Abstract. Dioscin, an extract from traditional Chinese herbal plants, displays various biological and pharmacological effects on tumors, including inhibition of cell proliferation and induction of DNA damage. However, the effects of dioscin on oral squamous cell carcinoma (OSCC) cells are not completely understood. The present study aimed to evaluate the impact of dioscin on OSCC cell proliferation. Cell Counting Kit-8 and 5-ethyl-2'-deoxyuridine incorporation assays were performed to assess cell proliferation. Flow cytometry was conducted to detect alterations in the cell cycle and cell apoptosis. Western blotting and coimmunoprecipitation were performed to determine protein expression levels. In SCC15 cells, dioscin treatment significantly induced cell cycle arrest, increased apoptosis and inhibited proliferation compared with the control group. Mechanistically, the tumor suppressor protein Ras association domain-containing protein 1A (RASSF1A) was activated and oncoprotein yes-associated protein (YAP) was phosphorylated by dioscin. Furthermore, YAP overexpression and knockdown reduced and enhanced the inhibitory effects of dioscin on SCC15 cells, respectively. In summary, the results demonstrated that, compared with the control group, dioscin upregulated RASSF1A expression in OSCC cells, which resulted in YAP phosphorylation, thus weakening its transcriptional coactivation function, enhancing cell cycle arrest and apoptosis, and inhibiting cell proliferation. The present study indicated that dioscin may serve as a therapeutic agent for OSCC.

Introduction

Oral squamous cell carcinoma (OSCC) is a common oral disease. According to the most recent GLOBOCAN estimate, in Europe between 2012 and 2015, there was an overall increasing incidence of oral cancer and of oral cancer-associated mortality (1,2). At present, OSCC treatment regimens include a combination of surgery, radiotherapy and chemotherapy (3,4). Despite the rapid development of modern medicine, the cure rate for OSCC is still poor, displaying an overall 5-year survival rate of 60% (5). Therefore, identifying additional safe and effective targeted drugs with low cytotoxicity is important.

It has been reported that natural plant products display tumor-inhibitory activities when used as chemopreventive or therapeutic agents against human cancer cells (6,7). Dioscin, which is extracted from the traditional Chinese medicinal herb Dioscorea nipponica, has been reported to display various biological effects, including renal ischemia/reperfusion injury-alleviating, anti-inflammatory and antiallergic effects (8,9). Additionally, in a nephrotoxicity and cardiotoxicity model, dioscin displayed protective effects via regulating oxidative stress (10-12). Meanwhile, dioscin has been reported to display potent effects against different types of cancer. For example, dioscin markedly inhibited hepatocellular carcinoma proliferation and migration, but induced apoptosis, autophagy and DNA damage (13). Si et al (14) reported that dioscin suppressed laryngeal cancer cell proliferation via inducing cell cycle arrest and inhibiting tumor invasion. Furthermore, dioscin inhibited human lung cancer cell proliferation (15). However, to the best of our knowledge, the effect of dioscin on OSCC has not been previously reported.

The Hippo signaling pathway is highly conserved across species. In mammals, the macrophage stimulating...
**Materials and methods**

**Cell line and culture.** The tongue squamous cell line SCC15 was provided by Professor Shaohua Liu (Qilu Hospital of Shandong University, Jinan, China). SCC15 cells were cultured in DMEM (HyClone; Cytiva) supplemented with 10% FBS (Gibco; Thermo Fisher Scientific, Inc.) at 37˚C with 5% CO₂.

**Reagents and antibodies.** Dioscin (purity ≥98%) was purchased from Beijing Solarbio Science & Technology Co., Ltd. Cells were treated with different concentrations of dioscin (0, 0.5, 1 or 2 µM) for 24 or 48 h, or with 1 µM dioscin for subsequent analysis via western blotting, according to the manufacturer's instructions (Active Motif, Inc.). Following centrifugation at 4˚C for 5 min at 137 x g, cells were suspended in 100 µl freshly prepared complete RIPA buffer and incubated for 30 min on a rotating platform at 4˚C. Whole cell extract was combined with 5 µg IgG (cat. no. 3900) or anti-MST2 (cat. no. 3952) antibodies (both from Cell Signaling Technology, Inc.) in a final volume of 500 µl and incubated overnight at 4˚C on a rotating platform. Subsequently, 25 µl protein G magnetic beads were added to the tube and incubated overnight at 4˚C on a rotating platform. The protein G magnetic beads were washed carefully and the protein samples were obtained. The samples were resuspended in loading buffer (cat. no. P1040; Beijing Solarbio Science & Technology Co., Ltd.) and boiled for subsequent analysis via western blotting, according to the aforementioned protocol.

