian hormone (AMH), and decreased antral fol-

licle count [4]. A follicle is comprised of oocytes and two sorts of somatic cells surrounding the oocytes, granulosa and theca cells [5]. Granulosa cell apoptosis has been shown to be disordered in DOR, which has significant consequences on the generation of empty follicles, oocyte retrieval, and the quality of oocytes and embryos [6]. Currently, hormone replacement therapy (HRT) is a commonly used treatment for pathologic DOR, but the long-term outcome of HRT is poor [7]. Although there have been a number of studies on DOR, the optimal treatment for pathologic DOR is still undetermined.



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Forkhead Box O (FOXO) family is a cluster of transcription factors that mediate downstream target genes involved in cellular differentiation, growth, survival, and cell cycle [8]. Depletion of FOXO was associated with impaired oocyte maturation and blocked ovarian development [9]. FOXO1 was reported to regulate most of the FSH responsive genes in ovarian granulosa cells [10]. Increased expression of FOXO1 played a positive role in relieving follicle death [11]. KLOTHO is a type of β -glucuronidase engaged in the development and progression of reproductive diseases with anti-aging and antioxidant properties [12]. Decreased KLOTHO expression was found in women with DOR [13], and overexpression of KLOTHO was related to higher blastocyst formation frequency in porcine cell lines [14]. Our bioinformatics prediction with Jasper database indicated that FOXO1 bound to KLOTHO promoter. Previously, KLOTHO enhanced FOXO1-dependent antioxidant activity and thereby inhibited apoptosis in podocytes [15]. However, the function of the FOXO1-KLOTHO axis in DOR is largely elusive.

Rehmanniae radix preparata is an extract processed by wine-steaming the Rehmannia root, and possesses antioxidant, anti-senescence, anti-inflammatory, and neuroprotective features [16]. Kuntai capsule, a type of herb formulas consisting of six traditional Chinese herbs including Rehmanniae radix preparata, demonstrated to exert a protective role in ovarian reserve and fertility in premature ovarian failure [17]. Rehmannioside D (RD) is an iridoid compound that can be isolated from the extracts of Rehmannia glutinosa [18]. Reportedly, RD was implicated in restraining the migration of human umbilical vein endothelial cells, suggesting that it might possess an anti-angiogenesis property [19]. In addition, Zuogui Pill, in which RD is the effective component of its monarch drug rehmannia glutinosa, could significantly reduce the levels of FSH and LH, and promote AMH and endometrial repair in patients with DOR [20]. Above evidence suggests that Rehmanniae radix preparata can serve as a candidate drug for treating DOR. However, the influences of RD on DOR are little to be known. Moreover, a previous study found that catalpol, a prevalent chemical component in rehmannia glutinosa, extends lifespan and increases stress tolerance of *Caenorhabditis elegans*, partly depending on DAF-16, a FOXO family member [21]. Herein, we sought to investigate the relationship between RD and FOXO1/KLOTHO axis in relieving the progression of DOR.

METHODS

A rat model of DOR

Specific pathogen-free (SPF) female Sprague–Dawley (SD) rats (8 weeks old) in estrus were purchased. All animal experiments were ratified by the Ethics Committee of Maternal and Child Health Hospital of Jiangxi Province and performed in accordance with guide for the use and care of laboratory animals and related

regulations. After acclimation, vaginal smears were prepared using cells exfoliated from rat vagina. Next, a total of 90 female rats with an estrus cycle of 4–5 days were selected and randomly divided into sham, DOR, DOR + RD (19 mg/kg), DOR + RD (38 mg/kg), DOR + RD (76 mg/kg), DOR + RD + sh-NC, DOR + RD + sh-FOXO1, DOR + RD + sh-FOXO1 + LV-NC, DOR + RD + sh-FOXO1 + LV-KLOTHO groups (n = 10 per group).

Rats in the sham group underwent intraperitoneal injection of normal saline (1 ml) once. Rats in the model group were induced with DOR by intraperitoneal injection of 90 mg/kg cyclophosphamide (CTX; Beyotime) once [22]. Fourteen days after CTX administration, the DOR rats were gavaged with dimethyl sulfoxide (DMSO; Beyotime) or different doses of RD dissolved in DMSO (low: 19 mg/kg; medium: 38 mg/kg; high: 76 mg/kg) once a day for consecutive 2 weeks [18], during which lentiviral vectors of sh-FOXO1, sh-NC, LV-NC, or LV-KLOTHO (20 μ l, virus titer: 10^9 TU/ml) were additionally injected into both ovaries of the DOR rats according to experimental design [23,24]. Fourteen days after injection of normal saline, sham-operated rats were intragastrically administrated with DMSO.

Observation of general data

The survival of rats in each group was observed daily, and the changes in appearance signs, behaviors and activities, mental state, eating, and urine and feces were recorded.

Euthanasia of rats and sample collection and preservation

After the last intragastric administration, all rats were fasted (without water deprivation). In the next morning, the rats were anesthetized by intraperitoneal injection of 40 mg/kg sodium pentobarbital, and weighted. Rat blood was collected through cardiac puncture. Both ovaries of the rats were extracted rapidly after cervical dislocation, and adipose tissues around the ovaries was dissected under a microscope. The ovaries were rinsed with normal saline, dried, and weighed. Part of the left ovary was immediately preserved at -80°C for biochemical analysis. The right ovaries were at once fixed in 4% paraformaldehyde solution for histological analysis.

