Senotherapy Protects against Cisplatin-Induced Ovarian Injury by Removing Senescent Cells and Alleviating DNA Damage

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Ovarian damage induced by platinum-based chemotherapy seriously affects young women with cancer, manifesting as infertility, early menopause, and premature ovarian insufficiency. However, effective prevention strategies for such damage are lacking. Senescent cells may be induced by chemotherapeutic agents. We hypothesized that cisplatin can lead to senescence in ovarian cells during the therapeutic process, and senolytic drugs can protect animals against cisplatin-induced ovarian injury. Here, we demonstrated the existence of senescent cells in cisplatin-treated ovaries, identified the senescence-associated secretory phenotype, and observed significant improvement of ovarian function by treatment with metformin or dasatinib and quercetin (DQ) independently or in combination. These senotherapies improved both oocyte quality and fertility, increased the ovarian reserve, and enhanced hormone secretion in cisplatin-exposed mice. Additionally, attenuated fibrosis, reorganized subcellular structure, and mitigated DNA damage were observed in the ovaries of senotherapeutic mice. Moreover, RNA sequencing analysis revealed upregulation of the proliferation-related genes Ki, Prx2, Sfrp4, and Megf10, and the antioxidative gene H2-Q10 after metformin plus DQ treatment. Gene ontology analysis further revealed that combining senotherapies enhanced ovarian cell differentiation, development, and communication. In this study, we demonstrated that metformin plus DQ recovered ovarian function to a greater extent compared to metformin or DQ independently, with more follicular reserve, increased pups per litter, and reduced DNA damage. Collectively, our work indicates that senotherapies might prevent cisplatin-induced ovarian injury by removing senescent cells and reducing DNA damage, which represent a promising therapeutic avenue to prevent chemotherapy-induced ovarian damage.
1. Introduction

With the rapid development of modern medicine, multiple cancers have been rapidly, accurately diagnosed and treated in the early stage. Thus, the survival time of cancer patients has been significantly prolonged [1]. Acute lymphoblastic leukemia and breast cancer are the most common malignancies among female populations of adolescent and reproductive ages, respectively. The 10-year survival rate of the former exceeds 80%, and the five-year survival rate of the latter has reached 70% [2, 3]. Since a variety of tumor incidences gradually approach the young population, the quality of life of cancer survivors has become a growing problem needing urgently to be solved.

At present, chemotherapy remains the most prevalent treatment for cancer. Cisplatin is a first-line drug extensively used for various solid tumors, such as acute lymphoblastic leukemia, breast cancer, and cervical cancer, and it has been widely studied for its multisystem side effects [4]. Female gonads are particularly sensitive to cisplatin. Evidence suggests that the principal manifestation of cisplatin-induced ovarian damage is a loss of the healthy ovarian reserve and female fertility, frequently accompanied by increased atretic follicles [5]. The precise mechanisms underlying this drug’s ovarian toxicity involve irreversible DNA damage, overloaded oxidative stress, and uncontrolled apoptosis activation [5–7]. Fertility-preservation strategies, such as ovarian tissue cryopreservation, embryo cryopreservation, stem cell transplantation, and oocyte donation, might benefit a proportion of tumor survivors. However, practical problems, such as safety and ethical issues, have restricted their wide use [8, 9]. Currently, no better method exists for recovering cisplatin-induced ovarian damage, and effective interventions can only be carried out if the involved mechanisms are uncovered.

Cellular senescence can be triggered by various stress stimuli and is characterized by permanent cell cycle arrest, macromolecular damage, deregulated cell metabolism, and a particular senescence-associated secretory phenotype (SASP), which can be detected by short, dysfunctional telomeres, persistent DNA damage response (DDR), 

2. Methods and Materials

2.1. In Vitro Primary Granulosa Cells (GCs) Culture and Drug Administration. Ovaries were isolated from euthanized 3-week-old female C57BL/6 mice and punctured thoroughly to release GCs. The harvested GCs were centrifuged at 800 revolutions per minute (rpm) for 5.5 minutes, incubated in DMEM/F12 (12400-024; Gibco Laboratories, Gaithersburg, MD, USA) culture medium, and then divided into 4 groups. The GCs of the blank control group (NC group) were treated with DMEM/F12 growth medium without any other treatment. For the cisplatin group (Cis group), cisplatin plus metformin group (Cis+M group), and cisplatin plus DQ group (Cis+DQ group), cisplatin (30 μmol/L) was added to the medium after GC attachment. Following 24 hours, the whole medium of each group was replaced. GCs in the NC group and Cis group were cultured in DMEM/F12 growth medium for another 24 hours, and the other two groups were treated with metformin (5 mmol/L) or D+Q (D: 1 nmol/L; Q: 20 μmol/L) for another 24 hours.

2.2. Cell Culture Medium Cytokine Detection. After drug administration, the cell culture medium of the NC group and Cis group was collected. The cytokines accumulated in culture medium after different medications were determined using a Luminex mouse 23 cytokine array (Wayen Biotechnologies, Shanghai, China).

