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> Authors' Contribution: Study Data Col

Manuscript Prepa Literature Funds Col **Ginsenoside Rg1 Inhibits Cell Proliferation** and Induces Markers of Cell Senescence in CD34+CD38- Leukemia Stem Cells Derived from KG1 α Acute Myeloid Leukemia Cells by Activating the Sirtuin 1 (SIRT1)/Tuberous Sclerosis Complex 2 (TSC2) Signaling Pathway

Study Design A Data Collection B Statistical Analysis C Data Interpretation D Aanuscript Preparation E Literature Search F Funds Collection G	BCDEF 2 BCDF 2 ABDEG 3 BCDF 4 BCF 3 BF 3	Cheng-Gui Zhang Heng Liu Yue Zhou Ya-Ping Wang Yuan Li Yan-Jun Han	 Yunnan, P.R. China 2 Key Laboratory of Insect Biomedicine Research and Development in Yunnan Province, Dali University, Dali, Yunnan, P.R. China 3 Department of Histology and Embryology, Key Laboratory of Cell Biology, Dali University, Dali, Yunnan, P.R. China 4 Laboratory of Stem Cell and Tissue Engineering, Department of Histology and Embryology, Chongqing Medical University, Chongqing, P.R. China 	
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Background:		Clinical relapse in acute myeloid leukemia (AML) is associated with the reduced treatment response of leukemia stem cells (LSCs). This study aimed to investigate the effects of the ginseng derivative, ginsenoside Rg1 (Rg1), on CD34+CD38– LSCs derived from KG1α human acute myeloid leukemia cells.		
Material/Methods:		CD34+CD38– LSCs were isolated from KG1 α human acute myeloid leukemia cells by cell sorting. CD34+CD38– KG1 α LSCs were divided into the control group and the Rg1 group (treated with Rg1). The cell counting kit-8 (CCK-8) assay evaluated the proliferation of CD34+CD38– KG1 α LSCs and flow cytometry stud- ied the cell cycle. The mixed colony-forming unit (CFU-Mix) assay and staining for senescence-associated beta- galactosidase (SA- β -Gal) evaluated cell senescence. Expression of sirtuin 1 (SIRT1) and tuberous sclerosis complex 2 (TSC2) were evaluated using Western blot and quantitative reverse transcription-polymerase chain reaction (gRT-PCR).		
Results:		CD34+CD38– KG1 α LSCs were isolated at 98.72%. Rg1 significantly reduced the proliferation of CD34+CD38– KG1 α LSCs compared with the control group (p<0.05). Cells in the G0/G1 phase were significantly increased, and cells in the G2/M and S phase were significantly reduced compared with the control group (p<0.05). Rg1 significantly increased SA- β -Gal and reduced CFU-Mix formation compared with the control group (p<0.05), significantly down-regulated SIRT1 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05), and significantly reduced TSC2 expression in CD34+CD38– KG1 α LSCs compared with the control group (p<0.05).		

Rg1 inhibited cell proliferation and induced cell senescence markers in CD34+CD38-KG1α LSCs by activating **Conclusions:** the SIRT1/TSC2 signaling pathway.

MeSH Keywords: Adult Stem Cells • Ginsenosides • Leukemia, Myeloid, Acute

trol group (p<0.05).

Full-text PDF:

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Background

Acute myeloid leukemia (AML) arises from progenitor hematopoietic cells and is associated with molecular changes that affect cell proliferation and differentiation [1,2]. Clinical relapse in AML is associated with the reduced treatment response of leukemia stem cells (LSCs) [1,2]. Clinically, adult patients with AML have a poor clinical outcome with a 5-year overall survival (OS) rate of 50% for younger adults, and less than oneyear OS for older patients [3]. Although chemotherapy may reduce the symptoms of AML, relapse is common due to chemoresistance [4]. Therefore, novel treatment strategies continue to be investigated.

In AML, LSCs are associated with disease occurrence and disease persistence [5,6]. The LSC phenotype was originally identified as CD34+ CD38- [7]. The identification of this LSC phenotype has resulted in the ability to label and select LSCs by cell sorting techniques, which may lead to the removal of this cell population as a novel therapeutic approach in patients with AML [8,9]. Cell aging, or senescence, limits cell survival and the proliferative capabilities of cells by inducing the dysfunction of cellular metabolism [10]. Although cancer is a disease of aging and the incidence increases with age [11], cancer cells show properties of cell immortalization, and inducing the senescence of malignant cells may suppress tumor growth, including in leukemias such as AML.

