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Endosulfan induced the arrest of cell cycle through
inhibiting the signal pathway mediated by PKC-α and
damaging the cytoskeleton in spermatagonial cell of mice in
vitro

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Abstract
The previous studies have showed that endosulfan has adverse effects on male
reproductive system and the oxidative stress induced by endosulfan is related to its
toxicity. However, the molecular mechanism of endosulfan reproductive toxicity is
still unknown. To investigate its mechanism of toxicity, the GC-1spg cells were
exposed to 0, 6, 12 and 24μg/mL endosulfan for 24h, respectively. Results showed
that endosulfan resulted in a dose-dependent reduction in cell viability and increases
in LDH release, apoptosis rate, the malondialdehyde (MDA) level, the reactive oxygen species (ROS) production and DNA damage degree including the percentage of tail DNA, tail length, tail moment (TM) and Olive tail moment (OTM). Endosulfan induced the arrests of both S and G2/M phase and proliferation inhibition. The expression of PKC-α, CDK2, Cyclin E, RAF-1, MEK1/2, p-MEK1/2, ERK1/2 and p-ERK1/2 in GC-1spg cells declined remarkably after treatment with endosulfan. Endosulfan also damaged the microfilament, microtubule and cell nucleus, and blocked the mitosis process. The results suggested that endosulfan could induce the cell cycle arrest and proliferation inhibition by inhibiting the protein expression of cellular signaling pathway mediated by PKC-α because of DNA damage resulted from oxidative stress; meanwhile, endosulfan could also lead to mitotic arrest through directly damaging the cytoskeleton and cell nucleus resulted from oxidative stress; which therefore induced cytotoxicity of GC-1spg cells.

**Key Words:** Spermatagonial cells, cytotoxicity, cell cycle arrest, cytoskeleton, cellular signaling pathway

**Introduction:**

Endosulfan(6,7,8,9,10-hexachloro-1,5,5a,6,9,9a-hexahydro-6,9-methano-2,4,3-benzodioxathiepin-3-oxide,C9H6Cl6O3S), which is an organochlorine insecticide, is still extensively used in some developing countries despite it is defined as a persistent organic pollutant (POP) by the Stockholm Convention on POPs in April 2011.1, 2
Contamination of the air, water and soil environments by endosulfan occurs generally as a result of runoff from the agricultural fields and/or discharge during its manufacture. Endosulfan exists in water with a half-life of more than 14 d and persists longer in soil with a half-life of approximately 60–800 d. Because endosulfan is highly lipophilic, it bioaccumulates and biomagnifies along the alimentary chain. It has been detected in different matrices (water bodies: 1.7, soil: 0.3–34.9, fruits: 0.36–212.3, milk: 0.4–56.2, butter: 0.6–13.4; Laddu: 676.0, human blood in Kasargod district of Kerala, India: 0.69–176.2 μg/mL). The data obtained from the global monitoring network programme for POPs also showed that endosulfan was still abundant in the environment and that it was used a lot in some countries.

Endosulfan has been implicated in genotoxicity, neurotoxicity, hepatotoxicity, immunotoxicity, reproductive toxicity and teratogenicity on exposed organisms. With the increasingly serious environmental pollution and reduction of biodiversity, there is growing concern that chemicals, whether natural or human-made, may be causing damage to reproductive system of wildlife and human. Moreover, the male reproductive system has raised major concerns from both the research community and the public. The previous studies showed that endosulfan caused reductions in the motility, viability and daily sperm production (DSP), damaged the integrity of sperm chromatin, decreased the level of serum testosterone and testicular weight, inhibited the spermatogenesis and increased sperm abnormalities. Moreover, accumulating evidence suggested that oxidative stress induced by...
endosulfan was related to its toxicity and endosulfan could indirectly cause a decline in reproductive function by damaging the structure of mitochondria, resulting in energy metabolism dysfunction. Study also reported that endosulfan upregulates AP-1 binding and ARE-mediated transcription via ERK1/2 and p38 activation in HepG2 cells. However, the molecular signaling pathways of endosulfan-induced reproductive toxicity on spermatagonial cells have maintained unknown. Besides, the data which associate cell-cycle signaling pathways with endosulfan have not been reported so far. Therefore, the present experiment was designed to research the effects of endosulfan on the cell cycle and related cellular signaling pathways mediated by PKC-α in spermatagonial cell in vitro, so as to explain the toxic mechanism of endosulfan.