2.3. Animal Care and Drug Administration. One hundred sex-mature 6-week-old female mice were provided by the Tongji Hospital Experimental Animal Center, which is affiliated with the Tongji Medical College of Huazhong University. Animals were acclimated to the environment for one week before drug administration on an alternating 12-hour light/dark cycle with free access to food and water. Then, 20 mice were administered dasatinib (5 mg/kg) and
quercetin (50 mg/kg) intragastrically 2 times weekly for 4 weeks and/or received metformin-containing water (2 mg/mL) or pure water over a period of 4 weeks (Cis+DQ, Cis+M, cisplatin plus metformin, and DQ [Cis+M+DQ] groups, 20 mice per group, 60 mice in total). One week after senotherapy administration, the aforementioned 60 mice were simultaneously subjected to intraperitoneal injection with 2.5 mg/kg of cisplatin 3 times weekly for the next 3 weeks. Another 20 mice received saline or cisplatin injection only at 7 weeks of age (NC and Cis groups, 20 mice per group, 40 mice in total). All animal operations were performed in compliance with the guide approved by the ethics board.
DQ (D: 5 mg/kg; Q: 50 mg/kg) M+DQ combined treatment

Ovary extraction for ovarian function and cell senescence analysis (n = 15)

M (2 mg/mL)

7w 8w Cis (2.5 mg/kg) 11w Mating test (n = 5) 24w

18y 20y 24y–25y 35y

Sβ-gal

Cis Cis+M Cis+DQ Cis+M+DQ

(a)

(b)

p16

p21

IL-6

(c)

Figure 2: Continued.
committee of Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology, in the People’s Republic of China.

2.4. Senescence-Associated- (SA-) β-Gal Staining. SA-β-gal staining of GCs or frozen ovarian slices was conducted according to the product manual. In brief, for GCs, samples were pre-washed with phosphate-buffered saline and then fixed with the fixative solution provided in the Senescence β-Galactosidase Staining Kit (C0602; Beyotime Biotechnology, Haimen, China). For frozen ovarian sections, ovarian slices were dewaxed and hydrated, and then we processed for the staining step.

2.5. Sirius red Staining. A series of representative ovarian sections in different medication groups were chosen to identify the fibrosis level using Sirius red staining. Initially, we dewaxed and hydrated the ovarian slices and processed them with 10-minute celestine blue staining. After 3 washes with ddH₂O, the ovarian sections were subjected to Sirius red solution for 30 minutes. Following another round of dehydration, we mounted ovarian slices with neutral resins for subsequent imaging and quantitative analysis.

2.6. Immunoblot and Quantitative Analyses. In brief, total ovarian proteins or proteins from cell lysates were extracted and separated by 10% sodium dodecyl sulphate–polyacrylamide gel electrophoresis (SDS-PAGE) gel or 12% SDS-PAGE gel according to the molecular weight of the proteins to be measured. Targeted proteins were subsequently transferred to polyvinylidene fluoride membranes and blocked with rapid blocking buffer for 30 minutes. After 3 washes with Tris-Buffered Saline and Tween 20 (TBST), targeted proteins were incubated with specific primary antibodies against nuclear factor kappa-light-chain-enhancer of activated B-cells (NF-κB) (Cat# 10745-1-AP, 1:1000; Proteintech, Rosemont, IL, USA), P16ink4a (Cat# 10883-1-AP, 1:500; Proteintech, Rosemont, IL, USA), P21WAF1/Cip1 (Cat# A1438, 1:1000; ABclonal Technology, Woburn, MA, USA), interleukin (IL)-6 (Cat# A0286, 1:1000; ABclonal Technology, Woburn, MA, USA), IL-1β (Cat# A19635, 1:500; ABclonal Technology, Woburn, MA, USA), proliferating cell nuclear antigen (PCNA) (Cat# sc-56, 1:1000; Santa Cruz Biotechnology, Dallas, TX, USA), heme oxygenase (HO)-1 (Cat# 10701-1-AP, 1:1000; ABclonal Technology, Woburn, MA, USA), and superoxide dismutase (SOD) 2 (Cat# Ag21388; ABclonal Technology, Woburn, MA, USA) diluted in TBST overnight in a 4°C refrigerator. After 30 minutes of rewarming, target blots were washed with TBST 3 times and incubated with appropriate secondary antibodies for 1 hour the next day. Ultimately, blots were imaged using the chemiluminescence method after 3 washes.

2.7. Immunohistochemistry (IHC) and Immunofluorescence (IF) Staining. Three representative sections of ovaries in distinct groups were selected for IHC and IF analyses. The former was used to detect the localization and quantitative expressions of ovarian P16ink4a, P21WAF1/Cip1, IL-6, and 8-hydroxy-2-deoxyguanosine (8-OHdG) (Cat# sc-393871, 1:100; Santa Cruz Biotechnology, Dallas, TX, USA), and the latter was applied to identify zona pellucida protein 3 (ZP3) protein expression (ab70384, 1:1000; Abcam, Cambridge, UK). IHC and IF were performed using routine protocols as previously described [25].