Ovarian index

Ovarian index was calculated by the following formula: ovarian index (%) = ovarian wet weight (g) / body weight before euthanasia (g) \times 100.

Assessment of estrous cycle

Vaginal smears were collected daily at 9:00 a.m. Glass slides were numbered in advance, and 20 ml of normal saline was pi-

petted, gently injected into rat vagina, and repeatedly sucked for 3–5 times. The collected solution was then evenly smeared on the glass slides. After drying, the slides were fixed with 95% alcohol for 30 min. After staining with Wright's-Giemsa stain, the slides were washed and dried. The morphology and number of exfoliated cells on each glass slide were visualized under an optical microscope.

Serum hormone analysis

Enzyme-linked immunosorbent assay (ELISA) was implemented to measure FSH, LH, and E₂ levels. Rat serum samples and ELISA kits for FSH, LH, and E₂ were taken out from a refrigerator and equilibrated for 30 min, and the levels of FSH, LH, and E₂ were tested according to the manufacturer's instructions.

Hematoxylin and eosin (H&E) staining

Ovarian tissues were fixed with 4% paraformaldehyde solution for 24 h, after which the fixative solution on the tissues was washed away with distilled water. The ovarian tissues underwent conventional dehydration in alcohol gradient (70%, 80%, 90%, 95%, and 100%) for 1 min per time, permeabilization by xylene twice (5 min per time), and paraffin embedding. Paraffin-embedded tissues were sliced into serial sections (5 μm). The sections were heated in an oven at 80°C for 1 h, hydrated by conventional alcohol gradient, permeabilized by xylene, and washed. Next, hematoxylin staining was performed for 4 min. After washing, the sections were immersed in hydrochloric acid alcohol for 10 sec and washed for 5 min. Eosin solution was appended for 2 min of staining. Histopathological changes in the tissues were observed under an optical microscope after the sections were sealed with neutral balsam. The number of primordial, mature, and atretic follicles in the ovaries was calculated.

TUNEL staining

Paraffin sections (5 μm) of ovarian tissues were dewaxed in xylene and hydrated in alcohol gradient. The sections were stained with a fluorescein-conjugated TUNEL in situ cell death assay kit, and fluorescence images were obtained under an Olympus IX-73 fluorescence microscope (Olympus).

Quantitative reverse transcription polymerase chain reaction (qRT-PCR)

Total RNA from ovarian tissues was extracted using TRIzol (15596026; Invitrogen). A PrimeScript RT reagent kit (RR047A; Takara) was used to convert total RNA into cDNA. A real-time qPCR instrument (ABI7500; ABI) was used for qRT-PCR analyses. The cDNA templates (2 μl) were added with SYBR Mix (9 μl), forward primer (0.5 μl), reverse primer (0.5 μl), and RNase Free

dH₂O (8 μl), and then amplified (95°C for 10 min, 95°C for 15 sec, and 60°C for 1 min; in a total of 40 cycles). Each sample had three duplicates. Ct value of each well was recorded, and GAPDH served as the internal control. Relative gene expression was calculated by the 2^{-ΔΔCt} method. ΔΔCt = (average Ct_{target gene in experimental group} – average Ct_{housekeeping gene in experimental group}) – (average Ct_{target gene in control group} – average Ct_{housekeeping gene in control group}). Each test was repeated thrice (see in Table 1 for primer sequences).

Western blotting

Ovarian tissues were washed thrice with pre-cooled PBS buffer and added with RIPA lysis buffer (Beyotime). A BCA protein assay kit (Beyotime) was used to evaluate protein concentration, after which the corresponding volume of protein was mixed with loading buffer (Beyotime) and heated in a boiling-water bath for 3 min. Electrophoresis was performed at 80 V for 30 min, and turned to 120 V for additional 1–2 h after bromophenol blue entered separation gels. The protein was blotted onto membranes in an ice-bath at a current of 300 mA for 60 min, after which the membranes were rinsed with washing buffer for 1–2 min and blocked in blocking buffer at room temperature for 60 min or at 4°C overnight. Incubation with anti-KLOTHO (PA5-99961, 1:1,000; Thermo Fisher Scientific), anti-FOXO1 (ab179450, 1:1,000; Abcam), anti-Bcl-2 (ab196495, 1:1,500; Abcam), or anti-Bax (ab32503, 1:1,000; Abcam) was performed on a shaker at room temperature for 1 h. The membrane was washed thrice with washing buffer for 10 min each, incubated with goat anti-rabbit immunoglobulin G (IgG) for 1 h at room temperature, and washed thrice for 10 min each. A chemiluminescence imaging system (Bio-Rad) was used to visualize the immunoblots after developer was added dropwise onto the membranes. All experiments were repeated in triplicate.

Chromatin immunoprecipitation (ChIP)-qPCR

ChIP assay was implemented with a SimpleChIP Plus sonication chromatin IP Kit (Cell Signaling Technology) according to the manufacturer's protocol. Rat ovarian tissues were fixed in 1% formaldehyde for DNA-protein crosslinking. Microson Ultrasonic Cell DisruptorXL (Misonix Inc.) was used to soni-

Table 1. Primer sequence

Name of primer	Sequences
FOXO1-F	CCCAATCTCGGAGCGACACT
FOXO1-R	CAGAGAGTACCAGGAGACGC
KLOTHO-F	GGCTCTTGCTGCTCCGTTT
KLOTHO-R	TCCGGGAAGGTGTCATGGAG
GAPDH-F	GCATCTTCTTGTCAGTGCC
GAPDH-R	GATGGTGATGGGTTTCCCGT

FOXO1, Forkhead Box O1; F, forward; R, reverse.

cate chromatin and fragmented chromatin was incubated with FOXO1 antibody or IgG antibody (negative control) at 4°C overnight. Thereafter, the immunoprecipitate was bound to protein G magnetic beads, and the crosslinking of protein and DNA was reversed and eluted DNA was purified DNA. The enrichment of DNA fragments was measured using qPCR.