Materials and Methods

Cell Culture and Treatment

The spermatagonial cell GC-1spg line was purchased from American Type Culture Collection (ATCC® Number: CRL-2196™). The cells were maintained in Dulbecco’s Modified Eagle’s Medium (DMEM) (Gibco, USA) supplemented with 10% fetal bovine serum (Gibco, USA), 100 U/mL penicillin and 100μg/mL streptomycin, and cultured at 37°C in 5% CO₂ humidified environment. For experiments, the cells were seeded in 6-well plates (except MTT assay using 96-well plates) at a density of 1×10⁵ cells/mL and allowed to attach for 24 h, then treated with various concentrations of endosulfan (3, 6, 12, 24, 36, 48, 60 and 72μg/mL) for
another 24 h, respectively. Endosulfan (analytical standard, purity: 96%) was obtained from Jiangsu Kuaida Agrochemical Co., Ltd. (Nantong, China), and dissolved in DMSO and diluted with DMEM. Vehicle controls were supplied with an equivalent volume of DMEM which included 0.24 % DMSO. Each group had five replicate wells. All experiments were repeated at least three times.

**Determination of Cell Viability**

The cell viability was determined using 3-(4, 5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide, a yellow tetrazole (MTT) assay performed as previously described. The absorbance of formazan was measured at 492 nm by a microplate reader (Thermo Multiskan MK3, USA).

**Determination of LDH Release**

The integrity of the cell membrane, which is identified by the measurement of lactate dehydrogenase (LDH) activity in the extracellular medium, was measured using LDH Kit (Jiancheng, China) according to the manufacturer’s protocols. After exposure to different concentrations of endosulfan for 24 h, 100μl of cell culture media was used to analyze LDH activity, and the absorbance at 440 nm was determined using a UV-visible spectrophotometer (Beckman DU-640B, USA).

**Determination of Cell Apoptosis**

Apoptosis in GC-1spg cells was determined using the Annexin V-propidium iodide (PI) apoptosis detection kit (KeyGen, China). After centrifugation at 1000 rpm for 5 min, the cells were washed twice with PBS and suspended with 500 μL binding buffer, then incubated with 5 μL Annexin V-FITC and 5μL PI for 15 min in the dark at room
temperature and analyzed using a flow cytometer (Millipore, USA), and at least $1 \times 10^4$ cells were counted for each sample. The different cell populations were identified by the different labeling patterns in the Annexin V-PI analysis. For example, FITC negative and PI negative were designated as live cells in the lower left quadrant; FITC positive and PI negative as early apoptotic cells in the lower right quadrant; FITC positive and PI positive as late apoptotic cells in the upper right quadrant; and FITC negative and PI positive as cells fragments in the upper left quadrant. The sum of early apoptosis rate and late apoptosis rate was finally calculated as the total rate of apoptosis.

**Determination of Oxidative Damage**

The malondialdehyde (MDA) (an end product of lipid peroxidation) content was determined using commercially available kits (Jiancheng, China) according to the manufacturer’s instructions. After exposure to endosulfan for 24 h, the cells were washed once with ice-cold PBS, and lysed in ice-cold RIPA lysis buffer which contained 1 mM phenylmethylsulphonyl fluoride (PMSF) (Dingguo changsheng biotech CO.LTD, China) for 30 min. After centrifugation at 12,000 rpm, 4°C for 10 min, the supernatants were collected to measure the MDA content.

The oxidative stress is related to overproduction of reactive oxygen species (ROS). The generation of intracellular ROS was measured through fluorescence intensity of dichlorofluorescein (DCF) by flow cytometry using the 2, 7-dichlorofluorescein diacetate (DCFH-DA) (Beyotime, China). After exposure to endosulfan for 24 h, the
cells were incubated with medium which contained DCFH-DA at a concentration of 5 μM for 40 minutes at 37 °C in the dark. Then the cells were washed twice with cold PBS and resuspended in the PBS for further analysis. At last, fluorescent intensity of at least 1×10^4 cells for each sample were measured by flow cytometry (Becton-Dickison, USA).