Ginsenoside Rg1 (Rg1), derived *Panax ginseng* and *Panax japonicus*, has previously been reported to have antitumor effects by inhibiting radiation-resistance and oxidation [12]. Previous studies have shown that Rg1 could affect cell senescence and cell proliferation of normal hematopoietic progenitor cells (HPCs) and hematopoietic stem cells (HSCs) [13,14]. Rg1 has been shown to inhibit the proliferation of cells of the AML cell line, K562, and to accelerate cell senescence [15]. However, the effects of Rg1 on the senescence of LSCs remain poorly understood.

Sirtuin 1 (SIRT1) participates in many biological functions, including cell senescence, and is involved in the pathogenesis of many age-related disorders [16]. The tuberous sclerosis complex 2 (TSC2) molecule and gene are involved in both cell autophagy and senescence [17].

Therefore, this study aimed to investigate the effects of the ginseng derivative, ginsenoside Rg1 (Rg1), on CD34+CD38–LSCs derived from KG1 α acute myeloid leukemia cells *in vitro*. Cell senescence was studied using a mixed colony-forming unit (CFU-Mix) assay and detection of senescence-associated betagalactosidase (SA- β -Gal).

Material and Methods

Cell culture of KG1 α human acute myeloid leukemia (AML) cells and cell sorting to isolate CD34+CD38- leukemia stem cells (LSCs)

The human acute myeloid leukemia (AML) cell line, KG1 α , was purchased from American Type Culture Collection (ATCC) (Manassas, VA, USA). The KG1 α cells were cultured in the Iscove's Modified Dulbecco's Medium (IMDM) (Gibco BRL Co. Ltd., Grand Island, NY, USA) supplemented with 10% fetal bovine serum (FBS) (Gibco BRL Co. Ltd., Grand Island, NY, USA), 100 µg/ml of streptomycin, and 100 units/ml of penicillin (Beyotime Biotech., Shanghai, China) at 37°C and 5% CO₂.

Cell sorting for CD34+CD38– KG1 α cell was performed by immunostaining with an antibody to CD38 conjugated to allophycocyanin (CD38–APC) (eBioscience, Thermo Fisher Scientific, San Diego, CA, USA) and an antibody to CD34 conjugated with fluorescein isothiocyanate (CD-34-FITC) (BD Bioscience, Oxford, UK) at 4°C for 30 min, according to the previously published method [18]. The stained cells were washed using phosphate-buffered saline (PBS) (Beyotime Biotech., Shanghai, China) and sorted using a FACS Aria III flow cytometer (BD Bioscience, Oxford, UK).

Study groups

Ginsenoside Rg1 (Rg1) was purchased from Hongjiu BioTech (Dalian, China) (Cat. No. 060427), and had a purity >95%. In this study, the sorted CD34+CD38–LSCs were divided into the control group and the Rg1 group. The CD34+CD38–LSCs in the control group were cultured in IMDM for 48 h. The CD34+CD38–LSCs in the Rg1 group were cultured in IMDM for 48 h and treated with Rg1 dissolved in dimethyl sulfoxide (DMSO), at a final concentration of 40 μ mol/l. The control group consisted of CD34+CD38–LSCs treating with DMSO alone.

Cell counting kit 8 (CCK-8) assay

The cytotoxic effect and cell proliferative activity of CD34+CD38– LSCs were studied using the CCK-8 kit (Cat. No. C0037) (Beyotime Biotech. Shanghai, China), according to the manufacturer's instructions. Briefly, the cells were seeded onto the 96-well plates at a density of 1×10^8 cells/well and cultured at 37°C and 5% CO₂ for 48 h. The CCK-8 reagents were added to the culture plates at a dose of 20 µl/well and incubated for another 2 h. The optical density (OD) of cells was examined using an ELx808 microplate reader (Bio-Tek Inc., Winooski, VT, USA) at a wavelength of 450 nm. The inhibition rate (%)=(the OD of the control group–the OD of the Rg1 group)/(the OD of the control–the OD of the blank group)×100%.

Table 1. Primers used in this study.