Statistical analysis

GraphPad Prism software version 8.0 (Graph Pad Software Inc.) was used for the statistical analysis. Measuring data were expressed as mean \pm standard deviation. For samples with normal distribution, t-test was used for comparisons between two groups; one-way analysis of variance was implemented to compare data among multiple groups, and post-hoc analysis was conducted by

Tukey's multiple comparisons test. In terms of samples not conforming to normal distribution, Mann-Whitney U test was used for two independent samples, and statistical analysis for multiple independent samples was performed by Kruskal-Wallis. $p < 0.05$ was considered to have significant differences.

RESULTS

High-dose administration of RD improved ovarian function and reduced granulosa cell apoptosis in DOR rats

The DOR rat model was established with CTX, and the rats were gavaged with different doses of RD (low: 19 mg/kg; me-

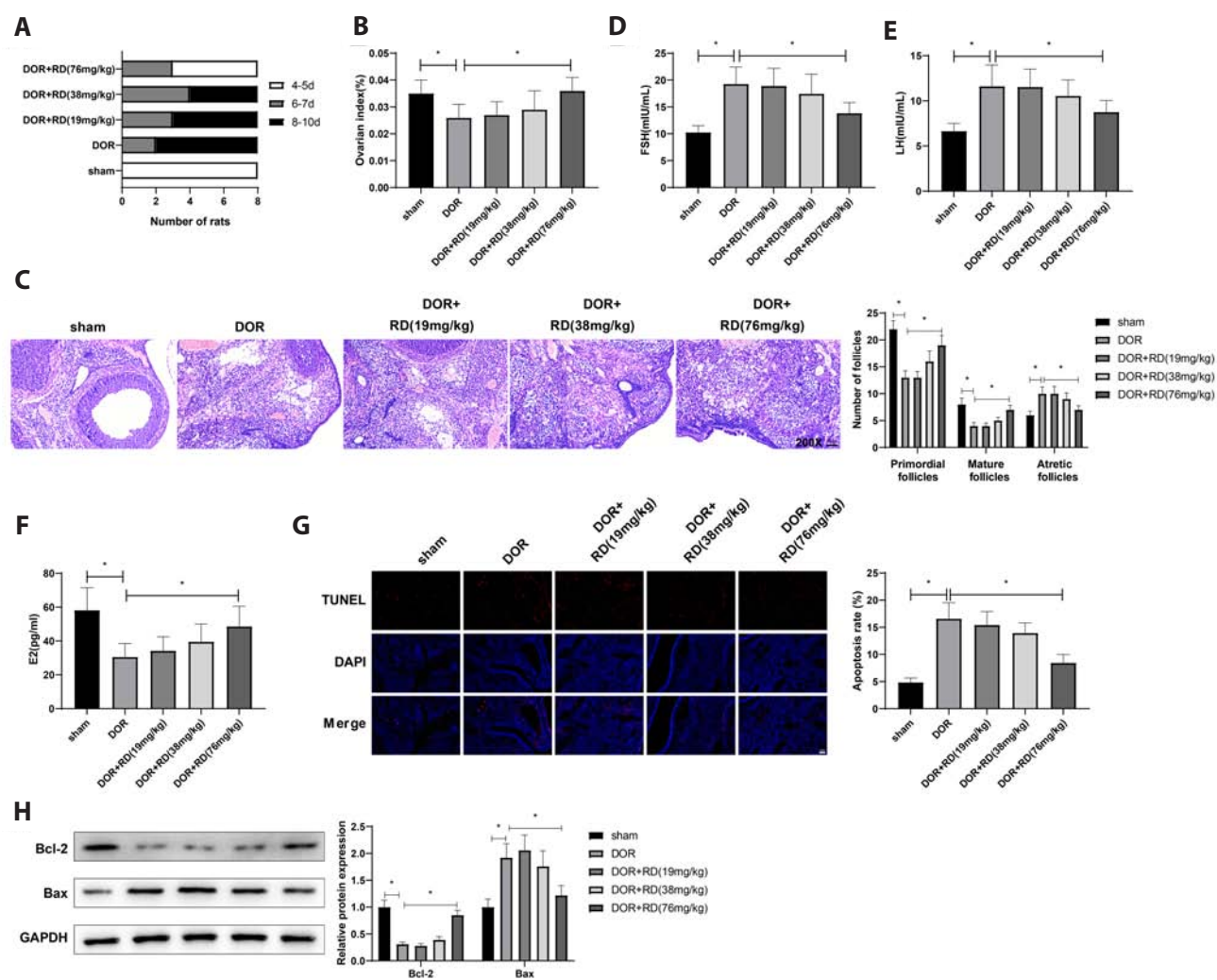


Fig. 1. High-dose administration of RD improved ovarian functions and decreased apoptosis of granulosa cells in DOR rats. (A) Vaginal smear was used to investigate estrous cycles of rats. (B) Ovarian index. (C) H&E staining of ovary, and the number of primordial, mature, and atretic follicles. (D–F) ELISA assay was used to determine the levels of FSH (D), LH (E), and E₂ (F). (G) Cell apoptosis in rat ovaries was tested by TUNEL staining. (H) The expression of Bcl-2 and Bax was measured by Western blotting. Values are presented as mean \pm SD. DOR, diminished ovarian reserve; RD, Rehmannioside D; FSH, Follicle-stimulating hormone; LH, luteinizing hormone; E₂, estradiol. $n = 10$. * $p < 0.05$.