**Cell proliferation.** Cell proliferation was assessed by performing Cell Counting Kit-8 (CCK-8) assays (Dojindo Molecular Technologies, Inc.) according to the manufacturer's protocol. SCC15 cells (3x10⁴ cells/well) were seeded into a 48-well plate and cultured for 24 h, lysates were prepared according to the manufacturer's instructions (Active Motif, Inc.). Following centrifugation at 4˚C for 5 min at 137 x g, cells were suspended in 100 µl freshly prepared complete RIPA buffer and incubated for 30 min on a rotating platform at 4˚C. Whole cell extract was combined with 5 µg IgG (cat. no. 3900) or anti-MST2 (cat. no. 3952) antibodies (both from Cell Signaling Technology, Inc.) in a final volume of 500 µl and incubated overnight at 4˚C on a rotating platform. Subsequently, 25 µl protein G magnetic beads were added to the tube and incubated overnight at 4˚C on a rotating platform. The protein G magnetic beads were washed carefully and the protein samples were obtained. The samples were resuspended in loading buffer (cat. no. P1040; Beijing Solarbio Science & Technology Co., Ltd.) and boiled for subsequent analysis via western blotting, according to the aforementioned protocol.

**Western blotting.** After washing twice with ice-cold PBS, the total protein in cells was lysed using RIPA (cat. no. R0020; Beijing Solarbio Science & Technology Co., Ltd.) lysis buffer for 30 min on ice. Cells were harvested and centrifuged at 20,664 x g at 4˚C for 15 min. In order to extract nuclear protein from the cells, cells were lysed using cytoplasmic protein extraction reagent lysis buffer for 30 min on ice. The proteins were collected from the supernatant after centrifugation at 20,664 x g at 4˚C for 15 min, and the nuclear protein extraction reagent was added to the cell precipitation for centrifugation at 20,664 x g at 4˚C for 15 min and collection of nuclear protein. Protein concentrations were determined using a BCA protein assay. Subsequently, proteins (20 µg) were separated via SDS-PAGE on 10% gels and transferred to PVDF membranes (EMD Millipore). After blocking with non-fat milk for 1 h at room temperature, the membranes were incubated overnight at 4˚C with the appropriate primary antibodies (all 1:1,000). After washing three times with TBS-0.1% Tween-20 (TBST) for 15 min per wash, the membranes were incubated with a goat anti-mouse or anti-rabbit HRP-conjugated secondary antibody (1:20,000) for 1 h at room temperature. The membranes were washed three times with TBST and then protein bands were visualized using an enhanced chemiluminescence kit (EMD Millipore) under using an Imager 600 (Amersham; Cytiva). GAPDH and histone H3 were used as the loading controls. Protein expression levels were analyzed by ImageJ (1.8.0) software (National Institutes of Health).

**Coimmunoprecipitation.** Following treatment with dioscin for 24 h, lysates were prepared according to the manufacturer's instructions (Active Motif, Inc.). Following centrifugation at 4˚C for 5 min at 137 x g, cells were suspended in 100 µl freshly prepared complete RIPA buffer and incubated for 30 min on a rotating platform at 4˚C. Whole cell extract was combined with 5 µg IgG (cat. no. 3900) or anti-MST2 (cat. no. 3952) antibodies (both from Cell Signaling Technology, Inc.) in a final volume of 500 µl and incubated overnight at 4˚C on a rotating platform. Subsequently, 25 µl protein G magnetic beads were added to the tube and incubated overnight at 4˚C on a rotating platform. The protein G magnetic beads were washed carefully and the protein samples were obtained. The samples were resuspended in loading buffer (cat. no. P1040; Beijing Solarbio Science & Technology Co., Ltd.) and boiled for subsequent analysis via western blotting, according to the aforementioned protocol.