Genes		Sequences	Length (bp)	
CIDT1	Forward	CAAGGGATGGTATTTATGCTCG	172	
SIRT1	Reverse	CAAGGCTATGAATTTGTGACAGAG	173	
TSC2	Forward	TCTGTCTGCTGTGCGTCCG		
1502	Reverse	AGCTTCCAGCAAGGCTCG	172	
0 actin	Forward	TGACGTGGACATCCGCAAAG CTGGAAGGTGGACAGCGAGG		
β-actin	Reverse			

Flow cytometry assay measurement for cell cycle

CD34+CD38– LSCs were harvested and washed twice in PBS. CD34+CD38– LSCs were fixed overnight using 70% ice-cold ethanol (Beyotime Biotech., Shanghai, China), and washed twice using PBS. Then, the CD34+CD38– LSCs were treated using bovine pancreatic ribonuclease (Sigma-Aldrich, St. Louis, MO, USA) at a dose of 1 mg/ml for 30 min at 37°C. The cells were stained with the propidium iodide (PI) at a concentration of 50 µg/ml in the dark for 30 min. The stained CD34+CD38– LSCs were viewed and imaged using the FACS Aria IIU flow cytometer (BD Bioscience, Oxford, UK) and the percentages of cells in each phase of the cell cycle were analyzed using MultiCycle software (Phoenix Flow Systems, Tokyo, Japan).

Colony formation assay

The colony formation assay was performed, as previously described, but with some modifications [19]. Briefly, the CD34+CD38-LSCs were adjusted to the density of 5×10³ cells/well in 96-well plates containing IMDM, and incubated with 1×10⁻⁴ mol/l 2-mercaptoethanol (Sigma-Aldrich, St. Louis, MO, USA), 3% L-glutamine (Sigma-Aldrich, St. Louis, MO, USA), interleukin 3 (IL-3) (Sigma-Aldrich), 0.8% methylcellulose (Sigma-Aldrich, St. Louis, MO, USA), recombinant human granulocyte macrophage colony-stimulating factor (rhGM-CSF), and human erythropoietin (rhEPO) (Beijing Kylin Culture Co. Ltd. Shanghai, China). The treated CD34+CD38-LSCs were cultured at 37°C and 5% CO, for two weeks. The colony formation units (CFU) consisting of 50 or more CD34+CD38-LSCs were assigned to the mixed colony forming unit (CFU-Mix) assay, which were counted using a BX51 light microscope (Olympus, Tokyo, Japan).

Senescence-associated beta-galactosidase (SA- β -Gal) staining

CD34+CD38– LSCs were stained with the SA- β -Gal detection kit (Cell Signaling Technology Inc., Beverly, MA, USA), according to the manufacturer's instructions. Briefly, CD34+CD38– LSCs were washed twice with PBS and stained with the staining

solution for 12 h at 37°C. The CD34+CD38– LSCs were centrifuged, and cytospin slides were prepared. The cytospin preparations were sealed with the 70% glycerol (Sigma-Aldrich, St. Louis, MO, USA). A total of 400 CD34+CD38– LSCs on the slides were randomly selected, and the positively-stained cells were counted.

Quantitative reverse transcription-polymerase chain reaction (qRT-PCR)

The total RNAs of CD34+CD38-LSCs were extracted with TRIzol regent (Beyotime Biotech., Shanghai, China). The complementary DNA (cDNA) was synthesized using a Reverse Transcription kit (Cat. No. 4366596) (Pierce Thermo Scientific, Rockford, IL, USA) and amplified using SYBR green I (Takara, Dalian, China). The gRT-PCR assay was conducted using 7900HT real-time PCR equipment (ABI, Foster City, CA, USA). The PCR conditions were as follows: initiation for 4 min at 94°C; amplification for 35 cycles at 94°C for 20 s; amplification at 60°C for 30 s; amplification at 72°C for 30 s; and termination at 72°C for 10 min. The qRT-PCR primers for SIRT1, TSC2, and β -actin are shown in Table 1. The amplified products of gRT-PCR were loaded onto 1.5% agarose gels and analyzed with the Quantity-One 1D gel imaging software (Bio-Rad Laboratories, Hercules, CA, USA). Relative gene expression was normalized to the expression of the β -actin gene using the 2^{- $\Delta\Delta CT$} method [20].

