

# A traditional Chinese medicine Huaier triggers G1 cell cycle arrest and apoptosis through cyclins-CDKs-CKIs machinery in MOLT4 cells

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**Abstract** *Objective:* The purpose of the study was to study the effect of Huaier, a traditional Chinese medicine, on the cell cycle adjustment in MOLT4 cells *in vitro*. *Methods:* We used MTT assay to test cell viability, flow cytometry to detect cell cycle and apoptosis and western blot to examine the expression of cell-cycle and apoptotic proteins in MOLT4 cells induced by Huaier. *Results:* Huaier could reduce the viability of MOLT4 cell by inducing G1 arrest and apoptosis. The induction of apoptosis after treatment with Huaier for 24 h was demonstrated in a dose- and time-dependent manner by flow cytometry analysis. G1 arrest induced by Huaier was modulated through the increased expression of Cdk proteins (p21<sup>cip/waf1</sup> and p27<sup>kip1</sup>) with a simultaneous decrease in Cdk2, Cdk4, Cdk6, cyclin D1 and cyclin E expression. Huaier also induced Bax and Bcl-2 expression and activation of Caspase-3. *Conclusion:* It is firstly demonstrated that Huaier can inhibit proliferation of MOLT4 cells via G1 arrest and apoptosis. These results suggest that Huaier is a cell-cycle anti-cancer drug.

**Key words** Chinese medicine; apoptosis; cell cycle arrest; MOLT4

The normal cell cycle core machinery is a family complexes of catalytic components called cyclin-dependent protein kinases (Cdks) and regulatory subunits named cyclins [1]. Distinct cyclin-Cdk complexes sequentially phosphorylate their respective substrates, thereby driving the cell through different phases of the cell cycle. Three essential classes of cyclin-Cdk complexes include the D-type cyclins (cyclins D1, D2 and D3), which activate Cdk4 and Cdk6 to execute critical events in G1 phase, the E-type and A-type cyclins, which activate Cdk2 to drive the cell through S phase, and the A-type and B-type cyclins, which activate Cdk1 to direct structural and regulatory events in mitosis [2-4]. Two families of Cdk inhibitors (CKIs) are known as p16<sup>INK4A</sup> family (i.e. p16<sup>INK4A</sup>, p15<sup>INK4B</sup>, p18<sup>INK4C</sup>, p19<sup>INK4D</sup>) and the p21<sup>CIP1</sup> family (i.e. p21<sup>CIP1</sup>, p27<sup>KIP1</sup> and p57<sup>KIP2</sup>) [4].

One hallmark of cancer is uncontrolled cell proliferation and/or insufficient apoptosis [5]. To inhibit cancer cell proliferation and induce apoptosis, most anti-tumor drugs were developed for targeting different cell cycle and apoptosis regulators [6-7].

Huaier, which come from an important traditional Chinese medicinal herb *Fungi* growing on the Chinese scholar tree, has a long history as an effective anti-can-

cer drug in China. Being extracted from the *trametes robiniohila murr*, the major active ingredient of Huaier is a polysaccharide protein which consists of 6 kinds of monosaccharides and 18 kinds of amino acids [8].

In this study, we planned to investigate the biological function and molecular targets of Huaier in human cancer cells.

## Materials and methods

### Preparation of Huaier dissolution

Huaier was provided by Gaitianli Pharmaceutical Co.Ltd. (China). Five mg powder was dissolved in 1 mL water, and thus its initial concentration was 5 mg/mL (each culture well containing 1 mL RPMI 1640 medium).

### Cell culture

The acute lymphocytic leukemia cell line MOLT4 (ATCC, USA) was cultured in a six-well plate with DMEM medium containing 10% fetal bovine serum, 100 U/mL penicillin, 100 µg/mL streptomycin, and 2 mM L-glutamine. All media, supplements, and sera were purchased from GIBCO (USA). Cells were incubated in a 5% CO<sub>2</sub> humidified incubator at 37 °C. The culture was diluted and re-plated every 3 days to keep them in an asynchro-