2.8. Terminal Deoxynucleotidyl Transferase dUTP Nick End Labeling (TUNEL) Cell Apoptosis Analysis and 5-Ethynyl-2'-Deoxyuridine (EdU) Cell Proliferation Analysis. Ovarian sections were first dewaxed and hydrated, and then we performed staining steps according to the manuals of the Colorimetric TUNEL Apoptosis Assay Kit (C1098; Beyotime Biotechnology, Haimen, China) and BeyoClick™ EdU Cell Proliferation Kit with Alexa Fluor 488 (C0071S; Beyotime Biotechnology, Haimen, China).

Figure 2: Metformin or DQ alone or in combination with each other alleviated the senescent stress induced by cisplatin in vivo. (a) Flowchart of drug administration animal experiment. (b, c) Light microscopic images of SA-β-gal staining and P16ink4a, P21WAF1/Cip1, and IL-6 IHC staining after different medications. Scar bars are marked on respective images (n=3). (d–g) Quantitative analysis of SA-β-gal–positive area and P16ink4a-, P21 WAF1/Cip1-, and IL-6-positive IOD (n=3). Results are presented using mean ± standard deviation values. *p < 0.05, **p < 0.01, ***p < 0.001.
Figure 3: Continued.
2.9. Transmission Electron Microscope. Three ovaries per group were preserved in electron microscope preservation solution (G1102; Servicebio, Wuhan, China). Transmission electron microscopy images of each ovary sample were taken by ServiceBio (Wuhan, China).

2.10. Estrous Cycle Analysis. Vaginal smears of all animals were sampled at 9:00 a.m. for 15 consecutive days from the day drug administration was finished. Vaginal secretions were collected with saline and then spread on slides, followed by hematoxylin and eosin (HE) staining after air drying. Then, the stage of the estrous cycle was determined by cytology by 2 people separately under a microscope.

2.11. Follicle Counting. Five ovaries from distinct individuals in each group were obtained for follicle counting. Dissected ovaries were preserved in 4% paraformaldehyde and then cut into 5 μm pieces. Follicles at different levels were counted by 2 people separately. The detailed procedures were performed as previously described [26].

2.12. Serum Anti-Müllerian Hormone (AMH) and Estradiol (E₂) Determination. Nonhemolytic blood samples of female mice were collected before sacrifice. Levels of serum AMH and E₂ were determined using the Mouse AMH ELISA Kit and Mouse E₂ ELISA Kit according to the manufacturer’s instructions (CSB-E13156 m and CSB-E05109 m; Cusabio Technology LLC, Houston, TX, USA).

2.13. Mating Test. Five female mice in each group were randomly selected for the mating test. The mating test began once the drug administration ended. In brief, 10 days were designated as a mating cycle, and 2 female mice with 1 male mouse were arranged in one cage for mating. We tested the vaginal plugs every morning at 9:00 a.m. to evaluate for a pregnant outcome. The pregnant mice were then allocated to an independent cage, and their pup conditions were recorded. Ten days later, nonpregnant mice were marked and separated from male mice. After the first round parturition ended, another mating cycle was repeated. The duration of the mating test was 6 months in total.

2.14. Statistical Analysis. Statistical analyses were conducted using the SPSS 26.0 software (IBM Corporation, Armonk, NY, USA). Group differences of continuous variables were calculated using one-way analysis of variance (nonparametric tests). Group differences in regular cycle numbers were analyzed using chi-squared analysis. p < 0.05 was considered statistically significant.