**Determination of DNA Damage**

Comet assay, also known as single cell gel electrophoresis (SCGE), is a sensitive technique for determining DNA breakage in individual cells. The DNA damage induced by endosulfan was measured by Single cell gel electrophoresis kit (Biolab, China). After GC-1spg cells were collected, they were resuspended in PBS, 20μL of the cells suspension and 80μL of low melting agarose were mixed, and 80μL of the suspension was pipetted onto a comet-slide. Then the slides were placed in dark at 4°C for 10 min, and then placed in pre-chilled lysing solution at 4°C for 2h. The slides were removed from lysing solution, tapped on a paper towel and immersed in alkaline solution for 45 min in dark at room temperature. After washed twice for 5 min per time, the slides were electrophoresed at low voltage (300 mA, 25 V) for 30 min. Then the slides were removed from the electrophoresis unit and tapped at room temperature. Subsequently, the air-dried slides were stained with propidium iodide (PI) and assessed by a fluorescence microscope (Olympus IX81, Japan). To prevent additional DNA damage, all the steps were performed under dimmed light. The data were analyzed using CASP software based on 100 randomly selected cells per sample. The percentage of tail DNA, tail length, tail moment (TM) and Olive tail moment (OTM)
were selected as indicators of DNA damage.

**Determination of Cell Cycle**

The cell-cycle distributions were measured by the cell cycle detection kit (KeyGen, China). After centrifugation at 1000 rpm for 5 min, the cells were washed once with PBS and fixed in 70% ethanol (placed in -20°C for 24 h in advance) for 24 h. Then the cells were washed twice with PBS and treated with 100μL of RNase A at 37°C for 30 min. Finally, the cells were stained with 400μl of propidium iodide (PI) and incubated in dark for 30 min. At least 1×10⁴ cells for each sample were analyzed by flow cytometry (Beckman Coulter, USA).

**Assessment of Microfilament, Microtubule and the Cell Nucleus Damage**

After washed twice with PBS and fixed with 3.7% formaldehyde solution at room temperature for 10 minutes, the cells were washed twice again with PBS (containing 0.1% TritonX-100) for 5 minutes per time, and stained with 5 μg/mL Hoechst 33258 solution (Beyotime, China) for 20 minutes. Then the cells were washed twice with PBS (containing 0.1% TritonX-100) for 5 minutes per time and stained with 200 nM of Actin-Tracker Green (Enogene Biotech. Co., Ltd, China) in the dark for 30 minutes at 37°C. Subsequently, the cells were washed twice with PBS (containing 0.1% TritonX-100) for 5 minutes per time and stained with 250 nM of Tubulin-Tracker Red (Enogene Biotech. Co., Ltd, China) in the dark for 30 minutes at 37°C. Finally, the cells were visualized using a laser confocal microscope (Leica, Germany) and selected randomly for further observation.

**Assessment of Mitosis Process**
After exposed to 24 μg/mL endosulfan, the GC-1spg cells were observed using a real-time inverted phase contrast microscope (UltraVIEW VoX, USA) for 24h to determine whether there were abnormal mitosis phenomenon. Six visual fields were selected randomly and took pictures every 10 minutes.

**Determination of Cell Proliferation**

After centrifugation at 1200 rpm for 5 min, the GC-1spg cells were marked with 5μM of CFDASE probe diluent（KeyGen, China）at 37 ºC for 15 min. Then the cells were washed twice with PBS, seeded in 6-well plates at a density of 1×10^5 cells/mL and allowed to attach for 24 h. After the cells were treated with endosulfan for 24 h and centrifuged at 1200 rpm for 5 min, 500μL of PBS were added in to suspend the cells. Finally, the average fluorescence intensity of cells per sample was detected by flow cytometry (BDFACSaria, USA).