**5-Ethynyl-2'-deoxyuridine (EdU) proliferation assay.** Cells (1x10⁴/well) were seeded into a 48-well plate and cultured for 48 h. The EdU Apollo DNA in vitro Kit (Guangzhou RiboBio Co., Ltd.) was used to assess cell proliferation. Cells were cultured with 200 µl EdU for 2 h and were then fixed with 4% paraformaldehyde for 30 min at room temperature. Subsequently, cells were treated for 10 min with 2 mg/ml glycine and 0.5% Triton X-100 at room temperature. Then, DAPI working solution was added to each well and incubated for 30 min in the dark at room temperature. Quantification
of EdU+ cells was expressed as a percentage of DAPI+ cells. The percentage of EdU+ cells was determined by fluorescence microscopy (Olympus Corporation).

**Lentiviral (LV) transduction.** Cells (1x10^5/well) were seeded into a 6-well plate and cultured for 24 h. At ~70% confluence, SCC15 cells were transduced with LV-homo-YAP particles [overexpression (OE) YAP group] or LV-3-negative control (NC; empty vector; OE NC group) for 12 h. Similarly, SCC15 cells were transduced with LV-homo-YAP-short hairpin (sh) RNA LV particles (sh YAP group) or a corresponding empty LV vector (sh NC group) for 12 h. The MOI for transduction was 20. All LV particles were purchased from Shanghai GenePharma Co., Ltd. After transduction for 48 h, cells were treated with 1 µg/ml puromycin for 2 weeks. Transduction efficiency was assessed via western blotting.

**Flow cytometric analysis of the cell cycle.** Cells (1x10^5) were harvested, fixed in 75% ethanol overnight at 4°C, washed with cold PBS and incubated with 0.5 ml propidium iodide/RNase (Tianjin Sungene Biotech Co., Ltd.) working solution for 30 min at room temperature. Subsequently, the cell cycle distribution was determined using an Accuri C6 flow cytometer (Becton, Dickinson and Company) and analyzed using FlowJo software (version 10; FlowJo LLC).

**Statistical analysis.** Data are presented as the mean ± SD of three independent experiments. Statistical analyses were performed using GraphPad Prism software (version 6; GraphPad Software, Inc.). Comparisons among multiple groups were analyzed using one-way ANOVA followed by Tukey’s post hoc test. P<0.05 was considered to indicate a statistically significant difference.

**Results**

**Dioscin reduces SCC15 cell proliferation.** To investigate the effects of dioscin on proliferation, SCC15 cells were treated with different concentrations of dioscin for 24 or 48 h, and cell proliferation was determined by performing CCK-8 assays. Compared with the control group (1.014±0.026), dioscin treatment significantly inhibited cell proliferation, even at 0.5 µM (0.820±0.004) (Fig. 1A). Similar results were obtained following treatment for 48 h (Fig. 1B). Moreover, the EdU incorporation assay results indicated that dioscin significantly decreased the ratio of EdU+ cells in a concentration-dependent manner (31.150±0.382, 22.92±0.352 and 18.100±0.311% at 0.5, 1 and 2 µM, respectively) compared with the control group (41.410±0.813%) (Fig. 1C and D). The results indicated that dioscin inhibited SCC15 cell proliferation in a dose-dependent manner.

**Dioscin induces SCC15 cell cycle arrest and apoptosis.** To explore the cellular mechanism underlying the antiproliferative effect of dioscin on SCC15 cells, flow cytometry was performed to detect alterations in the cell cycle and apoptosis. Following treatment with 0.5, 1 or 2 µM dioscin, the number of cells in the G2/M phase was significantly decreased (5.663±0.350, 1.816±0.304 and 0.5667±0.173%, respectively) compared with the control group (7.283±0.238%) (Fig. 2A and B). Moreover, following treatment with 1 or 2 µM dioscin, the number of cells in the G0/G1 phase was significantly increased (69.740±1.3960 and 74.200±0.305%, respectively) compared with the control group (65.830±0.644%). Subsequently, the effect of dioscin on cell apoptosis was evaluated. Compared with the control group (early apoptosis, 2.464±0.110%; late apoptosis, 6.254±0.078%), dioscin (0.5, 1 or 2 µM) significantly increased the population of early (3.100±0.112, 5.624±0.248 and 7.790±0.330%, respectively) and late (8.142±0.072, 15.10±0.353 and 21.020±0.213%, respectively) apoptotic cells (Fig. 2C and D). The apoptosis-inducing effect of dioscin was enhanced with increasing concentrations.