dium: 38 mg/kg; high: 76 mg/kg) for 2 weeks. None of the rats died at the end of intragastric administration. Rats in the sham group showed normal mental state, physical appearance, behavior and activity, food intake, respiration, as well as shape and color of urine and feces. Poor spirit, lusterless and thin fur, and decreased activities appeared in rats in the DOR group. After 2 weeks of RD treatment, the mental and physical states of the rats were re-

covered, but the mental state of rats treated with high dose of RD was better than those treated with low and medium doses of RD. Compared to the sham-operated rats, the estrous cycles of rats in the DOR group was disordered and notably prolonged, and rats in the DOR + RD (76 mg/kg) group had shorter and relatively regular estrous cycles compared to rats in the DOR group (Fig. 1A). Two weeks later, the ovarian index of rats with DOR was re-

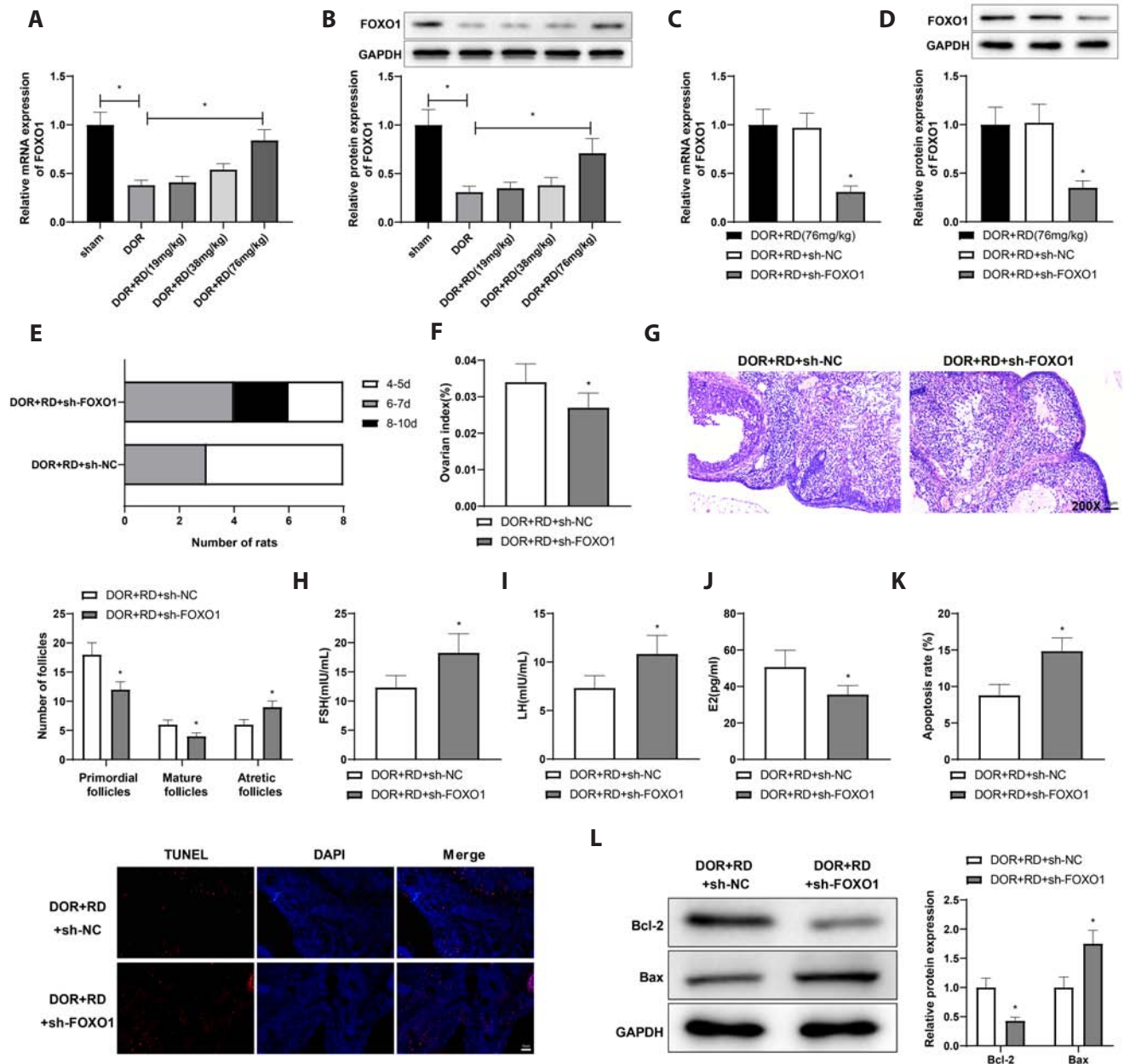


Fig. 2. RD treatment relieved DOR progression in rats by upregulating FOXO1. (A, B) qRT-PCR (A) and Western blotting (B) were implemented to investigate the expression of FOXO1 in ovarian tissues. DOR rats were injected with sh-FOXO1 or its negative control and administrated with a high dose of RD for 2 weeks. (C, D) qRT-PCR (C) and Western blotting (D) were used to determine FOXO1 in ovarian tissues. (E) The estrous cycles of rats were measured using vaginal smears. (F) Ovarian index. (G) Ovarian H&E staining, and the number of primordial, mature, and atretic follicles. (H–J) The levels of FSH (H), LH (I), and E₂ (J) were examined by ELISA assay. (K) Cell apoptosis in rat ovaries was investigated with TUNEL staining. (L) Western blotting was used to test the expression of Bcl-2 and Bax. Values are presented as mean ± SD. DOR, diminished ovarian reserve; RD, Rehmannioside D; FSH, Follicle-stimulating hormone; LH, luteinizing hormone; E₂, estradiol; FOXO1, Forkhead Box O1. n = 10. *p < 0.05.