**Determination of Protein Expression in Cellular Signaling Pathway**

The protein levels of PKC-α, Cyclin E, CDK2, RAF-1, ERK1/2, p-ERK1/2, MEK1/2, and p-MEK1/2 in GC-1spg cells were determined by western blot analysis. Total protein of GC-1spg cells was extracted using a Protein Extraction Kit (KeyGen, China) and measured by the bicinchoninic acid (BCA) protein assay (Dingguo changsheng biotech CO.LTD, China). The equal amounts of lysate proteins (20μg) were loaded onto SDS-polyacrylamide gels (12% separation gels) and electrophoretically transferred to polyvinylidene fluoride (PVDF) membranes (Millipore, USA). After blocked with 5% nonfat milk in TBST(Tris-buffered saline containing 0.05%Tween-20) for 1 h at room temperature, the membrane was
incubated with β-actin[1:1000, rabbit antibodies, Cell Signaling Technology (CST), USA], PKC-α (1:2000, rabbit antibodies, Abcam, USA), Cyclin E[1:1000, rabbit antibodies, Cell Signaling Technology (CST), USA], CDK2 (1:500, rabbit antibodies, Abcam, USA), RAF-1[1:500, rabbit antibodies, Santa Cruz Biotechnology (SCBT), USA], ERK1/2[1:1000, rabbit antibodies, Cell Signaling Technology (CST), USA], p-ERK1/2[1:1000, rabbit antibodies, Cell Signaling Technology (CST), USA], MEK1/2 (1:1000, rabbit antibodies, Abcam, USA) and p-MEK1/2 (1:1000, rabbit antibodies, Abcam, USA) overnight at 4°C, respectively, washed three times with TBST for 10 min per time, and incubated with a horseradish peroxidase-conjugated anti-rabbit IgG secondary antibody [1:5000, Immunology Consultants Laboratory (ICL), USA] for 1 h at room temperature. After washed three times with TBST for 10 min per time, the antibody-bound proteins were detected with the ECL chemiluminescence reagent (Pierce, USA).

**Statistical Analysis**

All data were analyzed using SPSS (Statistical Package for the Social Sciences software, version 17.0 for Windows). One-way analysis of variance (ANOVA) was used to test the significant difference among all groups, followed by least significant difference (LSD) test to detect significant difference between two groups. All values were expressed as mean ± standard error (S.E.), and the results were considered significant at P < 0.05. Based on the MTT results, the values from LC10 to LC90 were calculated by probit analysis using SPSS software, and the confidence intervals were 95% according to the method used by Nath and Kannan.22, 23
Results

Cytotoxicity

Cell viability gradually decreased with the increase of endosulfan level in a dose-dependent manner compared with the control group, and in endosulfan treated groups (>3μg/mL) were remarkably lower than that of control group and significantly different from each other (Figure 1A). In addition, the LC50 was 32.829μg/mL (Table 1). Based on the MTT results, endosulfan concentrations were chosen at 6, 12 and 24μg/mL for the next experiments. Moreover, the MTT result was in accordance with the increased cytomembrane damage measured by LDH activity. The LDH activity increased apparently in 12 and 24μg/mL endosulfan treated group compared to control group (Figure 1B). The total apoptotic rates in GC-1spg cells increased with the increase of endosulfan level (Figure 4A), and significantly enhanced in 12 and 24μg/mL endosulfan treated group when compared to control group (Figure 1C). The total apoptosis levels of GC-1spg cells were 1.82%, 5.25%, 8.04% and 28.71% , respectively, after exposure to 0, 6, 12 and 24μg/mL endosulfan.

Oxidative Stress and DNA Damage

Oxidative damage was evaluated by measuring the intracellular MDA level and the generation of ROS. The MDA level gradually increased with the elevation of endosulfan level, and significantly enhanced in 12 and 24μg/mL endosulfan treated group when compared to control group (Figure 1D). The intracellular ROS levels of all endosulfan-treated groups were remarkably increased in a dose dependent manner.
when compared to control group. In 24μg/mL treated group, the fluorescence intensity was about 3-fold much higher than that of control (Figure 1E). The degree of DNA damage involving the length of tail, the percentage of tail DNA, tail moment (TM) and Olive tail moment (OTM) had no significant difference in 6 and 12μg/mL endosulfan treated group compared with the control group. While in 24μg/mL endosulfan treated group, the degree of DNA damage significantly elevated (Table 2).