**Dioscin-induced inhibition of cell proliferation is mediated by the Hippo signaling pathway.** Previous research has demonstrated that RASSF1 supports the phosphorylation of MST2 and SAV, both of which are upstream molecules of YAP (22). A previous study reported that dioscin induced RASSF1 demethylation and inhibited the proliferative activity of bladder cancer cell lines (21). Therefore, the present study aimed to investigate the role of the Hippo signaling pathway in dioscin-treated cells. SCC15 cells were treated with different concentrations of dioscin (0, 0.5, 1 or 2 µM) for 24 h or with 1 µM dioscin for different durations (0, 2, 4 or 8 h). The ratios of p-LATS (Ser909)/LATS, p-MST/MST2 and p-YAP (Ser397)/YAP were significantly increased by dioscin treatment in a concentration- and time-dependent manner compared with the control group (Fig. 3A-D). Compared with the control group, dioscin did not notably alter the expression levels of total LATS1 and MST2, but markedly decreased the expression levels of total YAP (Fig. 3A-D). In addition, the expression levels of cell cycle- and apoptosis-related proteins were assessed. Cyclin D1 and BCL-2 expression levels were markedly decreased by dioscin in a concentration- and time-dependent manner compared with the control group (Fig. 3E and F). However, BAX expression levels displayed the opposite pattern. To investigate whether dioscin treatment altered the localization of YAP, nuclear and cytoplasmic proteins were extracted using the Nuclear Protein Extraction kit. As dioscin concentrations increased, nuclear YAP expression levels were markedly decreased, whereas YAP cytoplasmic accumulation was notably increased compared with the control group (Fig. 3G).

To detect Hippo signaling pathway activation following dioscin treatment, the expression levels of RASSF1A, a protein consistently linked to tumor onset and poor cancer prognosis (23), were detected. The western blotting results indicated that RASSF1A expression levels were markedly increased...
by dioscin treatment in a concentration- and time-dependent manner compared with the control group (Fig. 3A and B). However, the mechanism underlying dioscin-induced inactivation of the Hippo signaling pathway is not completely understood. RASSF1A has been reported to form a complex with MST2 and SAV, which phosphorylates LATS1, leading to YAP phosphorylation (24). The aforementioned results indicated that dioscin treatment resulted in RASSF1A accumulation (Fig. 3A and B), and endogenous RASSF1A or LATS1 were coimmunoprecipitated with the MST2 protein (Fig. 3H). As shown in Fig. 3H, RASSF1A and LATS1 were detected in the anti-MST2 precipitate, but not in the IgG precipitate.

**YAP overexpression abrogates dioscin-induced inhibition of cell proliferation.** The aforementioned results indicated that the RASSF1A/MST2/YAP axis was involved in dioscin-induced inhibition of cell proliferation. To evaluate the hypothesis, SCC15 cells were transduced with OE YAP or OE NC and then treated with 1 µM dioscin. The flow cytometry results demonstrated that, compared with the OE NC group (3.450±0.886%), dioscin treatment significantly decreased the number of cells in the G2/M phase (1.097±0.112%), whereas the number of cells in the proliferative phases was significantly increased in the OE YAP group (7.887±0.206%) compared with the OE NC group. (Fig. 4A and B). Meanwhile, compared with the OE NC + dioscin group (1.097±0.112%), the increase of proliferation activity of OE YAP + dioscin group (6.890±0.237%), indicated that overexpression of YAP gene abrogated dioscin-induced inhibition of cell proliferation. (Fig. 4B). The apoptosis assay results demonstrated that dioscin treatment significantly increased the number of apoptotic cells (39.010±0.464%), whereas in the OE YAP group this increase was attenuated (19.150±0.831%) compared with the OE NC group (24.770±0.382%) (Fig. 4C and D). The EdU incorporation (Fig. 4E and F) and CCK-8 (Fig. 4G) assay results demonstrated that, compared with the OE NC group or OE YAP group, dioscin treatment significantly suppressed SCC15 cell proliferation in the OE NC + dioscin and OE YAP + dioscin groups, whereas YAP overexpression in the OE YAP or OE YAP + dioscin groups enhanced cell proliferation compared with the OE NC or OE NC + dioscin groups, respectively. The dioscin-induced suppression of proliferation
was inhibited in the OE YAP group. The protein expression levels of cell cycle- and apoptosis-related proteins were measured via western blotting. The western blotting results further verified the alterations in the cell cycle and apoptosis (Fig. 4H).