markedly increased by treatment with high dose of RD (Fig. 1B). Results of H&E staining revealed that the structures of ovarian cortex and medulla were clear, and the morphology and structure of follicles were intact in the ovaries of sham-operated rats; rat ovaries in the DOR, DOR + RD (19 mg/kg), and DOR + RD (38 mg/kg) groups showed medullary atrophy, cortical hyperplasia, interstitial fibrosis-like changes, reduced thickness of granulosa cell layer, less primordial and mature follicles, and more ovarian atretic follicles. Compared with those in the DOR group, the structures of ovarian tissues gradually recovered, the thickness of granulosa cell layer increased, the number of atretic follicles decreased, and the number of primordial and mature follicles increased in the DOR + RD (76 mg/kg) group after 2 weeks of RD treatment (Fig. 1C). The levels of FSH and LH were obviously higher, but the level of E_2 was distinctly lower in the DOR group compared to those in the sham group. On the contrary, FSH and LH were clearly restrained, and E_2 was upregulated in the DOR + RD (76 mg/kg) group compared to those in DOR group (Fig. 1D–F). TUNEL staining demonstrated that CTX induced apoptosis

of granulosa cells, and high-dose administration of RD inhibited the damage of CTX to ovarian granulosa cells (Fig. 1G). Immunoblots displayed that Bcl-2 evidently decreased while Bax significantly strengthened in ovarian tissues of the DOR group than those in the sham group, and high-dose administration of RD facilitated the expression of Bcl-2, but suppressed the expression of Bax (Fig. 1H). Taken together, rat DOR models were successfully induced by CTX, and treatment with a high dose of RD blocked the decline in ovarian function and granulosa cell apoptosis in CTX-induced rats.

RD treatment upregulated FOXO1 expression to ameliorate DOR progression in rats

Shown by qRT-PCR and western blotting, FOXO1 expressed at a low level in the DOR group, but its expression was notably strengthened by a high-dose administration of RD for two weeks (Fig. 2A, B). A FOXO1-deficiency DOR rat model was established by ovarian injection with sh-FOXO1 or its negative control, and