3. Results

3.1. Confirmation of Senescent Ovarian Cells Induced by Cisplatin and Alleviated by Senotherapies. To determine whether cisplatin could induce ovarian cell senescence, we exposed ovarian GCs of female mice to cisplatin in vitro. The flowchart of in vitro experiment is shown in Figure 1(a). We first screened senescent stress via SA-β-gal staining. As shown in Figures 1(b) and 1(c), SA-β-gal–positive staining was remarkably increased in cisplatin-treated GCs relative to the control group (p < 0.05). To better investigate the secretory phenotype, cell culture medium following cisplatin administration was collected and determined using a Luminex mouse 23 cytokine array (Wayen Biotechnologies, Shanghai, China). Compared to the control, IL-6 obviously accumulated in the Cis group culture medium (p < 0.05, Figure 1(d)). Other inflammatory cytokines, such as granulocyte colony-stimulating factor (G-CSF) and IL-12 (P40), showed an upward trend following cisplatin administration without statistical significance (Figures 1(e) and 1(f)). The above results indicated that cisplatin might induce granular cell senescence.
Figure 4: Metformin or DQ alone or in combination with each other pretreatment reorganized the ovarian subcellular structure and preserved mice fertility. (a) General state of pups in different groups. (b–d) Successful pregnant rate, pup mortality rate, and pups per litter of the different drug-administration groups (n = 5). (e) Transmission electron microscopy images revealed subcellular morphologies of different medication-treated ovaries (n = 3). Red arrow: mitochondria; white arrow: endoplasmic reticulum (ER); blue arrow: vacuolated cytoplasm; N: nucleus of the GC. The low magnification was 3,000x and the high magnification (insets) was 10,000x. (f) Immunofluorescence of ZP3 in ovaries treated with different medication regimens (n = 3). Results are presented using mean ± standard deviation values. **p < 0.01, ***p < 0.001.
To further explore whether senotherapies metformin and/or DQ may alleviate the senescence stress caused by cisplatin in vitro, we undertook western blot analysis of senescence-related markers. Quantitative analysis of senescence-related proteins in each group revealed significant upregulation of the senescence marker P21<sub>WAF1/Cip1</sub> and activation of NF-κB in cisplatin-exposed GCs lysates (p < 0.05, Figures 1(g)–1(l)); however, another senescence effector, P16<sub>link4a</sub>, only shows an increased tendency (Figure 1(k)). Conversely, DQ administration led to an obvious downregulation of the senescence markers P21<sub>WAF1/Cip1</sub> and P16<sub>link4a</sub> (p < 0.05) (Figures 1(k) and 1(l)), and metformin treatment only reduced P21<sub>WAF1/Cip1</sub> expression (p < 0.05, Figure 1(l)), suggesting metformin had a less profound senolytic effect than DQ. Western blot analysis of the SASP markers indicated that metformin excessively prohibited inflammatory cytokine secretion, and IL-6 and IL-1β in cell lysates were highly enriched after metformin treatment (p < 0.05, Figures 1(h) and 1(i)). Collectively, these results indicated that metformin and DQ variously attenuated the cisplatin-induced ovarian cell senescence burden in vitro. Of note, DQ exhibited a prominent senescent GC elimination capacity, and metformin probably acted via effective blockade of SASP secretion.

3.2. Removal of Senescent Cells by Metformin or DQ Alone or Both in Combination following Cisplatin Chemotherapy. To further reveal whether metformin and/or DQ intervention could mitigate senescent stress in vivo, we established a cisplatin-induced ovarian damage animal model to assess SNCs after the aforementioned medication. The flowchart of the animal experiment is shown in Figure 2(a). SA-β-gal staining of ovarian sections showed a trend consistent with that of the in vitro experiments. In particular, enlarged SA-β-gal–positive staining of mouse ovaries was recorded after cisplatin administration, and metformin and/or DQ intervention resulted in various shrinkages of the SA-β-gal–positive staining area in the ovaries (p < 0.05, Figures 2(b) and 2(d)). Likewise, we determined senescence-associated markers in ovaries post cisplatin exposure. IHC results revealed remarkably increased positive expressions of P16<sup>link4a</sup>, P21<sub>WAF1/Cip1</sub>, and IL-6 in cisplatin-treated ovaries, suggesting that senescence stress was also induced by cisplatin in vivo (Figure 2(c)). Alone or combined treatment with metformin and DQ significantly inhibited the expression of P16<sup>link4a</sup>, P21<sub>WAF1/Cip1</sub>, and IL-6 (Figure 2(c)). Quantitative analysis of integral optical density (IOD) indicated that metformin plus DQ treatment removed the maximum senescence signal (p < 0.05, Figures 2(e)–2(g)). In summary, these data validated that senescent cells do exist in cisplatin-treated mouse ovaries, and ovarian senescence stress induced by cisplatin can be effectively removed by metformin and/or DQ treatment in vivo. In particular, metformin plus DQ treatment maximally eliminated the senescent burden in the cisplatin-exposed ovarian setting.

3.3. Reversal of Ovarian Function by Metformin or DQ Alone or Both in Combination against Cisplatin-Induced Damage. To further uncover whether metformin and/or DQ senotherapies protect ovaries from cisplatin-induced ovarian dysfunction, we undertook ovarian fibrosis evaluation, ovarian follicle counting, and estrous cycle and serum hormone determination in each group. Series of ovarian Sirius red staining indicated obvious fibrosis in cisplatin-treated ovaries, which was significantly alleviated following metformin and/or DQ administration (p < 0.05, Figures 3(a) and 3(h)). Ovarian follicle counting directly reflected the ovarian reserve. In this study, primordial follicles and growing follicles were largely reduced in cisplatin-treated ovaries, and atretic follicles were significantly elevated (p < 0.05, Figure 3(c)). However, metformin and/or DQ treatment restored the growth and development of ovarian follicles (Figure 3(c)). Notably, combination administration of both maximally preserved ovarian reserve and prohibited follicle atresia caused by cisplatin (p < 0.05, Figure 3(c)).