YAP knockdown enhances the antitumor activity of dioscin. To further understand the role of YAP in mediating the suppressive effects of dioscin on SCC15 cells, SCC15 cells were transduced with sh YAP or sh NC and then treated with 1 µM dioscin. Flow cytometry was performed to assess alterations in cell cycle distribution and apoptosis. Both YAP knockdown and dioscin treatment led to a decrease of cells in G2/M phase, and the combination of these interventions augmented this effect (G2/M phase cells: sh NC, 11.770±0.0880%; sh NC + dioscin, 11.000±0.152%; sh YAP, 10.300±0.152%; and sh YAP + dioscin, 6.263±0.489%) (Fig. 5A and B). Moreover, compared with the sh NC group (19.860±0.474%), the proportion of apoptotic cells was increased following dioscin treatment (21.600±0.462%) (not statistically significant) or YAP knockdown (22.010±0.353%), and the combination of these two interventions was synergistic (27.020±0.277%) (Fig. 5C and D). Subsequently, SCC15 cell proliferation was evaluated by performing EdU incorporation (Fig. 5E and F) and CCK-8 (Fig. 5G) assays. Compared with the sh NC group, both dioscin and YAP knockdown significantly decreased cell proliferation. In addition, the combination of dioscin treatment and YAP knockdown resulted in further inhibition of cell proliferation compared with either intervention alone, as determined by flow cytometry, EdU staining, western blotting and CCK-8 assay at 24 h.

Subsequently, the expression levels of YAP, cyclin D1, BCL-2 and BAX were measured. Similarly, the combination of dioscin treatment and YAP knockdown led to more notable suppressive effects on protein expression compared with either intervention alone. However, the expression levels of the proapoptotic protein BAX displayed the opposite trends (Fig. 5H).
Figure 3. Dioscin inhibits SCC15 cell proliferation via inactivating the Hippo signaling pathway. Following treatment with (A) 0, 0.5, 1 or 2 µM dioscin for 24 h or (B) 1 µM dioscin for 0, 2, 4 or 8 h, Hippo signaling pathway-related protein expression levels were measured via western blotting. Hippo signaling pathway-related protein expression levels were semi-quantified in SCC15 cells treated with (C) 0, 0.5, 1 or 2 µM dioscin for 24 h or (D) 1 µM dioscin for 0, 2, 4 or 8 h. Following treatment with (E) 0, 0.5, 1 or 2 µM dioscin for 24 h or (F) 1 µM dioscin for 0, 2, 4 or 8 h, cell cycle and apoptosis-related protein expression levels were determined via western blotting. (G) Following treatment with 0, 0.5, 1 or 2 µM dioscin for 24 h, nuclear and cytoplasmic proteins were extracted and detected via western blotting. (H) Following treatment with 1 µM dioscin for 24 h, MST2 immunoprecipitation was detected via coprecipitation with RASSF1A and LATS1 followed by western blotting analysis. Data are presented as the mean ± SD of three independent experiments. *P<0.05 and ***P<0.001. RASSF1A, Ras association domain-containing protein 1A; p, phosphorylated; LATS, large tumor suppressor kinase; MST, macrophage stimulating; YAP, yes-associated protein; Cyto, cytoplasmic; ip, immunoprecipitation.
Figure 4. YAP OE reverses dioscin-mediated inhibition of cell proliferation. Effects of YAP OE and dioscin treatment on the cell cycle distribution were (A) determined via flow cytometry and (B) quantified. Effects of YAP OE and dioscin treatment on cell apoptosis were (C) determined via flow cytometry and (D) quantified. Cell proliferation was (E) detected by performing EdU staining (magnification, x200) and (F) quantified. (G) Cell proliferation was also assessed by performing Cell Counting Kit-8 assays. (H) Protein expression levels were measured via western blotting. Data are presented as the mean ± SD of three independent experiments. *P<0.05, **P<0.01 and ***P<0.001. YAP, yes-associated protein; OE, overexpression; NC, negative control; OD, optical density; ns, not significant; EdU, 5-ethynyl-2'-deoxyuridine.
Figure 5. YAP knockdown enhances dioscin-induced inhibition of cell proliferation. Effects of YAP knockdown and dioscin treatment on SCC15 cell cycle distribution were (A) determined via flow cytometry and (B) quantified. Effects of YAP knockdown and dioscin treatment on SCC15 cell apoptosis were (C) determined via flow cytometry and (D) quantified. Cell proliferation was (E) determined by performing EdU staining (magnification, x200) and (F) quantified. (G) Cell proliferation was also assessed by performing Cell Counting Kit-8 assays. (H) Protein expression levels were determined via western blotting. Data are presented as the mean ± SD of three independent experiments. *P<0.05, **P<0.01 and ***P<0.001. YAP, yes-associated protein; sh, short hairpin; NC, negative control; APC, allophycocyanin; OD, optical density; EdU, 5-ethynyl-2'-deoxyuridine.
Discussion