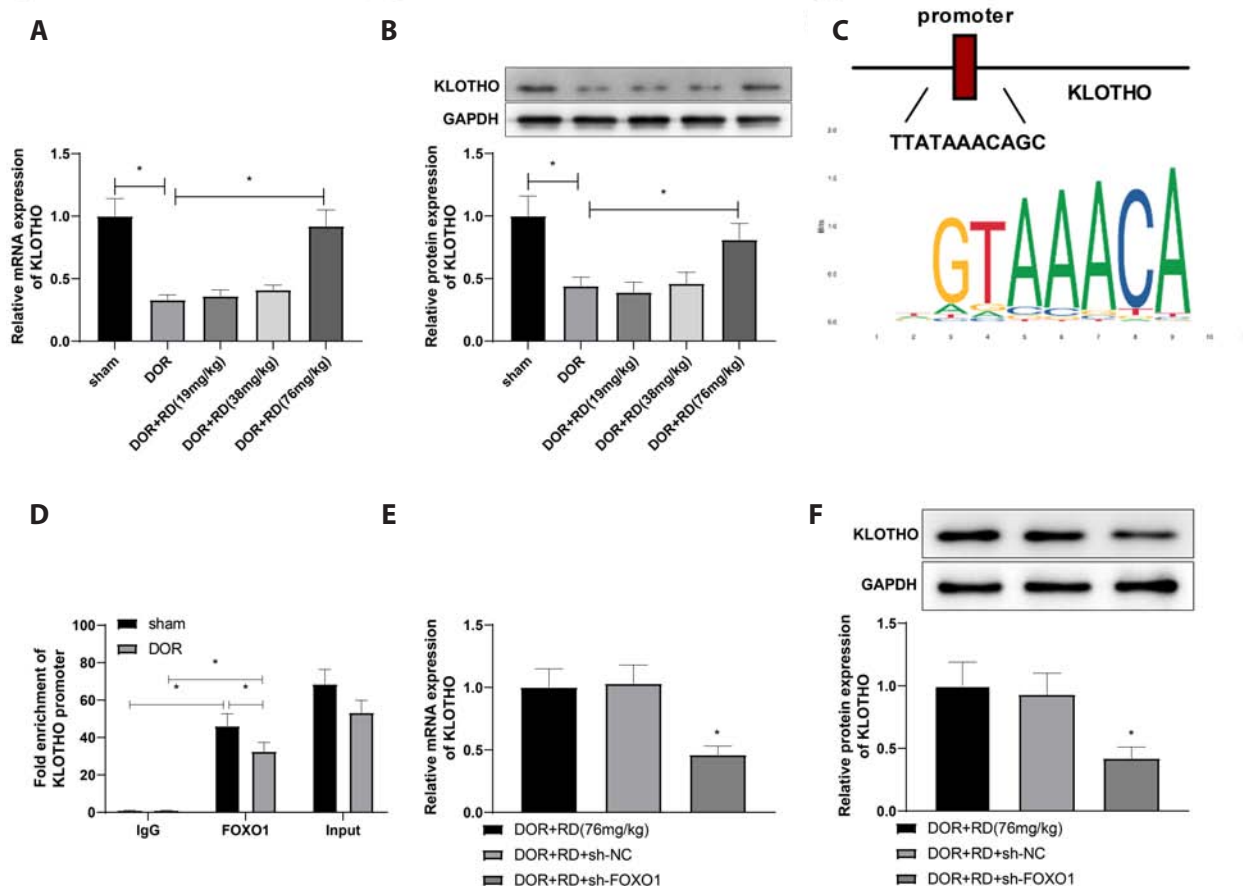


Fig. 3. FOXO1 enhanced KLOTHO transcription by binding to the KLOTHO promoter. (A, B) The expression of KLOTHO in ovarian tissues was measured by qRT-PCR (A) and Western blotting (B). (C) The binding site of FOXO1 and the KLOTHO promoter was predicted by Jasp database. (D) ChIP assay was used to detect the binding of FOXO1 to the KLOTHO promoter. (E, F) qRT-PCR (E) and Western blotting (F) were used to investigate the expression of KLOTHO in rat ovarian tissues. Values are presented as mean \pm SD. DOR, diminished ovarian reserve; RD, Rehmnnioside D; FOXO1, Forkhead Box O1; ChIP, Chromatin immunoprecipitation. $n = 10$. * $p < 0.05$.

these rats were administrated with a high dose of RD for 2 weeks. Results of qRT-PCR and western blotting indicated that ovarian FOXO1 expression was dramatically weakened in the DOR + RD + sh-FOXO1 group compared to that in the DOR + RD + sh-NC group (Fig. 2C, D). Rats in the DOR + RD + sh-FOXO1 group showed languid mental state and lackluster fur, as well as prolonged and irregular estrous cycles, compared to those in the DOR + RD + sh-NC group (Fig. 2E). The ovarian index of rats was evidently decreased in the DOR + RD + sh-FOXO1 group than that in the DOR + RD + sh-NC group (Fig. 2F). Results of H&E staining suggested slight medullary atrophy and mild gran-

ulosa cell loss in the DOR + RD + sh-NC group (Fig. 2G). Also, the histological staining showed aggravated medullary atrophy in ovarian tissues, thinned granulosa cell layer, increased number of atretic follicles, and less primordial and mature follicles in the DOR + RD + sh-FOXO1 group (Fig. 2G). The expression of FSH and LH was significantly increased and the expression of E₂ was remarkably reduced in the DOR + RD + sh-FOXO1 group compared with those in the DOR + RD + sh-NC group (Fig. 2H-J). Ovarian TUNEL staining illustrated that the apoptotic rate in granulosa cells was distinctly higher in the DOR + RD + sh-FOXO1 group than that in the DOR + RD + sh-NC group (Fig.