Western blot

The CD34+CD38–LSCs cells were lysed using protein lysis buffer (Applygen Tech. Inc., Beijing, China). The protein concentration of the lysates was evaluated using the bicinchoninic acid (BCA) protein concentration detection kit (Cat. No. P0012) (Beyotime Biotech., Shanghai, China). The lysates were separated with 12% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) (Sigma-Aldrich, St. Louis, MO, USA) and transferred onto polyvinylidene fluoride (PVDF) membranes (Beyotime Biotech., Shanghai, China). The PVDF membranes were incubated overnight at 4°C in the primary antibodies, which were all purchased from Abcam Biotech (Cambridge, MA, USA). The primary antibodies included rabbit anti-human

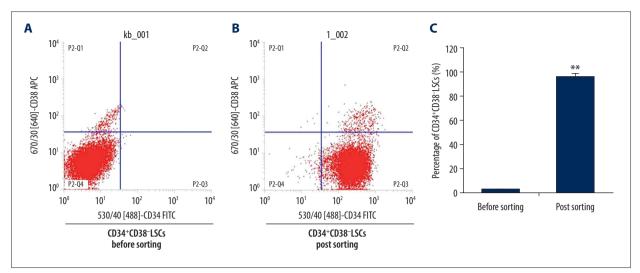


Figure 1. Cell sorting of the CD34+CD38– leukemia stem cells (LSCs) derived from KG1α human acute myeloid leukemia (AML) cells.
 (A) Flow cytometry of CD34+CD38– LSCs derived from KG1α human acute myeloid leukemia cells before cell sorting. (B) Flow cytometry of CD34+CD38– LSCs following cell sorting. (C) Statistical analysis of the sorted CD34+CD38– LSCs. * p<0.05 vs. the control group.

SIRT1 monoclonal antibody (Cat. No. ab32441) (1: 2000), rabbit anti-human TSC2 monoclonal antibody (Cat. No: ab190349) (1: 2000), rabbit anti-human polyclonal antibody (Cat. No. ab8227) (1: 2000). The membranes then incubated in the secondary horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (Cat. No. ab6721) (1: 1000) (Abcam Biotech., Cambridge, MA, USA) at 37°C for 1 h. Western blot images were visualized using the electrochemiluminescence (ECL) kit (Pierce Thermo Scientific, Rockford, IL, USA).

Statistical analysis

Data were expressed as the mean±standard deviation (SD) and analyzed using SPSS version 11.0 software (SPSS Inc., Chicago, III, USA). Experiments were performed in triplicate. One-way analysis of variance (ANOVA) was validated by Tukey's post hoc test to compare data between groups. A P-value <0.05 was considered to be statistically significant.

Results

Isolation and survival rate of cell sorting for CD34+CD38leukemia stem cells (LSCs)

The flow cytometry results showed that following cell sorting, the purity of CD34+CD38– leukemia stem cells (LSCs) was significantly greater compared with the cells before cell sorting (95.86% vs. 2.36%) (Figure 1) (p<0.05). The survival rate of the sorted CD34+CD38– LSCs was 98.72%. The findings demonstrated the successful isolation of CD34+CD38– LSCs.

Ginsenoside Rg1 (Rg1) reduced the proliferation rate of CD34+CD38- LSCs

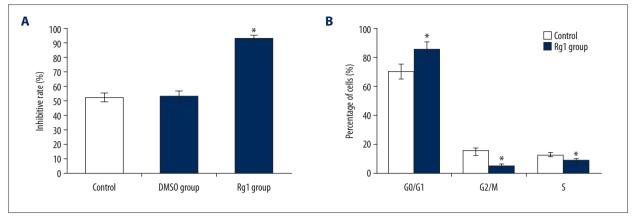
The cell-counting kit-8 (CCK-8) assay was performed to determine the effects of Rg1 on the proliferation of CD34+CD38–LSCs. Rg1 treatment significantly inhibited the CD34+CD38–LSC proliferation compared with the control group (Figure 2A) (p<0.05). There were no significant differences in the proliferation rates of CD34+CD38–LSCs between the control group and the DMSO group, which indicated that DMSO was safe and had no significant cell toxicity.

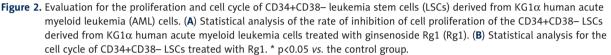
Rg1 modulated the phases of the cell cycle in CD34+CD38- LSCs

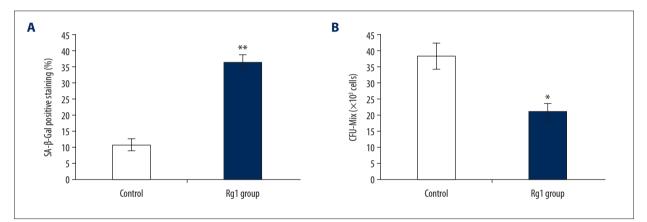
Cell cycle analysis showed that CD34+CD38–LSCs in the G0/G1 phase of the cell cycle were significantly increased, and cells in the G2/M and S phases were significantly reduced compared with that of the control group (Figure 2B) (p<0.05).