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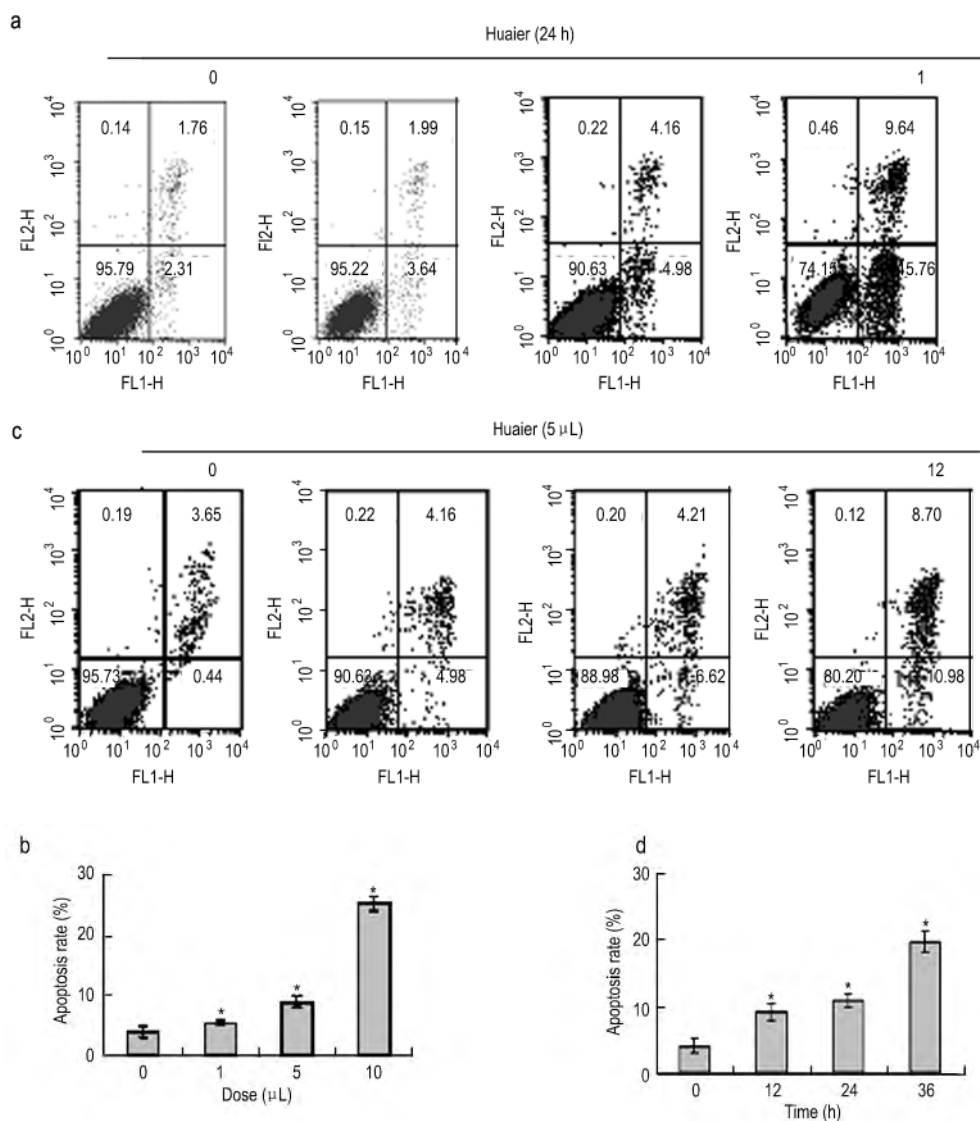


Fig. 1 Effect of Huaier on MOLT4 cells apoptosis with different doses or at different times. (a) and (c) showed Annexin V/PI assay was performed to detect apoptosis with different doses or at different times; (b) and (d) display histogram was used to show an increase of apoptosis rates as doses rised or time went by. \*  $P < 0.05$

nous and exponential phase of growth.

Huaier solution was added into 4 wells with 0  $\mu\text{L}$ , 1  $\mu\text{L}$ , 5  $\mu\text{L}$  and 10  $\mu\text{L}$ , respectively. After the cells grew for 24 h, they were then rinsed with phosphate buffered saline (PBS), centrifugated and harvested, while Huaier solution was also added to another 4 wells with 5  $\mu\text{L}$ . After cells were incubated for 0 h, 12 h, 24 h and 36 h and then rinsed with PBS, centrifugated and harvested.

### Flow cytometric analysis

#### *Annexin V/PI assay for apoptosis*

Five  $\mu\text{L}$  FITC-Annexin V and 10  $\mu\text{L}$  PI were added to harvested fresh cells that had been resuspended in 100  $\mu\text{L}$  cold binding buffer at a density of  $10^6$  cells/mL. The cells

were then incubated at room temperature in the dark for 30 min, and finally detected by flow cytometry and analyzed with Cellquest software (FACSVantage, Becton Dickinson, USA).

