Deguelin Induces the Apoptosis of Lung Squamous Cell Carcinoma Cells through Regulating the Expression of Galectin-1

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Abstract

Lung cancer is the leading cause of cancer mortality around the world. Despite advances in the targeted therapy, patients with lung squamous cell carcinoma (SCC) still benefit few from it, and the search for potential effective therapies is imperative. Here, we demonstrated that deguelin induced significant apoptosis of lung SCC cells in vitro. Importantly, we found deguelin down-regulated the expression of galectin-1, which was involved in a wide range of tumorous physiologic process. Thus, we both over-expressed and down-regulated galectin-1 to perform its role in deguelin-induced apoptosis. We found that increased galectin-1 attenuated apoptosis of SCC cells exposed to deguelin, while galectin-1 knockdown sensitized lung cancer cells to deguelin treatment. Additionally, we observed that down-regulation of galectin-1 resulted in suppression of Ras/Raf/ERK pathway which was involved in deguelin-induced cell apoptosis. We also found that deguelin had a significant anti-tumor ability with decline of galectin-1 in vivo. In conclusion, these findings confirm that deguelin may act as a new chemo-preventive agent through inducing apoptosis of lung SCC cells in a galectin-1 dependent manner.

Key words: Deguelin; Lung squamous cell carcinoma; Apoptosis; Galectin-1

Introduction

Lung cancer represents one of the most frequently diagnosed cancers worldwide. The data from global cancer statistics indicate that lung cancer is the leading cause of cancer-associated death among males, and has surpassed breast cancer as the leading cause of female cancer mortality in many developed countries[1]. Approximate 80%-85% of lung cancers are categorized as non-small-cell lung cancer (NSCLC), which contains three main subtypes including adenocarcinoma, SCC and large cell carcinoma[2]. Although intensive efforts are devoted to improving lung cancer treatment, it still remains a worldwide dilemma for clinical therapy. Targeted therapies such as EGFR-TKIs have been widely applied in patients with adenocarcinoma, however, patients with lung SCC, which account for 30% of NSCLC, benefit few from the treatment. Therefore, potential alternative therapies for lung SCC are encouraging.

Deguelin, a rotenoid extracted from Derris trifoliata Lour. or Mundulea sericea, has been reported to be effective in inhibiting the proliferation, metastasis and invasion of various types of tumors[3-6]. Previous studies have shown that deguelin can induce the apoptosis of cancer cells by targeting the NF-κB pathway[7-9] and PI3K/Akt pathway[10, 11]. In addition, deguelin has been demonstrated to mediate its chemopreventive effects through targeting cell cycle arrest[12, 13] and anti-angiogenesis[13, 14]. Although the application of deguelin has been explored in several tumors[15-17], the effects of deguelin on lung SCC and the cellular
mechanisms are still not fully studied.

Recently, Hung-Sheng Shang et al. reported that deguelin could decrease the Ras protein level in human osteosarcoma cells [5]. H-Ras belongs to the RAS family of GTPases and contributes to numerous cellular functions including control of cell apoptosis and proliferation [18]. Several reports have shown that H-Ras biological activity requires membrane anchorage which can be strengthened by galectin-1 [19], a β-galactoside-binding protein, which had been reported to be highly secreted in lots of malignancies and was widely involved in tumorous physiologic processes such as cell proliferation, adhesion, migration and apoptosis [20-24]. Increasing clinical and preclinical evidences have confirmed that galectin-1 is correlated with poor prognosis in a variety of tumors [25-27], and galectin-1 inhibition may directly lead to anti-proliferation in cancer cells, indicating that galectin-1 may be a promising drug target for cancer treatment [28]. Although various possible mechanisms related to degelin-induced apoptosis have been proposed, yet the function of galectin-1 in deguelin-treated cancer cells remains to be elucidated.

In the present study, we focused on the potential effects of deguelin on human lung SCC and identified that deguelin can be an apoptosis inducer for lung SCC cells. It is noteworthy that we found deguelin reduced the expression of galectin-1. For further study, we knocked down and overexpressed galectin-1, respectively, and found the expression of galectin-1 was related to apoptotic effect of deguelin by regulating the Ras/Raf/ERK pathway. These results indicated that deguelin could induce apoptosis of lung SCC through regulating expression of galectin-1. Our preliminary studies innovatively explore the role of galectin-1 in deguelin-induced tumor cells apoptosis, and contribute to better understanding with the effect of deguelin on lung SCC cells.