The estrous cycle is thought to be a reflection of ovarian endocrine function. We next assessed irregular estrous cycle frequencies in each group. Remarkably, cisplatin treatment disturbed estrous cycles in mice, as the percentage of irregular estrous cycles was up to 78.9% (p < 0.05, Figure 3(d)), with irregular estrous cycle rates of 50%, 47.4%, and 38.9% in the Cis+M group, Cis+DQ group, and Cis+M+DQ group, respectively (p < 0.05, Figure 3(d)). Moreover, the ovarian index was obviously reduced in cisplatin-treated mice, whereas senotherapies (Cis+M, Cis+DQ, and Cis+M+DQ) led to a significant recovery (p < 0.05, Figure 3(d)). To evaluate murine ovarian endocrine function post medication accurately, we measured the serum concentrations of E2 and AMH. Consistent with previous reports, serum E2 levels dramatically decreased after cisplatin exposure (p < 0.05, Figure 3(g)). Combination treatment with metformin and DQ reversed the decline in serum E2 levels in mice; however, no significant improvement in serum E2 levels was observed in the Cis+M and Cis+DQ groups (Figure 3(g)). Markedly decreased serum AMH (a predictive biological marker of ovarian reserve) in cisplatin-treated mice was also observed, while elevated serum AMH was recognized in the Cis+M, Cis+DQ, and Cis+M+DQ groups compared with cisplatin-exposed mice (p < 0.05, Figure 3(f)). Similarly, serum AMH was significantly improved in the Cis+M+DQ group, albeit remaining barely different from the NC group mice (Figure 3(f)). Taken together, cisplatin severely impaired mouse ovarian health reserve and function, and senotherapies metformin and/or DQ effectively reversed the injury.

3.4. Preservation of Fertility and Reshaping of Ovarian Subcellular Structure by Metformin or DQ Pretreatment. To identify whether metformin and/or DQ mitigates cisplatin-induced reproductive dysfunction and oocyte impairment, we performed a mating test. The general status of pups in each medication group is shown in Figure 4(a). The successful pregnancy rate in cisplatin-treated mice was significantly lower than that in the NC group, which was recovered in the Cis+M, Cis+DQ, and Cis+M+DQ groups (Figure 4(b)). Meanwhile, the pup mortality rate of cisplatin-treated mice increased compared to that of NC mice, and administration of metformin or DQ alone or both in combination seemed to decrease pup mortality (Figure 4(c)). Additionally, the
Figure 5: Continued.
average pup number per litter in the Cis+M, Cis+DQ, and Cis+M+DQ groups mice was significantly greater than that in cisplatin-treated mice ($p < 0.05$, Figure 4(d)). Notably, the number of pups per litter was indistinguishable between the combo medication group and the NC group (Figure 4(d)).

Healthy follicles constitute the elementary units of female ovaries. GCs serve as vital compartments of the follicle unit, suffering severe damage during cisplatin exposure. To disclose the influence of SNCs on granular subcellular structure, transmission electron microscopy was conducted to record subcellular morphologic alternations of GCs in distinct groups (Figure 4(e)). The majority of GCs were well aligned with integrated nuclear membranes, visible nucleoli, and evenly distributed chromatin in the NC group. Abundant good shapes of the subcellular organelles were captured by cameras, including oval and complete mitochondria with clear cristae and plentiful and enlarged reticulum with ribosome attachments, suggestive of functional protein synthesis. Conversely, the GCs in the Cis group showed a morphology of severely damaged subcellular ultrastructure, characterized by expansive perinuclear space, inhomogeneous distributed chromatin, increased heterochromatin, and even emergence of plasmarrhexis. The functional organelles also suffered from cisplatin exposure. We noticed that mitochondria were dissolved into vacuoles accompanied by discrete cristae, and the endoplasmic reticulum was swollen and even ruptured in cisplatin-exposed GCs. However, metformin, DQ, and metformin plus DQ combined treatment attenuated the subcellular deterioration caused by cisplatin. Compared to cisplatin-treated GCs, the distribution of chromatin was relatively uniform, along with rare heterochromatin, clear nuclear membrane, and limited perinuclear space after senotherapies. The subcellular organelles of the senotherapeutic groups appeared to have a good outline under transmission electron microscopy. The above results illustrated that cisplatin exerted a robust toxic effect on granular subcellular structure, and metformin, DQ, and metformin plus DQ in combination may counteract the subcellular toxic effect caused by cisplatin.

Essentially, oocyte quality is the primary determinant of female fertility and reproductive health. Previous literature has reported the direct impairment of oocytes caused by cisplatin [27]. Ovarian section ZP3 IF was further applied to assess oocyte quality via evaluation of zona pellucida conditions. Remarkably reduced healthy follicles but increased shrinking atretic follicles in cisplatin-treated ovaries were recorded via ZP3 staining. The administration of metformin and/or DQ was associated with an upregulation of normal ZP3 staining, indicating a restoration of oocyte quality following senotherapies (Figure 4(f)). These above results indicated that metformin and/or DQ administration reorganized the granular subcellular structure, preserved female mice fertility, and prohibited oocyte deterioration after cisplatin treatment.