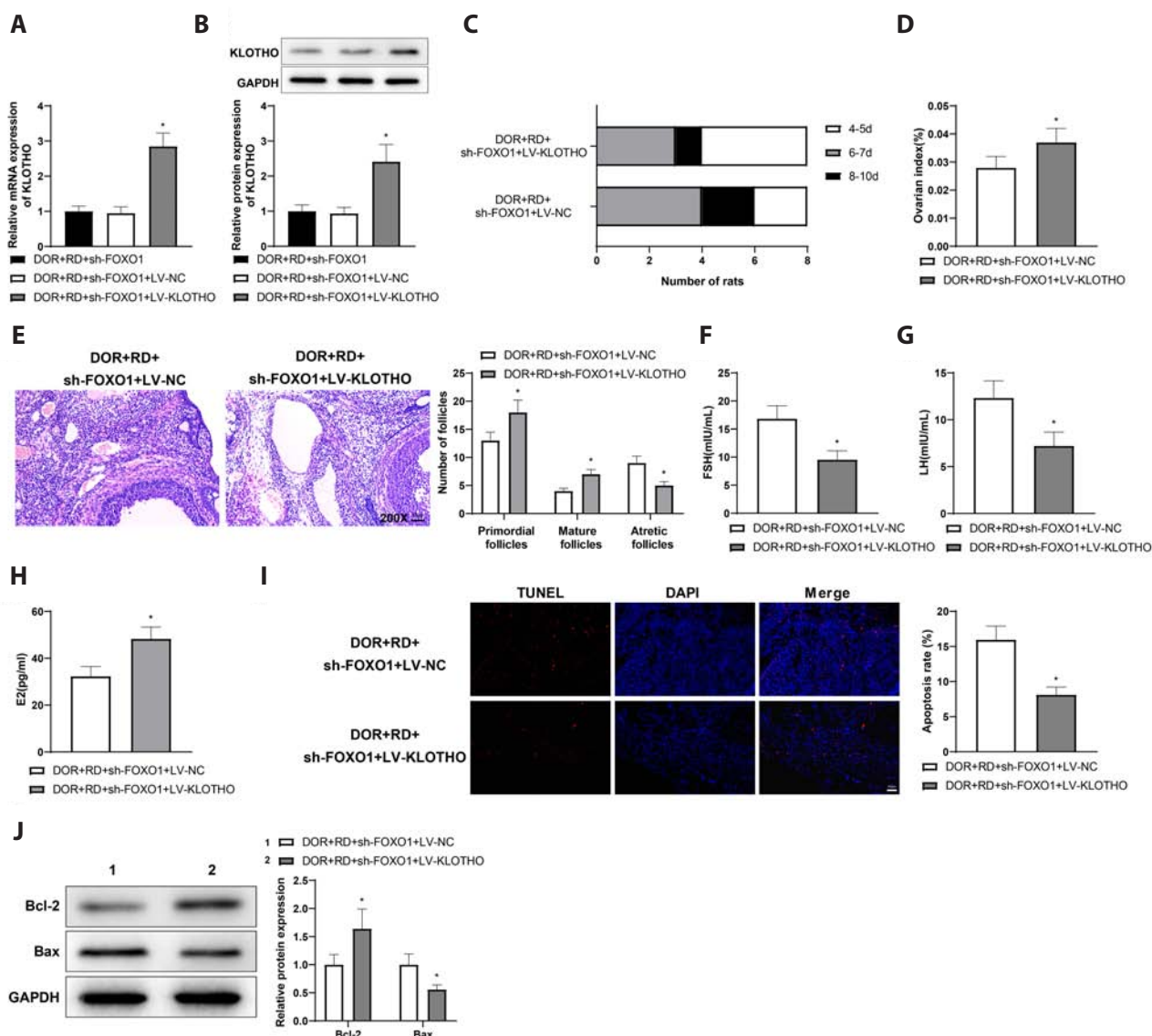


Fig. 4. RD restored ovarian functions in DOR rats through the FOXO1/KLOTHO axis. DOR rats were injected with sh-FOXO1, LV-KLOTHO, and their negative controls, in addition to intervention with a high dose RD for 2 weeks. (A, B) The expression of KLOTHO in ovarian tissues was determined by qRT-PCR (A) and Western blotting (B). (C) Vaginal smear test was used to measure the estrous cycles of the rats. (D) Ovarian index. (E) Ovarian H&E staining, and quantitative histograms of primordial, mature, and atretic follicles. (F-H) The levels of FSH (F), LH (G), and E₂ (H) were tested by ELISA assay. (I) Ovarian TUNEL staining was used to detect cell apoptosis. (J) The expression of Bcl-2 and Bax was investigated using Western blotting. Values are presented as mean ± SD. DOR, diminished ovarian reserve; RD, Rehmannioside D; FSH, Follicle-stimulating hormone; LH, luteinizing hormone; E₂, estradiol; FOXO1, Forkhead Box O1. n = 10. *p < 0.05.

2K). The results of immunoblots revealed decreased ovarian Bcl-2 expression and increased ovarian Bax level in the DOR + RD + sh-FOXO1 group (Fig. 2L). Collectively, FOXO1 was downregulated in the ovaries of rats with DOR, and RD intervention upregulated the expression of FOXO1 to improve ovarian functions and inhibit granulosa cell apoptosis.

FOXO1 bound to the KLOTHO promoter to facilitate KLOTHO transcription

qRT-PCR and western blotting demonstrated that the expression of KLOTHO was significantly downregulated in the DOR group, but remarkably increased after 2 weeks of intervention with a high dose of RD (Fig. 3A, B). Jaspar database predicted that a transcription factor FOXO1 bound to the KLOTHO promoter (Fig. 3C). For further verification, ChIP assay was conducted and displayed that FOXO1 antibody evidently enriched the KLOTHO promoter, and the binding of FOXO1 to the KLOTHO promoter was reduced in the DOR group compared to that in the sham group (Fig. 3D). Results of qRT-PCR and western blotting manifested that KLOTHO level was remarkably decreased in the DOR + RD + sh-FOXO1 group in contrast to that in the DOR + RD + sh-NC group (Fig. 3E, F). All results above illustrated that FOXO1 bound to the KLOTHO promoter to promote KLOTHO transcription.

RD treatment ameliorated ovarian functions in DOR rats via the FOXO1/KLOTHO axis

The impact of KLOTHO on DOR progression was further investigated by injection with sh-FOXO1 and LV-KLOTHO into rat ovaries. qRT-PCR and western blotting indicated that KLOTHO level was notably increased in the DOR + RD + sh-FOXO1 + LV-KLOTHO group than that in the DOR + RD + sh-FOXO1 + LV-NC group (Fig. 4A, B). When compared to the DOR + RD + sh-FOXO1 + LV-NC group, rats of the DOR + RD + sh-FOXO1 + LV-KLOTHO group showed a better mental state, as well as shorter and more regular estrus cycles (Fig. 4C). The ovarian index of rats was dramatically enhanced in the DOR + RD + sh-FOXO1 + LV-KLOTHO group (Fig. 4D). H&E staining demonstrated that the medullary atrophy in ovarian tissue was alleviated, the number of atretic follicles was reduced, and the number of primordial and mature follicles was increased in the DOR + RD + sh-FOXO1 + LV-KLOTHO group in contrast with those in the DOR + RD + sh-FOXO1 + LV-NC group (Fig. 4E). The DOR + RD + sh-FOXO1 + LV-KLOTHO group had downregulated expression of FSH and LH and increased E_2 level in rat serum (Fig. 4F–H). Ovarian TUNEL staining manifested that apoptosis of granulosa cells was significantly blocked in the DOR + RD + sh-FOXO1 + LV-KLOTHO group than that in the DOR + RD + sh-FOXO1 + LV-NC group (Fig. 4I). Western blotting displayed that Bcl-2 expression was evidently enhanced while Bax expression was distinctly

weakened in the DOR + RD + sh-FOXO1 + LV-KLOTHO group (Fig. 4J). Above results suggested that RD administration ameliorated ovarian functions in DOR rats by the FOXO1/KLOTHO axis.