#### *PI assay for cell cycle analysis based on DNA content and distribution*

Harvested cells were re-suspended in 1 mL of 0.1% sodium citrate containing 0.3% NP-40, 0.0002 mg/mL RNase and 50  $\mu\text{g}/\text{mL}$  propidium iodide, and then were incubated on ice in the dark for 30 min. Cell cycle were analyzed on a FACSort with Cellquest (Becton Dickinson, USA) and ModFit software (Verity Software House, USA).

#### *$\gamma\text{H2AX}$ for DNA damage*

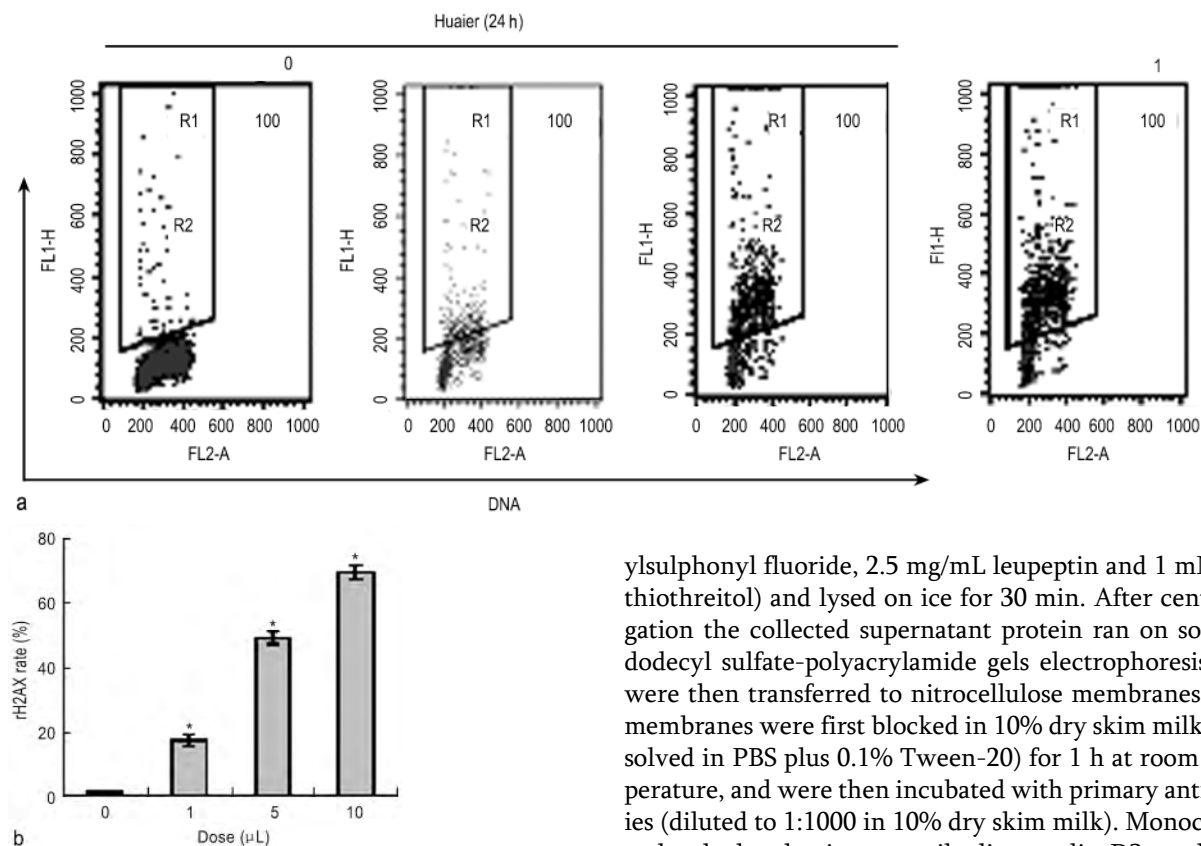


Fig. 2 Effect of different doses Huaier on MOLT4 cell cycle distribution after 24 h. (a) PI assay was performed to detect every percentage of G1, S and G2/M; (b) histogram was used to show an increase of G1 percentage and a decrease of G2/M percentage. \*  $P < 0.05$