3.5. Alleviation of Cisplatin-Induced Ovarian DNA Damage by Metformin or DQ Alone or Both in Combination. As reported, DDR is an important inducer of cellular senescence [28]. Previous studies have acknowledged DNA damage in other tissues caused by cisplatin and the emerging therapeutic effects of senolytics and senomorphics on organ dysfunction [13, 29]. Thus, we measured DNA damage levels via 8-OHdG IHC staining to identify the possible ovarian genotoxicity caused by cisplatin and the protective
GO enrichment analysis (Cis vs Cis+M)

- Regulation of vasoconstriction
- Regulation of blood vessel diameter
- Regulation of tube diameter
- Cholesterol transport
- Regulation of tissue size
- Sterol transport
- Extracellular region
- Vascular process in circulatory system
- Regulation of smooth muscle contraction
- Positive regulation of heat generation
- Intercellular cholesterol transport
- Intercellular sterol transport
- Circulatory system process
- Regulation of heat generation
- Extracellular structure organization
- Positive regulation of blood circulation
- Phospholipid efflux
- Positive regulation of multicellular organismal process
- Regulation of blood circulation
- Apolipoprotein binding
- Response to endogenous stimulus
- Ovulation
- Negative regulation of lipid catabolic process
- Sterol transfer activity
- Cholesterol transfer activity
- Cholesterol metabolic process
- Response to organonitrogen compound
- Response to lead ion
- Secondary alcohol metabolic process

Figure 6: Continued.
GO enrichment analysis (Cis vs Cis+DQ)

- Positive regulation of development process
- Regulation of developmental process
- Positive regulation of cell differentiation
- Regulation of differentiation
- Positive regulation of multicellular organismal process
- Anatomical structure development
- Cell communication
- Regulation of multicellular organismal process
- Collagen-containing extracellular matrix
- Extracellular matrix
- Extracellular region
- Extracellular region part
- Signaling receptor binding
- G protein-coupled receptor binding
- Development process
- Positive regulation of biological process
- Negative regulation of cellular process
- Regulation of signal transduction
- Regulation of cell communication
- Regulation of cellular process
- Protein binding
- Movement of cell or subcellular component
- Cellular development process
- DNA-binding transcription factor activity, RNA polymerase II-specific
- Regulation of biological process
- Regulation of interleukin-2 production

Figure 6: Continued.
effects of senotherapeutic agents. Increased positive staining of 8-OHdG was observed in the Cis group, and western blot analysis revealed a significant decline in the antioxidative markers HO-1 and SOD2 in ovaries following cisplatin chemotherapy, indicating dramatic DNA oxidative injury and unmatched antioxidative capacity in cisplatin-treated ovaries ($p < 0.05$, Figures 5(a), 5(d), 5(g), and 5(h)). The above indexes were variously restored in ovaries treated by Cis+M, Cis+DQ, and Cis+M+DQ; among these, the combination treatment (Cis+M+DQ) ensured the largest elimination of ovarian DNA oxidative damage.

Additionally, the senolytics are suggested to take effect via the modulation of cell apoptosis [30]. To determine whether cisplatin-induced cell senescence affected ovarian cell viability, we performed TUNEL- and EdU-specific staining of ovarian sections to assess the status of cell apoptosis and proliferation in ovaries subjected to distinct treatments. Cisplatin chemotherapy resulted in severe apoptosis of ovarian cells (primarily GCs, small fraction of interstitial cells), while metformin, DQ, and metformin plus DQ co-administration treatment decreased the TUNEL staining ($p < 0.05$, Figures 5(b) and 5(e)). Positive EdU staining was almost invisible in cisplatin-
3.6. Positive Regulation of Cell Development in Cisplatin-Induced Ovaries by Senotherapeutic Treatment. To explore the molecular events induced by senotherapies in cisplatin-treated ovaries, we employed transcriptome sequencing of ovaries in the different drug-administration groups. Venn diagrams show the number of differentially expressed genes (DEGs) between NC and Cis ovaries along with the rescued genes in different senotherapeutic groups (Figures 6(a) and 6(b)). Specifically, cisplatin exposure resulted in the downregulation of 73 genes in the ovaries, among which metformin, senolytic DQ, and metformin and DQ together rescued 1, 3, and 70 of them, respectively (Figure 6(b)). Thirty-five cisplatin-upregulated genes were modulated following senotherapies—in particular, metformin, senolytic DQ, and metformin and DQ together reset 13, 5, and 31 of them (Figure 6(a)). A heat map unveils the specific DEGs involved. In comparison to Cis-treated ovaries, cell proliferation-related DEGs, including Ki, Prrx2, Sfrp4, and Megfl0, and the antioxidative gene H2-Q10, were significantly upregulated in ovaries of the Cis+M+DQ group (Figure 6(f)). Tnfrsf9 and another proliferation activator of endothelial cells, Tnfrsf12a, were modulated upwards following single metformin treatment (Figure 6(f)). Strikingly, senolytic DQ treatment downregulated the inflammatory-related gene Cebpβ and the fibrosis-related gene Loxl4 (Figure 6(f)). Moreover, DEGs between Cis-treated ovaries and different senotherapeutic groups were enriched in a series of gene ontology (GO) terms related to multiple cellular processes (Figures 6(c)–6(e)). Compared to the Cis group, senolytic DQ co-administration led to a significant enhancement of steroid hormone and lipid biosynthesis, and the genes associated with the cell-development process were also enriched after DQ treatment (Figure 6(d)). In addition, combination senotherapies (Cis+M+DQ group) facilitated cell differentiation, cell development, and cell signal communication (Figure 6(e)). The above comparative analysis of cisplatin and senotherapeutic DEGs in the ovaries supports the promotion effects of senotherapies on steroid hormone, cell proliferation, differentiation, and cell communication pathways, all of which are pathways that may contribute to fertility preservation in female cancer survivors.