Materials and Methods

Cell Lines and Reagents

Human lung SCC cell SK-MES-1 was purchased from the Committee on Type Culture Collection of Chinese Academy of Sciences (Shanghai, China). NCI-H520 was a kind gift from Dr. Ying (Department of Respiratory Diseases, Sir Run Run Shaw Hospital, Zhejiang University, Hangzhou, China). Both cell lines were maintained in RPMI-1640 (Gibco BRL Co.Ltd., MD, USA), supplemented with 10% fetal bovine serum (FBS, Gibco BRL Co.Ltd., MD, USA), 100U/ml penicillin and 100μg/ml streptomycin (Gibco BRL Co.Ltd., MD, USA) at 37°C with 5% CO2. Deguelin was obtained from Sigma (St. Louis, MO, USA). Galectin-1 shRNA plasmid, negative control shRNA plasmid, shRNA plasmid transfection reagent and shRNA plasmid transfection medium were purchased from Santa Cruz Biotechnology (CA, USA). Galectin-1 expression plasmid pCMV3-SP-N-His-Gal-1 and empty vector pCMV3-SP-N-His plasmid were from Sino Biological Inc. (Beijing, China). OPTI-MEM was purchased from Gibco BRL Co.Ltd. (MD, USA) and HiPerFect transfection reagent was obtained from Qiagen (Hilden, Germany). Antibodies against rabbit Galectin-1, H-Ras, Raf-1, p-Raf-1 and Raf kinase inhibitor L779450 were purchased from Abcam (Cambridge, MA, USA). GAPDH, Bax, Bcl-2, cleaved caspase-3, cleaved caspase-9, phospho-p44/42 MAPK(T202/Y204), p44/42 MAPK antibodies were purchased from Cell Signaling Technology (Danvers, MA, USA).

Cell Viability Assay

Cells were seeded in a 96-well plate at a density of 4x10^3/well. Different concentrations of deguelin were added into each well, and incubated for 12, 24, 48 and 72 hours, respectively. After incubation, WST-8 (Dojindo Laboratories, Tokyo, Japan) was added into each well. The absorbance of solutions was measured spectrophotometrically at 450nm with an automatic microplate analyzer after incubation for an additional two hours.

Cell Apoptosis Assay

With deguelin treatment for 24 hours, cells were harvested and detected for apoptosis by using FITC Annexin V Apoptosis Detection Kit (BD Biosciences, NJ, USA) according to the manufacture’s protocols. Briefly, cells were washed twice with cold PBS and resuspended in binding buffer at a concentration of 1x10^5 cells/ml. Then, cells were stained with 5ul of FITC Annexin V and 5ul of Propidium Iodide followed by incubation for 15 minutes in dark at room temperature. Finally, cells apoptosis were analyzed by BD FACSVerse (BD Biosciences, NJ, USA).

Quantitative real-time PCR

Total cellular RNA was extracted by using RNeasy Mini Kit (Qiagen, Hilden, Germany) in accordance to the manufacturer’s protocol and the cellular RNA concentration was measured by NanoDrop 2000 (Thermo Scientific, Waltham, MA, USA). The reverse transcription into complementary DNA was performed by using the PrimeScript 1st Strand cDNA Synthesis Kit (Takara, Dalian, China), and for quantitative RT-PCR analysis, cDNA amplification was performed by QuantiFast SYBR Green PCR Kit (Qiagen, Hilden, Germany). The comparative Ct method (2^-ΔΔCt) was used to analyze.
data. The specific primers of RT-PCR are shown in Table 1.

| mRNA       | Oligonucleotides (5’ to 3’) |
|------------|-----------------------------|
| Bax-F      | 5-CCCCGAGGCTTGTTCGGAG-3     |
| Bax-R      | 5-CCAGG gagAACCTGTTGCTG-3   |
| Bcl-2-F    | 5-GTGAGGGTGATGGTGAGGG-3     |
| Bcl-2-R    | 5-GTGGTCAGGTACTCGTCAATCC-3  |
| Galectin-1-F | 5-TGGCGCAAACCTGATCTCTC-3  |
| Galectin-1-R | 5-GGACGAGATCCCTCCAAAT-3   |
| GAPDH-F    | 5-GGCTGTTGTCATACTTCTGAG-3   |
| GAPDH-R    | 5-GGCTGTTGTCATACTTCTGAG-3   |