Harvested cells were fixed by 1% formaldehyde in PBS without methanol on ice for 15 min, and rinsed, then centrifugated. The collected cells were fixed by ice-cold 80% ethanol at  $-20^{\circ}\text{C}$  for at least 24 h. These cells were washed with PBS twice and permeated with 0.5% Triton X-100 in PBS on ice for 5 min. After centrifugation, they were incubated overnight at  $4^{\circ}\text{C}$  in the presence of primary antibody  $\gamma\text{H2AX}$  [BD PharMingen; diluted in PBS containing 1% bovine serum albumin (BSA)]. The next day cells were rinsed and incubated with the secondary FITC-conjugated antibody (DAKO, Denmark; diluted in PBS containing 1% BSA) for 30 min. Finally, cells were rinsed and resuspended in propidium iodide solution (50  $\mu\text{g}/\text{mL}$  PI) and incubated at room temperature for 30 min. Cell fluorescence was measured by a FACSVantage flow cytometry (Becton Dickinson).

### Western blotting detection

Harvested cells were re-suspended in lysis buffer (50 mM HEPES, pH 7.4, 250 mM NaCl, 2 mM ethyleneglycoltetraacetic acid, 2 mM ethylenediamine-tetraacetic acid, 1 mM NaF, 0.1 mM vanadate, 0.1 mM phenylmeth-

ylsulphonyl fluoride, 2.5 mg/mL leupeptin and 1 mM dithiothreitol) and lysed on ice for 30 min. After centrifugation the collected supernatant protein ran on sodium dodecyl sulfate-polyacrylamide gels electrophoresis and were then transferred to nitrocellulose membranes. The membranes were first blocked in 10% dry skim milk (dissolved in PBS plus 0.1% Tween-20) for 1 h at room temperature, and were then incubated with primary antibodies (diluted to 1:1000 in 10% dry skim milk). Monoclonal and polyclonal primary antibodies: cyclin D3, cyclin E, CDK2, CDK4, CDK6, p16, p21, p27, Bcl-2, Bax and Beta-actin antibodies (BD PharMingen) overnight at  $4^{\circ}\text{C}$ . Next day the membranes were rinsed twice and incubated with the horseradish peroxidase-coupled secondary antibody (diluted to 1:5000 in 10% dry skim milk. Wuhan Boster Biological Technology Ltd., China) at room temperature for 2 h. Detection was performed by ECL system (Amersham Pharmacia, Tokyo).

### Statistical analysis

The study was repeated thrice. The statistical significance of all results was evaluated by paired Student's  $t$ -test. Data were presented with  $P < 0.05$  accepted as significance.

## Results

### Huaier treatment induces a dose- and time-dependent apoptosis in MOLT4

To examine its apoptotic effect on MOLT4 cells, different dosages of Huaier (0  $\mu\text{L}$ , 1  $\mu\text{L}$ , 5  $\mu\text{L}$ , 10  $\mu\text{L}$ ) were treated for 24 h, and cell apoptosis was determined by Annexin V/PI assay. With the increased dosage of Huaier, the percentage of cell apoptosis also increased from 4.07% to 25.40% (Fig. 1a and 1b). In addition, apoptosis at different time courses (0 h, 12 h, 24 h, 36 h) was detected. Similarly, as the time went by, the apoptosis rate increased