**Cell Cycle Arrest and Proliferation Inhibition**

Results showed that the percentage of GC-1spg cells in G0/G1 phase declined in a dose-dependent manner, whereas the percentages of GC-1spg cells in S and G2/M phase increased when compared with the control group (Figure 5C). The percentages of GC-1spg cells had a significant increase in S phase in 24μg/mL endosulfan treated group and G2/M phase in 6, 12, 24μg/mL endosulfan treated group when compared to control group (Figure 2B). Results also showed that the average fluorescence intensity of cells in groups exposed to 12μg/mL and 24μg/mL endosulfan increased significantly in comparison with the control group (Figure 2A), which indicated that cell proliferation in this two groups were significantly inhibited.

**Changes of the Microfilament, Microtubule, Nucleus and Mitosis process**

Results indicated that compared with the control group, the fluorescence intensity of microfilaments and microtubules weakened gradually in cells after treatment with
various concentrations of endosulfan, which showed that the number of microfilaments and microtubules reduced gradually with the increase of endosulfan concentration. Consistently, the maldistribution phenomena of microfilaments and microtubules were more obvious with the increase of endosulfan concentration, even there were accumulations of microfilaments and microtubules, and extinctions of microfilaments and microtubules(Figure 3A and 3B). Moreover, compared with the control group, the size and morphology of nuclei was obviously different in cells after exposure to endosulfan. The cells in control group had round and homogeneously stained nuclei, whereas the phenomenon of nucleus pycnosis was frequently observed in 12 and 24μg/mL endosulfan treated group(Figure 3C).

Results also indicated that, after treatment with 24μg/mL endosulfan, some cells failed to complete mitosis process normally and died finally (Figure 3E).

Changes of Protein Expression in Cellular Signaling Pathway Mediated by PKC-α

Results showed that the expression of PKC-α, CDK2, Cyclin E, RAF-1, MEK1/2, p-MEK1/2, ERK1/2 and p-ERK1/2 in GC-1spg cells declined gradually with the increase of endosulfan concentration (Figure 4A). The expression of CDK2 in GC-1spg cells was obviously inhibited in 6, 12 and 24μg/mL endosulfan treated group (Figure 4B). The expression of PKC-α, Cyclin E, RAF-1, MEK1/2, p-MEK1/2, ERK1/2 and p-ERK1/2 in GC-1spg cells decreased remarkably in 12 and 24μg/mL
endosulfan treated group (Figure 4B).

**Discussions**

The purpose of this study was to investigate the molecular mechanism by which endosulfan induced the male reproductive toxicity. Health of spermatagonial cells may affect the quality of gametes. So we studied the molecular mechanism *in vitro* with the GC-1spg cells.

First of all, to demonstrate the potential cytotoxicity of endosulfan on GC-1spg cells, the cell viability, LDH activity and apoptosis rates were measured. Results showed that endosulfan resulted in reduction in cell viability and increases in LDH activity and the apoptotic rates in GC-1spg cells. To further confirm whether the oxidative stress was one of mechanisms by which endosulfan induced male reproductive toxicity, we measured the levels of MDA (a biomarker of oxidative stress) and ROS which mediated oxidative stress. The result showed that endosulfan could raise the MDA and ROS level, suggesting that endosulfan induced oxidative stress. Takhshid also showed that endosulfan could increased testis MDA, while supplementation of vitamin C and vitamin E to endosulfan-treated rats could reduce the toxic effect of endosulfan on lipid peroxidation in the testis. Recognizing that the reactive oxygen could directly induce DNA base oxidation, deamination and indirectly lead to base alkylation via lipid peroxidation (LPO), we sought to examine whether the DNA was damaged in GC-1spg cells treated with endosulfan. We did find the higher
concentration of endosulfan produced more severe damage on DNA of GC-1spg cells, which was similar to those of Tao, who showed that the DNA damage in both tissues had excellent correlation with endosulfan concentration.\textsuperscript{26}