**Immunoblotting**

Cells were lysed with lysis buffer, the lysate supernatants were harvested in 1.5ml microcentrifuge tubes and boiled for 10 minutes in loading buffer. Protein concentration was measured by using Pierce BCA Protein Assay Kit (Thermo Scientific, Waltham, MA, USA) according to the manufacture’s instructions. Cell lysates were subjected to SDS-PAGE, and then transferred to polyvinylidene difluoride (PVDF) membrane (Millipore, Bedford, MA, USA) for immunoblotting analysis. After blocking with 5% nonfat milk for 1 hour, membranes were incubated with the appropriate primary antibodies overnight at 4°C followed by incubation with the corresponding secondary antibody for 2 hours at room temperature. Immunoreactive bands were visualized using the enhanced chemiluminescence (ECL) detection system.

**Cell Transfection**

NCI-H520 and SK-MES-1 cells were seeded in six-well plates before transfection. In order to down-regulate galectin-1 expression, NCI-H520 and SK-MES-1 cells were transfected with galectin-1 shRNA or negative control shRNA using shRNA plasmid transfection reagent according to the procedures provided by Santa Cruz Biotechnology (CA, USA). For galectin-1 overexpression, 0.7μg pCMV3-SP-N-His-Gal-1 plasmid or negative control pCMV3-SP-N-His plasmid and 20μl HiPerFect transfection reagent were diluted separately in 40μl OPTI-MEM at room temperature for 5 minutes. The two mixtures were re-mixed and incubated for 30 minutes. Finally, the complex was added to the wells along with 2ml of medium. After incubation for another 48 hours, western blotting was performed to test the expression level of galectin-1 to confirm transfection efficiency. In order to obtain stable transfected cell lines, the positive transfectants were selected and expanded for further study.

**Colony Formation Assay**

Cells transfected with shGal-1, negative control shRNA, galectin-1 plasmid vector and empty plasmid vector were counted and seeded in six-well plates at a density of 400 cells/well. After incubation of 24 hours, the cells were treated with 10μM deguelin. Meanwhile, the culture medium was refreshed every three days. After 14 days, the cells were stained with crystal violet, and the number of colonies was counted only if they contained more than 50 cells. The following equation: (number of colonies/ number of seeded cells) ×100% was performed for the rate of colony formation[29].

**Co-immunoprecipitation Assay**

Mock- or deguelin-treated cells were washed twice with cold PBS and lysed with NP-40 lysis buffer (50mM Tris-HCl pH 7.4, 150 mM NaCl, 1 mM EDTA, 1% NP-40 and 5% glycerol) with 1mM PMSF for 30 minutes. The lysate was centrifuged at 12000 rpm at 4°C for 10 minutes and the supernatants were incubated with primary antibodies or non-specific IgG as a negative control at 4°C overnight, then the Protein A+G Sepharose beads were applied and rotated for 3 hours at 4°C. The immunocomplexes were washed for 5 times with lysis buffer followed by boiling with SDS-PAGE buffer and then analyzed by western blotting.

**Animal Experiments**

Six-week-old female BALB/c-nude mice obtained from Shanghai Experimental Animal Center (Chinese Academy of Sciences, Shanghai, China) were used for human tumor xenograft models. After one week’s acclimatization, they were randomized and allocated into groups of six mice. The control group was injected subcutaneously into the left armpit with 3.5×10^6 NCI-H520 cells per mouse. Other groups were injected with an equal number of NCI-H520 cells, NCI-H520 cells transfected with galectin-1 plasmid vector or NCI-H520 cells transfected with empty plasmid vector, respectively. When palpable tumors (~200–300mm³) arose, the control group was orally treated with physiological saline, while other groups were treated with deguelin (4 mg/kg) by oral gavage on days 1, 3 and 5 of each week for three weeks. Tumor size was measured by caliper through measurements of the two perpendicular diameters every three days using the formula: Volume = (width^2 × length)/2. All procedures were performed according to the Regulations for the Administration of Affairs Concerning Experimental Animals. The experiments were approved by the State Council of the People’s Republic of China and the Experimental Animal Ethics Committee of Zhejiang University.
Statistical Analysis

All data are presented as mean ± SD of three independent experiments. SPSS 19.0 (SPSS Inc., Chicago, IL, USA) were used for statistical analysis. Data were analyzed using the Student’s t-test, and significance was chosen as P values < 0.05.