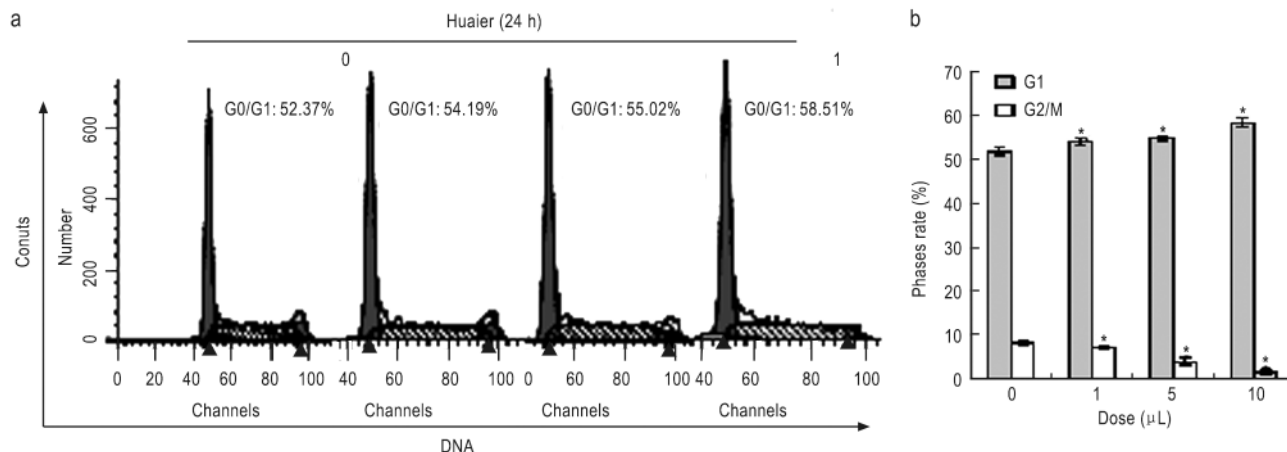


Fig. 3 Effect of different doses Huaier on MOLT4 cells DNA damage after 24 h. (a)  $\gamma$ H2AX/DNA multiparameter assay was performed to detect the expression of  $\gamma$ H2AX foci; (b) histogram was used to show an aggravation of DNA damage degree as doses increased. \*  $P < 0.05$

from 4.09% to 19.68% (Fig. 1c and 1d).

To examine the effect of Huaier on cell proliferation, we analyzed cell cycle distribution of MOLT4 cells treated at different dosages (0  $\mu$ L, 1  $\mu$ L, 5  $\mu$ L, 10  $\mu$ L) for 24 h. Flow cytometry data showed that cell cycle was arrested in G1 phase, which was indicated by increasing in percentage G1 phase from 52.37% to 58.51% with a dose-dependent

manner and concomitant decreasing in the percentage of G2/M cells from 8.22% to 1.50%, whereas the S phase did not change significantly (Fig. 2).

To examine the DNA damage induced by Huaier, we measured the expression of  $\gamma$ H2AX protein by multiparameter flow cytometric analysis. After MOLT4 cells were treated with Huaier (0  $\mu$ L, 1  $\mu$ L, 5  $\mu$ L, 10  $\mu$ L) for 24 h, the expression of  $\gamma$ H2AX protein increased significantly compared to control treatment (from 0.96% to 69.71%, Fig. 3).

We further determined the molecular mechanism of Huaier in inducing G1 arrest and apoptosis. Expression of G1- and apoptosis-associated proteins in MOLT4 cells treated with Huaier was detected by Western blotting. Our data showed that Huaier induced a dramatic upregulation of p16, p21 and p27 and downregulation of cyclin D3, cyclin E, CDK2, CDK4 and CDK 6 in a dose-dependent manner (Fig. 4). The apoptosis-associated Bcl-2 family includes anti-apoptotic (such as Bcl-2) and pro-apoptotic proteins (such as Bax) [9]. Western blotting analysis showed that the expression of Bax protein was significantly increased, whereas the expression of Bcl-2 protein was markedly decreased in the cells treated with Huaier (Fig. 4). The ratio of Bax/Bcl-2 showed a significant increase, which is believed to induce apoptosis in MOLT4 cells [10].

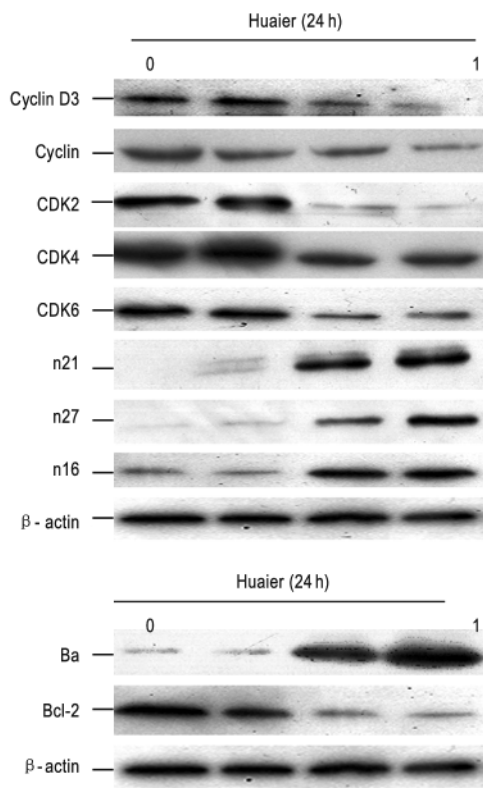


Fig. 4 Effect of different doses Huaier on the expression of G1- and apoptosis-associated proteins after 24 h by Western blot analysis