Studies had shown that, in response to transient or low levels of DNA damage, cells had a reversible cell cycle arrest whereas a sustained cell cycle arrest followed prolonged or high levels of DNA damage.\textsuperscript{27} For DNA damage was closely related to cell cycle arrest, especially G1 phase, S phase and G2 phase, and cell cycle was regulated by Cell-cycle checkpoints.\textsuperscript{28} The function of these checkpoints was to test the damaged or abnormally structured DNA, and to control cell-cycle progression with DNA repair.\textsuperscript{28} To get a closer insight into the toxic mechanisms of endosulfan on GC-1spg cells, cell cycle distribution was measured. We found that endosulfan blocked the GC-1spg cells at both S and G2/M two phases. Several evidences indicated that the main components of cell cycle control machinery were the Cyclins and CDKs, which constituted active kinase complexes during specific phases of the cell cycle.\textsuperscript{29} Among them, Cyclin E/CDK2, had an indispensable function in the transition from G1 into S phase.\textsuperscript{15} Besides, there were protein kinases C (PKC) as key regulators of critical cell cycle phases, including G1/S transition and G2/M progression.\textsuperscript{30-33} Among them, PKC-\alpha was known for its involvement in G1/S transition.\textsuperscript{34} So the expression of Cyclin E, CDK2 and PKC-\alpha were measured. Results showed that endosulfan induced down-regulation of Cyclin E, CDK2 and PKC-\alpha expression in GC-1spg cells. The results above suggested that endosulfan blocked the
cell cycle at S phase through the down-regulation of Cyclin E/CDK2 and PKC-α expression, which was similar to those of Guo, who showed that P.linteus(PL) and Ganoderma lucidum (GL) suppressed the growth of Bel-7404 cells through S phase cell cycle arrest, mediated by the inhibition of Cyclin E/CDK2 activities.

Microfilament and microtubule were the main components of the cytoskeleton. The cytoskeleton played an important role in the process of mitosis and M phase was the mitotic phase. To examine the mechanism of cell cycle arrest at G2/M phase, the changes in microfilament, microtubule, the nucleus and the process of mitosis in cells after exposure to endosulfan were observed. Results showed that the number of microfilaments and microtubules significantly reduced, the maldistribution phenomenon of microfilaments and microtubules obviously increased, and the nucleus pycnosis was more frequently observed with the increase of endosulfan concentration. Moreover, some cells in 24μg/mL endosulfan treated group failed to complete mitosis process normally and died finally, which was similar to those of William. They showed that treatment with the ILK Inhibitor QLT-0267 resulted in a significant G2/M arrest as well as aberrant cytokinesis and mitotic spindle disorganization. Hence, we infered that endosulfan might inhibit the process of mitosis by causing the damages of microfilament, microtubule and nuclear. Blocking cell cycle progression may result in proliferation inhibition. So the proliferation were assessed. Results showed that endosulfan inhibited the proliferation of GC-1spg cells. But the result of Tellez-Banuelos showed that low concentration of endosulfan
increased cellular proliferation in spleen cells, which was different with our results. This may be because the doses of endosulfan and cell lines used in the experiments were different. To investigate the possible mechanism of cell proliferation inhibition induced by endosulfan, the related signaling pathways of cell cycle were further studied. The mitogen-activated protein kinases (MAPKs) were serine/threonine kinases which played a vital role in controlling proliferation. Among them, the ERK family including ERK1 and ERK2 were proven to be implicated. In proliferation, the PKC-α–RAF-1–MEK1/2–ERK1/2 pathway was involved. So the expression of ERK1/2, p-ERK1/2, MEK1/2, p-MEK1/2 and RAF-1 were measured. Our results showed that endosulfan induced the down-regulation of RAF-1, MEK1/2, p-MEK1/2, ERK1/2 and p-ERK1/2 expression, which was similar to those of Levinthal. The study of Levinthal using murine neuronal cells had shown that glutamate-induced oxidative stress specifically inhibited the phosphatase activity by the down-regulation of ERK1/2 expression (PP2A, MKPs). However, the previous study had shown that endosulfan could activate ERK1/2 via PKC-RAF-MEK1/2-dependent and PKC-MEK1/2-dependent pathways in human HaCaT cells. Our results suggested that endosulfan inhibited ERK1/2 via PKC-α–RAF-1–MEK1/2-dependent pathway in GC-1spg cells. So we deduced that different kinds of cells might have different responses to various concentration of endosulfan.