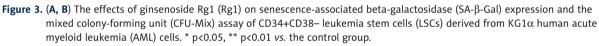
Rg1 increased the expression of senescence-associated beta-galactosidase (SA- β -Gal) in CD34+CD38- LSCs

Previous studies have shown that measurement of the expression of SA- β -Gal and the mixed colony-forming unit (CFU-Mix) assay are biomarkers of cell senescence [21,22]. Therefore, in this study, the levels of SA- β -Gal and CFU-Mix in CD34+CD38– LSCs were evaluated. Rg1 treatment significantly increased the levels of SA- β -Gal compared with the control group (Figure 3A) (p<0.05). Also, the CFU-Mix formation was significantly lower in the Rg1 group compared with the control group (Figure 3B) (p<0.05).









Rg1 down-regulated expression of sirtuin 1 (SIRT1) in CD34+CD38- LSCs

In this study, the expression of SIRT1 mRNA and protein were determined using quantitative reverse transcription-polymerase chain reaction (qRT-PCR) and Western blot, respectively. The qRT-PCR findings showed that the SIRT1 mRNA levels in the Rg1 group were significantly lower compared with the control group (Figure 4A) (p<0.05). Western blot showed that Rg1 treatment significantly down-regulated SIRT1 expression compared with the control group (Figure 4B, 4C) (p<0.05).

Rg1 down-regulated the expression of tuberous sclerosis complex 2 (TSC2) in CD34+CD38- LSCs

TSC2 is a downstream molecule in the SIRT1 pathway. The expression of TSC2 mRNA and protein were determined using quantitative reverse transcription-polymerase chain

reaction (qRT-PCR) and Western blot, respectively. The qRT-PCR findings showed that the TSC2 mRNA levels in the Rg1 group were significantly lower compared with the control group (Figure 5A) (p<0.05). Western blot showed that Rg1 treatment significantly down-regulated TSC2 expression compared with the control group (Figure 5B, 5C) (p<0.05).

The downstream molecule of SIRT1, TSC2, in the CD34+CD38–LSCs was also examined using qRT-PCR (Figure 5A) and Western blot (Figure 5B). The results showed that the Rg1 treatment significantly decreased the mRNA (Figure 5A) and protein expression (Figure 5B, 5C) of TSC2 compared with that in the control group (p<0.05).

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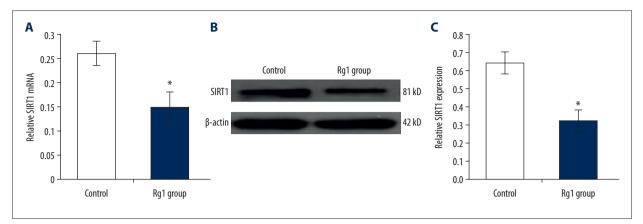


Figure 4. The effects of ginsenoside Rg1 (Rg1) on SIRT1 expression in CD34+CD38– leukemia stem cells (LSCs) derived from KG1α human acute myeloid (AML) leukemia cells. (A) The evaluation for mRNA expression of SIRT1 following treatment with ginsenoside Rg1 (Rg1) using the quantitative reverse transcription-polymerase chain reaction (qRT-PCR). (B) The evaluation of sirtuin 1 (SIRT1) protein expression following treatment with Rg1 using Western blot. (C) Statistical analysis of SIRT1 expression. * p<0.05 vs. the control group.</p>

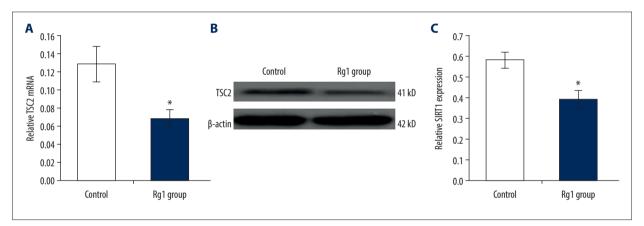


Figure 5. The effects of ginsenoside Rg1 (Rg1) on TSC2 expression in CD34+CD38– leukemia stem cells (LSCs) derived from KG1α human acute myeloid (AML) leukemia cells. (A) The evaluation for mRNA expression of TSC2 following treatment with ginsenoside Rg1 (Rg1) using the quantitative reverse transcription-polymerase chain reaction (qRT-PCR). (B) The evaluation of tuberous sclerosis complex 2 (TSC2) protein expression following treatment with Rg1 using Western blot. (C) Statistical analysis of TSC2 expression. * p<0.05 vs. the control group.</p>