DISCUSSION

DOR is clinically manifested as decreased quality of ovarian follicular reserve as a consequence of ovarian dysfunction, which is caused by various factors involving endocrine disorders and ovarian injury and leads to reduced fertility [25]. Current research has investigated the clinical therapeutics for DOR, including dehydroepiandrosterone supplementation, ovulation induction therapy, and HRT [26]. Nevertheless, the true benefit of HRT is under active debate as it has some side effects of overall increased risks of breast cancer, heart disease, stroke, and venous thromboembolism [27]. The complex pathophysiology and etiology of DOR urge more efficient treatment strategies. In the present study, the impact of RD treatment on moderating disease progression of DOR was unraveled in CTX-induced rats. In terms of specific mechanisms, RD administration suppressed apoptosis of granulosa cells and improved ovarian functions in DOR rats via the FOXO1/KLOTHO axis, ultimately mitigating DOR progression.

Western medicine believes that DOR can be attributed to advanced age and non-physiological parameters such as genetic background, surgical interventions, and therapies for cancer treatment [28]. In accordance with the theories of traditional Chinese medicine, oocytes derive from kidney essence and the basic pathogenesis of DOR is related to a deficiency of kidney essence [29]. Herein, a DOR rat model was established by CTX injection which has been widely used for DOR modelling [22]. *Rehmannia* is effective in cooling blood and hemostasis and tonifying the yin and the kidney [30]. *Rehmannia glutinosa* leaves were disclosed to restore the function of intestinal flora, mediate glycolipid level, and reduce early kidney damage in diabetes [31]. A more recent work reported that *Rehmannioside A*, derived from *Rehmannia glutinosa* Libosch, has a neuroprotective effect in cerebral ischemia-induced cognitive impairment by inhibiting ferroptosis and activating the PI3K/AKT pathway [32]. Our experimental results revealed that high-dose administration of RD inhibited the damage caused by CTX in ovarian reserve and granulosa cells. Of note, as a major cause of DOR, the apoptosis of ovarian granulosa cells has been a consistent research focus concerning the etiology of DOR [33]. FOXO1 expression was found to be downregulated in granulosa cells from patients with poor ovarian reserve [34]. Liu *et al.* [35] found an interesting association between abnormal follicular development and significantly reduced FOXO1 mRNA expression in mice. In DOR rats, FOXO1 expressed at a low level, but notably increased after 2 weeks of intervention with a high dose of RD, suggesting that RD played a positive role in amelio-

rating ovarian functions and reducing apoptosis of granulosa cells by upregulating FOXO1.

KLOTHO is an aging-suppressor gene predominantly found in the kidney and choroid plexus [36]. CTX-dependent microRNA-15 reduced the autophagy and the scavenging of reactive oxygen species in mouse ovarian granulosa cells by inhibiting α -KLOTHO expression, thereby promoting premature ovarian failure [37]. Besides, increasing β -KLOTHO expression contributed to inhibiting age-related malfunction in bovine granulosa cells [38]. In addition to the association between KLOTHO expression and granulosa cell behaviors, the relationship of KLOTHO and FOXO1 has been reported. Previously, KLOTHO and FOXO1 dephosphorylation were suppressed in tacrolimus-induced renal injury, and KLOTHO relieved renal injury by the PI3K/AKT/FOXO pathway [39]. Activated PPAR γ -KLOTHO-FOXO1 signaling pathway was participated in attenuating podocyte apoptosis [15]. Herein, FOXO1 was disclosed to facilitate KLOTHO transcription by binding to the KLOTHO promoter. To explore the effects of KLOTHO in DOR, we injected with KLOTHO over-expression lentiviruses into rat ovaries, and found the efficacy of those lentiviruses lasted for four weeks. However, considering the difficulty of blocking DOR development by KLOTHO over-expression alone, we intended to investigate the improvement of ovarian functions in DOR rats by RD treatment in relation to the FOXO1/KLOTHO axis. In DOR rats, RD treatment-promoted ovarian functions were improved by KLOTHO upregulation despite of FOXO1 silencing. FOXOs are deemed as downstream targets of PI3K/AKT, and FOXO phosphorylation is induced by phosphorylated AKT, which leads to suppression of FOXO transcription [40]. There has been convincing evidence manifesting that Zihuai recipe alleviated CTX-DOR by suppressing PI3K/AKT pathway-mediated ovarian cell apoptosis [7]. Likewise, Dingkun Pill enhanced ovarian reserve by suppressing the PI3K/AKT/mTOR signaling pathway in DOR rats [41]. The available findings suggested that the PI3K/AKT signaling pathway may be implicated in RD treatment enhancing the FOXO1/KLOTHO axis and protecting against granulosa cell apoptosis in DOR, but extensive experiments are required to validate this notion.

Collectively, this study suggested the role of RD in blocking DOR progression via the FOXO1/KLOTHO signaling pathway. To the best of our knowledge, the present study provided the first evidence for the beneficial effects of RD on alleviating DOR progression, paving a way for possible clinical trials of RD in the future. However, our finding still needs more clinical supports, and more attention could be paid on the downstream signaling pathways involved in RD-mediated protective effect in DOR, as well as the toxic effects of RD in DOR.

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CONFLICTS OF INTEREST

The authors declare no conflicts of interest.

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