The previous findings indicated that Cyclins and Cyclin dependent kinase (CDK)
were downstream mediators of the growth-regulatory activity of Irf-1,\textsuperscript{42-44} and blocking ERK1/2 pathway inhibited Irf-1-mediated proliferation.\textsuperscript{45} The results indicated that inhibition of ERK1/2 pathway might cause the down-regulation of Cyclins and CDK expression, which was the same with our results. Hence, the molecular mechanism may be that endosulfan could induce the cell cycle arrest and proliferation inhibition by inhibiting the protein expression of PKC-α–RAF-1–MEK1/2-ERK1/2 pathway because of DNA damage resulted from oxidative stress and down-regulating the protein expression of Cyclin E and CDK2; meanwhile, endosulfan could also lead to mitotic arrest through directly damaging the cytoskeleton and cell nucleus resulted from oxidative stress; which therefore induced cytotoxicity of GC-1spg cells.

In summary, endosulfan could cause the cytotoxicity of GC-1spg cells by the following two pathways (1) endosulfan could induce cell cycle arrest at S phase and proliferation inhibition by inhibiting the protein expression of cellular signaling pathway mediated by PKC-α; (2) endosulfan could lead to cell cycle arrest at G2/M phase through directly damaging the microfilament, microtubule and the cell nucleus. The finding provides new evidence to explain the endosulfan-induced male reproductive toxicity and reveals new potential downstream pathway. It is important to understand the mechanism of reproductive toxicity from endosulfan and prevent from it. However, additional work is required to explore the precise mechanism responsible for endosulfan-induced male reproductive toxicity \textit{in vivo}, such as
signaling pathway related to meiosis. Moreover, management should be more strict in additional risks of endosulfan exposure in certain subpopulations, for example, people who have the disease of reproductive system.

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References


Figure 1.

Cytotoxicity and Oxidative Damage of GC-1spg Cells Induced by Endosulfan

(Means±S.E.) . (A) Cell viability of GC-1spg cells treated with various concentrations of endosulfan for 24 h was measured by MTT assay. Values with entirely different superscripts were significantly different (P<0.05) (B) LDH leakage of GC-1spg cells exposed to various concentrations of endosulfan for 24 h. (C) The sum of early apoptosis rate and late apoptosis rate was finally calculated as the apoptosis rate. (D) The MDA level of GC-1spg cells exposed to various concentrations of endosulfan for 24 h. (E) The ROS level of GC-1spg cells exposed to various concentrations of endosulfan for 24 h.
* indicated significant difference compared to control group ($P < 0.05$)

Figure 2.

**Proliferation Inhibition and Cell Cycle Arrest of GC-1spg Cells Induced by Endosulfan** (Means±S.E.) . (A) After exposure to various concentrations of endosulfan for 24 h, the proliferation of GC-1spg cells were determined using flow cytometry. The increase of average fluorescence intensity in cells indicated cellular proliferation were inhibited. (B) After exposure to different concentrations of endosulfan for 24h, the cell cycle distribution of GC-1spg cells was measured by flow cytometry.

* indicated significant difference compared to control group ($P < 0.05$)
Figure 3.

Changes of the Microfilament, Microtubule, Cell Nucleus and the Mitosis Process Induced by Endosulfan. After exposure to various concentrations of endosulfan for 24 h, the changes in microfilament, microtubule, cell nucleus and the mitosis process were visualized using a laser confocal microscope. (A)
microfilament was stained with Actin-Tracker Green. The images from a5 to a8 were the images magnified double from a1 to a4, respectively. (B) The microtubule was stained with Tubulin-Tracker Red. The images from b5 to b8 were the images magnified double from b1 to b4, respectively. (C) The cell nucleus was stained with Hoechst 33258 solution. The images from c5 to c8 were the images magnified double from c1 to c4, respectively. (D) The images were the composite images of microfilament, microtubule and cell nucleus. The images from d5 to d8 were the images magnified double from d1 to d4, respectively. (E) The cells were incubated with 24 μg/mL endosulfan and observed for 24h by a real-time inverted phase contrast microscope. Six fields were selected randomly and taken pictures every 10 minutes. In the images, the white arrow pointed to the normal mitosis process and the black arrow pointed to the abnormal mitosis process.
Figure 4.