Discussion

Leukemia stem cells (LSCs) play crucial roles in initiating the occurrence and development of acute myeloid leukemia (AML), and are characterized by insensitivity to chemotherapy [24]. Therefore, LSCs have an important role in multidrug resistance and clinical relapse of AML [25]. LSCs also play critical roles in metastasis and treatment failure in leukemia [26,27]. Therefore, traditional chemotherapy may not treat AML effectively. Studies have shown that for patients who have refractory AML, allogeneic stem cell transplantation may be the only treatment option that is effective [26,27]. Therefore, targeted control of LSCs may be a future treatment strategy in patients with AML.

Cell senescence is inevitable, and LSCs also undergo senescence [28]. Recent studies investigated the role of increasing the rate of senescence of cancer cells or cancer stem cells. The traditional Chinese medicine theory has recommended the use of ginseng for its anti-aging, anti-apoptosis, anti-oxidant, and immune-enhancing properties [15,29]. Previous studies have reported that the ginsenoside Rg1 (Rg1) promoted the proliferation of hematopoietic progenitor cells (HPCs) and hematopoietic stem cells (HSCs) and reduced the longevity of mice by reducing DNA damage and down-regulating the expression of p16INK4a and p21CIP1/WAF1 [30,31]. Rg1 was previously shown to inhibit the proliferation of the leukemia cell, K562, by inducing cell senescence [32]. The findings from these previous studies indicated that Rg1 had dual regulatory effects on senescence in different cells.

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In the present study, the Rg1 was administrated to CD34+CD38– LSCs, with the typical LSC phenotype [18,19,33]. The findings showed that Rg1 reduced the proliferation rate of CD34+CD38– LSCs, which was consistent with the findings from a previous study [34]. Also, the results of the evaluation of the cell cycle showed that Rg1 triggered G1 phase cell cycle arrest, which suggests that the effects of Rg1 on inhibition of cell proliferation of CD34+CD38– LSCs were induced by G1 phase arrest.

A previous study reported that the senescence of tumor cells was associated with suppressed cell proliferation [35]. Therefore, the effects of Rg1 administration on the expression of senescence-associated beta-galactosidase (SA- β -Gal) by CD34+CD38–LSCs were studied [21,22]. The findings showed that Rg1 treatment significantly increased SA- β -Gal levels and significantly reduced the formation of the mixed colony-forming unit (CFU-Mix), which suggests that Rg1 was associated with changes of cell senescence of CD34+CD38–LSCs by reducing the renewal and proliferative ability of cells. These results support the previous findings on the effects of Rg1 on the K562 human leukemia cell line [36,37], but to our knowledge, these findings are the first on the effects of Rg1 on senescence of CD34+CD38–LSCs.

Sirtuin 1 (SIRT1) has a critical role in several molecular processes, including cell senescence, cell proliferation, and inflammation [38,39]. A previous study reported that the SIRT1 over-expresses in patients with AML was associated with

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treatment-resistance [40]. The effects of SIRT1 occur by its interaction with tuberous sclerosis complex 2 (TSC2) in the leukemia cells [41]. Therefore, in the present study, the expression of both SIRT1 and TSC2 in the CD34+CD38– LSCs treated with Rg1 was investigated. The findings showed that Rg1 down-regulated the expression of both SIRT1 and TSC2 in the CD34+CD38– LSCs. These results suggest that the Rg1 mediated senescence of CD34+CD38– LSCs is triggered through the down-regulation of the SIRT1/TSC2 signaling pathway.

Conclusions

This study aimed to investigate the effects of the ginseng derivative, ginsenoside Rg1 (Rg1), on CD34+CD38– leukemia stem cells (LSCs) derived from KG1 α human acute myeloid leukemia (AML) cells. The findings showed that Rg1 inhibited cell proliferation and induced the G1 cell cycle arrest of the CD34+CD38– LSCs. Rg1 significantly enhanced the expression of senescence-associated beta-galactosidase (SA- β -Gal) and reduced the mixed colony-forming unit (CFU-Mix), which were indicators of cell senescence, by activating the SIRT1/TSC2 signaling pathway. Future studies should be undertaken to investigate the effects of Rg1 on LSCs both *in vitro* and in *vivo*.

Conflict of interest

None.

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