4. Discussion

To date, chemotherapy is still the prevalent treatment for tumors and, among the available drugs, cisplatin is widely used in clinical practice. Cisplatin-induced ovarian damage primarily manifests as a loss of the healthy ovarian reserve and increased atretic follicles [5]. Currently, no effective medications are available to prevent such ovarian damage.

The underlying mechanisms of cisplatin-induced ovarian dysfunction involve irreversible DNA damage, over-loaded oxidative stress, and uncontrolled apoptosis activation [5–7]. Recent evidence suggests that cisplatin not only leads to senescence of tumor cells but also affects peripheral non-tumor renal and neuronal tissues [23, 24]. However, it remains unclear whether SNCs exist in cisplatin-exposed normal ovarian tissues. In our present study, we confirmed that cisplatin administration could induce ovarian cell senescence both in vitro and in vivo, and we further discovered that metformin and DQ treatment may improve cisplatin-injured ovarian function by removing senescent ovarian cells and reducing DNA damage.

DQ, a well-known senolytic, has shown a broad-spectrum senescent cell-killing effect. Emerging evidence suggests that senolytics targeting cell senescence may blaze a new trail for age-related diseases [13]. The cocktail of DQ has been reported to improve murine cardiac function and exercise capacity, thus delaying murine aging [30]. Metformin has been a widely used oral hypoglycemic medication for type 2 diabetes mellitus for five decades, and its nonglycemic benefits, including a prominent SASP elimination efficiency, have been consecutively identified in various in vitro and in vivo models [18, 19, 31]. In our present study, we confirmed the distinct senescence-removal effect and ovarian-preservation efficiency of metformin and/or DQ. Consistent with previous studies, metformin displayed a better SASP-inhibition ability, since western blot analysis showed a remarkable accumulation of SASP in cell lysates following metformin treatment. Importantly, dramatic downregulation of the senescent marker P21 WAF1/CIP1 was observed in Cis+DQ and Cis+M GCs, indicating a considerable senolytic capacity of DQ and metformin in the ovarian setting. Notably, in addition to SASP prohibition, metformin benefits ovaries by promoting Sirt1 expression, preventing ovarian oxidative damage, reducing degenerating follicles and interstitial cells, and improving ovarian angiogenesis [32, 33]. Understandably, metformin plus DQ combination treatment produced a synergistic effect in cisplatin-exposed ovarian protection considering the complementary anti-aging targets of these 3 drugs. These data also supported the notion that the cell senescence burden is negatively correlated with ovarian health reserve and may probably work as a useful marker to predict female fecundity after cisplatin chemotherapy.

A hallmark of senescent cells is the secretion of SASPs or the senescence messaging secretome [10]. SASPs serve as mediators and spread senescence or senescence phenotypes in autocrine and paracrine fashions, thus interfere with cell viability and induce apoptosis [34]. We first set out to determine the SASPs secreted by senescent GCs following cisplatin exposure in vitro. A mouse cytokine array indicated significant elevation of inflammatory cytokine secretion in the cisplatin-exposed group; in particular, representative...
SASP IL-6 tripled over that in the NC group after cisplatin administration. The senomorphic agent metformin prominently alleviated SASP secretion after cisplatin exposure, since IL-6 and IL-1β were remarkably accumulated in cell lysates of Cis+M GCs compared to Cis-treated GCs. DQ showed a less profound influence on senescent granular SASP secretion, and IL-1β quantitative analysis showed an upwards tendency in cell lysates but reached no statistical significance compared to the Cis group. In addition, an in vivo animal experiment further demonstrated that cisplatin-induced ovarian cell senescence was not confined to GCs, since SA-β-gal staining and P16ink4a, and P21WAF1/Cip1-positive staining were also present in ovarian theca-interstitial cells. However, to determine whether theca-interstitial cell senescence is attributed to direct cisplatin damage or influenced by the SASP paracrine activity of GCs, further investigation is required.