## Discussion

Huaier used to be extracted for the treatment of many kinds of carcinomas in the practice of Chinese traditional medicine, such as liver cancer [11], breast cancer [12], lung cancer [13]. Besides, we found it have also the same effect on colorectal cell line SW480 and human leukemic cell line HL-60 (data not shown), but better effect on MOLT4 cells. It is reported that Huaier can induce cells to secrete

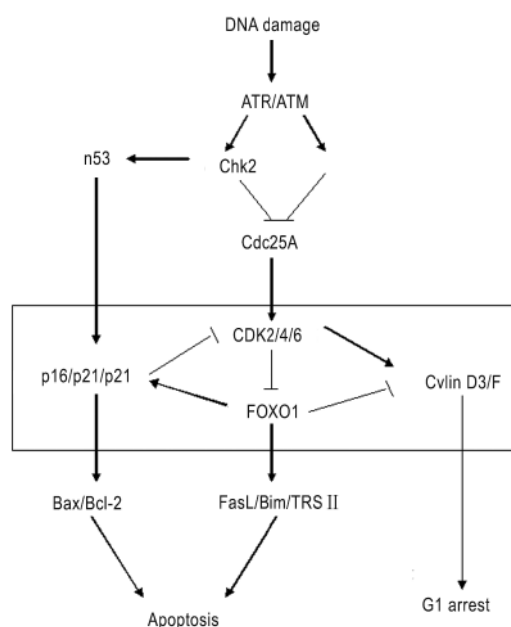


Fig. 5 Huaier triggers G1 cell cycle arrest and apoptosis probably through cyclins-CDKs-CKIs machinery in MOLT4 cells

IFN- $\alpha$  and IFN- $\gamma$ , then activate NK cells, and finally in turn kill cancer cells [14].

In our study, we showed that Huaier could inhibit cancer cell proliferation by triggering G1 cell cycle arrest and induce cell apoptosis, which provided the *in vitro* evidence for its anti-tumor effect. Huaier-induced DNA Stranded Break indicated by  $\gamma$ H2AX upregulation was measured by flow cytometry (Fig. 3). It can activate DNA checkpoints including Ataxia telangiectasia mutated (ATM) and/or ATM and Rad 3-related (ATR) [15–17], which inhibits CDKs to arrest cell cycle progression, allowing cells to repair the damaged DNA or removing those cells by inducing apoptosis whose DNA damage can not be repaired [18–20]. ATM and ATR were shown to activate Chk1 and Chk2 respectively [21–23], and the latter can cause ubiquitin-dependent degradation of protein phosphatase Cdc25A, which inhibits CDK2 [24–25], CDK4 and CDK6 (Fig. 5). It was reported that CDK2 phosphorylates FOXO1 at ser249 *in vivo* and inhibit FOXO1 function [26–27]. FOXO1, as a transcription factor, on the one hand up-regulates p21 [28], p27 [29–30] and p16; on the other hand, it represses cyclin D3 expression [31–32]. Our data showed that in Huaier-treated cells, CDK2 was down-regulated; p21, p27, and p16 were up-regulated, as well as cyclin D3 was down-regulated, which is consistent with previous studies. Activated FOXO1 can also trigger a complicated network of mitochondria apoptotic signaling, such as Bax/Bcl-2 pathway [11] activated indirectly by p21, p27, and p16 [28–30, 33–34] (Fig. 4), and/or death receptor apoptotic signaling, such as FasL/Bim/TRAIL pathway activated di-

rectly by FOXO1 [26–27]. Both apoptotic signaling can activate common downstream caspase-3 to trigger apoptosis (Fig. 5).

In summary, we have shown that Huaier can effectively inhibit cell proliferation by inducing G1 arrest through the down-regulation of CDK2/CDK4/CDK6, and cyclin D3/cyclin E, together with the up-regulation of CKIs including p16/p21/p27. Apoptosis was accompanied by the G1 cell cycle arrest. To our knowledge, this is the first study that shows the involvement of the cyclins-CDKs-CKIs machinery during the G1 arrest and apoptosis of MOLT4 cells treated with Huaier. These findings indicate that Huaier could be a potential therapeutic anti-cancer agent. In our future study, we will further purify the Huaier's monosomy effective ingredient and elucidate the mechanism of Huaier-triggered both mitochondria- and death receptor-mediated apoptotic pathways.

### Conflicts of interest

The authors indicated no potential conflicts of interest.

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