Effect of Endosulfan on the Cellular Signaling Pathway Mediated by PKC-α

(Mean±S.E.). (A) Effect of endosulfan on the protein expression of PKC-α, CDK2, Cyclin E, RAF-1, MEK1/2, p-MEK1/2, ERK1/2 and p-ERK1/2. β-actin was used as an internal control to monitor for equal loading. (B) Relative densitometric analysis of the proteins bands was carried out and presented.

* indicated significant difference compared to control group (P < 0.05)
Figure 5.

The Images of Apoptosis, DNA Damage, Cell Cycle Arrest and Proliferation Inhibition of GC-1spg Cells after Exposure to Endosulfan for 24h (A) a-d:

Control group(a), 6μg/mL endosulfan group(b), 12μg/mL endosulfan group (c), 24μg/mL endosulfan group (d). Apoptotic populations of cells double stained with PI- and FITC-labeled Annexin V were described by flow cytometry. (B) a-d: Control group 200×(a), 6μg/mL endosulfan group 200×(b), 12μg/mL endosulfan group 200×(c),
24μg/mL endosulfan group 200×(d). DNA damage of GC-1spg cells treated with various concentrations of endosulfan for 24 h was measured by Comet assay. More severe DNA damage was reflected by larger area of the comet tail. (C) a-d: Control group (a), 6μg/mL endosulfan group (b), 12μg/mL endosulfan group (c), 24μg/mL endosulfan group (d). The red area, green area and yellow area represented the G0/G1, S and G2/M phase, respectively. (D) a-d: Control group(a), 6μg/mL endosulfan group(b), 12μg/mL endosulfan group (c), 24μg/mL endosulfan group (d). The abscissa value of peak was the average fluorescence intensity of cells.

Table 1.

**Cytotoxicity of GC-1spg Cells Induced by Endosulfan**

<table>
<thead>
<tr>
<th>Concentration (μg/mL)</th>
<th>Cell viability (% of control)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>100</td>
</tr>
<tr>
<td>4.397</td>
<td>90</td>
</tr>
<tr>
<td>11.505</td>
<td>80</td>
</tr>
<tr>
<td>18.613</td>
<td>70</td>
</tr>
<tr>
<td>25.721</td>
<td>60</td>
</tr>
<tr>
<td>32.829</td>
<td>50</td>
</tr>
<tr>
<td>39.937</td>
<td>40</td>
</tr>
<tr>
<td>47.045</td>
<td>30</td>
</tr>
<tr>
<td>54.153</td>
<td>20</td>
</tr>
<tr>
<td>61.261</td>
<td>10</td>
</tr>
</tbody>
</table>
Table 2.

DNA Damage of GC-1spg Cells Induced by Endosulfan

<table>
<thead>
<tr>
<th>Concentration (μg/mL)</th>
<th>Tail length</th>
<th>Tail DNA (%)</th>
<th>Tail movement</th>
<th>Olive tail movement</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>18.30±0.90</td>
<td>8.20±0.47</td>
<td>1.59±0.22</td>
<td>2.19±0.23</td>
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<tr>
<td>6</td>
<td>22.47±0.81</td>
<td>9.24±0.46</td>
<td>2.52±0.20</td>
<td>2.64±0.19</td>
</tr>
<tr>
<td>12</td>
<td>22.53±1.18</td>
<td>9.25±0.50</td>
<td>3.56±0.40</td>
<td>3.34±0.29</td>
</tr>
<tr>
<td>24</td>
<td>69.65±2.04*</td>
<td>54.64±0.81*</td>
<td>58.85±2.04*</td>
<td>39.94±1.07*</td>
</tr>
</tbody>
</table>

* indicated significant difference compared to control group (P < 0.05)