In addition to the senescence removal of metformin and DQ discovered in our study, we surprisingly uncovered a metformin and/or DQ anti-DNA damage capacity in cisplatin-damaged ovaries. The antioxidative proteins SOD2 and HO-1 were significantly upregulated in the ovaries of the senotherapeutic groups, while subtle TUNEL staining and obvious EdU IF were noted in Cis+M, Cis+DQ, and Cis+M+DQ follicle units. Quantitative analysis of PCNA protein expression also revealed marked elevation after single or combined application of senotherapeutics. Transcriptome sequencing analysis further proved that the proportions of cisplatin-modulated genes could be rescued with single or combined senotherapies; of note, combination treatment rescued the largest proportion of genes. As expected, the heat map showed that cell proliferation-associated genes, such as Ki, Prrx2, Sfrp4, and Megfl0; and the antioxidative gene H2-Q10 were remarkably enhanced in Cis+M+DQ ovaries. Except for senolytic efficiency, DQ probably exerted anti-inflammatory and antifibrosis effects since related genes were upregulated following DQ co-administration. GO enrichment analysis revealed that the biological processes enhanced after combination treatment were positive regulation of cell development, cell differentiation, and cell communication. Taken together, these data verified that senotherapies not only fight against cell senescence but also participate in multiple cellular processes and promote ovarian cell survival after chemotherapy in multiple ways.

Several limitations of our study should be declared here. First, we did not comprehensively detect all senescence-related markers in mouse ovarian tissue and GCs. Future studies are required to cover more SASPs and senescence markers. Second, we did not confirm our discovery in clinical samples, and associated experiments will be undertaken in the near future.

In summary, our present study demonstrated that cellular senescence is involved in cisplatin-induced ovarian dysfunction and clearance of senescent cells by independent or combined administration of metformin and/or DQ ameliorated cisplatin-induced ovarian dysfunction in mice. Metformin plus DQ therapy further attenuated cisplatin-induced DNA damage. Transcriptomic sequencing revealed that the proportions of cisplatin-altered genes were rescued and biological processes, such as cell differentiation, cell development, and cell signal communication, were significantly upregulated after combination treatment. Additionally, we discovered synergistic effects of metformin and DQ in cisplatin-induced ovarian failure. Our work provides a previously unreported mechanism for cisplatin-induced ovarian damage and reveals that senotherapies targeting senescent cells could partially reverse this damage and may represent a promising therapeutic avenue to prevent chemotherapy-induced fertility loss.

5. Conclusion

In conclusion, our work demonstrates that senotherapies might prevent cisplatin-induced ovarian injury by removing senescent cells and reducing DNA damage, which may serve as a promising therapeutic avenue to prevent chemotherapy-induced ovarian damage.

Abbreviations

DQ: Dasatinib and quercetin
SASP: Senescence-associated secretory phenotype
DDR: DNA damage response
β-gal: β-Galactoside
SNCs: Senescent cells
GCs: Granulosa cells
rpms: Revolutions per minute
SDS-PAGE: Sodium dodecyl sulphate-polyacrylamide gel electrophoresis
TBST: Tris-Buffered Saline and Tween 20
NF-κB: Nuclear factor kappa-light-chain-enhancer of activated B-cells
IL: Interleukin
PCNA: Proliferating cell nuclear antigen
HO: Heme oxygenase
SOD: Superoxide dismutase
IHC: Immunohistochemistry
IF: Immunofluorescence
8-OHdG: 8-hydroxy-2-deoxyguanosine
ZP3: Zona pellucida protein 3
TUNEL: Terminal deoxynucleotidyl transferase dUDP nick end labeling
EdU: 5-Ethynyl-2'-deoxyuridine
HE: Hematoxylin and eosin
AMH: Anti-Müllerian hormone
E2: Estradiol
G-CSF: Granulocyte colony-stimulating factor
IOD: Integral optical density
DEG: Differentially expressed genes
GO: Gene ontology.

Data Availability

The data that support the findings of this study are available from the corresponding author upon reasonable request.
Disclosure

The funders had no role in the study design, data collection, analysis, and interpretation of the data or in writing the manuscript.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors’ Contributions

Dingfu Du, Jinjin Zhang, and Shixuan Wang designed the experiments. Dingfu Du and Xianan Tang accomplished the molecular-biological experiments in vitro and in vivo. Yueyue Gao, Runhua Chen, and Huan Lu took the responsibility of animal drug administration and data analysis. Qian Chen, Jingyi Wen, Tong Wu, and Yan Zhang revised the manuscript. Dingfu Du and Xianan Tang contributed equally to this work.

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