OSCC is one of the most common and malignant tumors in the oral and maxillofacial regions; however, following systemic treatment, the prognosis of patients with OSCC is still poor (25). In addition to the health and functional abnormalities that typically result from tumors, OSCC often affects the appearance of patients, leading to a heavy psychological burden (26).

Dioscin is a natural steroidal saponin that can be extracted from various traditional Chinese medicines, including Dioscorea. Dioscin has been reported to exert a variety of biological activities, including protection against liver damage (27) and anti-inflammatory activity (28). Moreover, dioscin may produce anticancer effects on various types of cancer cells (29-32). Previous studies have indicated that dioscin induced cervical carcinoma cell apoptosis (33) and suppressed ovarian cancer cell viability (34). The present study demonstrated that, compared with the control group, dioscin resulted in cell cycle arrest and increased apoptosis, which was consistent with a previous study that indicated that dioscin protected against cancer progression (35). The present study also assessed cell cycle- and apoptosis-related protein expression levels, including cyclin D1, BCL-2 and BAX. The alterations in protein expression levels were consistent with the alterations in biological activities.

YAP is the hub protein in the Hippo signaling pathway (36). Previous studies have demonstrated that the Hippo signaling pathway is inactivated and YAP protein expression is increased in numerous types of cancer, including cervical cancer, urothelial cell carcinoma and oral squamous cell carcinoma (37-39). Direct inhibition of the YAP oncoprotein is an anticancer therapeutic strategy (40). A previous study reported that dioscin downregulated the expression of tafazzin, a homolog protein of YAP, in hepatocellular carcinoma; however, alterations in YAP expression were not investigated (41). The present study demonstrated that, compared with the control group, dioscin notably deactivated YAP protein expression by increasing its phosphorylation, and nuclear YAP accumulation was markedly decreased following dioscin treatment. Therefore, it was hypothesized that dioscin inhibited OSCC cell viability by downregulating YAP expression. To further investigate whether the effects of dioscin on the biological characteristics of OSCC cells were mediated via YAP, YAP expression was altered in OSCC cells via LV transduction. YAP overexpression and knockdown reduced and enhanced dioscin-induced antitumor effects, respectively.

RASSF1A is one of the most commonly inactivated tumor suppressor genes in sporadic human malignant tumors (42). RASSF1A activation is an important factor in the pathogenesis and progression of solid tumors (43,44). A previous study reported that dioscin demethylated the RASSF1A gene in bladder cancer cells, thus inhibiting cancer cell proliferation (21). The present study demonstrated that RASSF1A protein expression levels were increased by dioscin treatment in a concentration- and time-dependent manner compared with the control group. Increases in RASSF1A expression provided a potential explanation for the inhibitory effects of dioscin on OSCC cell proliferation. A previous study demonstrated that ATM serine/threonine kinase directly phosphorylated the Ser131 site in RASSF1A, activating MST2 and LATS1 (22), which then phosphorylated YAP and induced cell apoptosis.

The results of the present study demonstrated that dioscin treatment enhanced the binding of MST2 with RASSF1A and LATS1, which could be explained by the augmentation of YAP phosphorylation.

To conclude, the present study demonstrated that dioscin resulted in OSCC cell cycle arrest and apoptosis, which involved the RASSF1A/MST2/YAP signaling axis. The present study indicated that dioscin may serve as a useful therapeutic agent for OSCC or other types of tumors.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

HT and XC performed the experiments, analyzed the data, prepared the figures and tables, and wrote the manuscript. YZ and YW assisted in the data collection and analyses. YWA and XF assisted in data analyses. HT and XC confirmed the authenticity of all the raw data. WG was involved in designing the experiments and drafting the manuscript. YX assisted in the data collection and analyses. YWW and YWX assisted in data analyses. HT and XC performed the experiments, analyzed the data, and YX assisted in drafting the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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