Results

Deguelin induced the apoptosis of human lung SCC cells

In two subpopulations of human lung SCC cells, NCI-H520 and SK-MES-1, deguelin administration greatly inhibited the cell proliferation in both dose- (Fig. 1A) and time-dependent (Fig. 1B and 1C) manners. Flow cytometry was applied to identify the cellular apoptosis with deguelin exposure, as shown in Fig. 2, a significant cell apoptosis was observed in NCI-H520 cells, similar result was obtained from SK-MES-1 (data not shown). Next, real-time PCR analysis indicated that deguelin down-regulated the expression of anti-apoptotic factor Bcl-2 (Fig. 3A and 3C), while apoptotic factor Bax was up-regulated (Fig. 3B and 3D) in mRNA level. Furthermore, consistent with the RT-PCR results, similar results were shown by immunoblotting that deguelin exposure increased Bax activity while caused caspase-3 and caspase-9 cleaved and down-regulated Bcl-2 expression (Fig. 3E and 3F). Thus, these results indicated that deguelin was able to induce the apoptosis of lung SCC cells.

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Figure 1. Deguelin inhibited the lung SCC cells proliferation in both dose-dependent and time-dependent manners. (A) NCI-H520 and SK-MES-1 cells were incubated with 0.1-100 μM deguelin for 72 hours. (B) NCI-H520 cells were exposed to deguelin (25 μM) for different time points (12, 24, 48 and 72 hours). (C) SK-MES-1 cells were exposed to deguelin (25 μM) for different time points (12, 24, 48 and 72 hours). Cell viability was measured using WST-8 assays. Each experiment was performed in triplicate, thrice independently. The data are presented as mean±SD. *P<0.05, **P<0.01 vs Control.

Figure 2. Deguelin induced the apoptosis of human lung SCC cells. (A) NCI-H520 cells were treated with indicated concentrations of deguelin for 24 hours. Flow cytometry was used for analysis of cell apoptosis. (B) Apoptosis percentages of NCI-H520 cells from three independent experiments are presented as mean±SD. *P<0.05, **P<0.01 vs Control.
Deguelin exposure down-regulated the expression of anti-apoptotic factors while apoptotic factors were up-regulated. (A, B) NCI-H520 cells were treated with various concentrations of deguelin for 24 hours. Total cellular RNA was extracted and analysed by real-time PCR. (C, D) SK-MES-1 cells were treated with various concentrations of deguelin for 24 hours. Total cellular RNA was extracted and analysed by real-time PCR. The data from three independent experiments are presented as mean±SD. *P<0.05, **P<0.01 vs Control. (E, F) NCI-H520 and SK-MES-1 cells were incubated with different concentrations of deguelin for 24 hours, cell lysates were harvested and the indicated proteins were determined by western blotting.

Deguelin down-regulated the expression of galectin-1 in lung SCC cells

To determine galectin-1 expression in lung SCC cell lines treated with deguelin, we performed western blotting results as shown in Fig. 4C and 4D that deguelin dose-dependently application could cause galectin-1 degradation. We also analyzed the relative galectin-1 mRNA expression by RT-PCR and the mRNA levels of galectin-1 were prominently down-regulated by deguelin in different doses as shown in Fig. 4A and 4B, which were consist with the immunoblotting results. In summary, these results indicated that deguelin significantly degraded galectin-1 expression at both protein and mRNA levels in lung SCC cells.

Deguelin induced apoptosis of lung SCC cells in a galectin-1 dependent manner

To confirm whether galectin-1 expression level contributed to the apoptotic effect induced by deguelin, we down-regulated galectin-1 expression by transfecting Gal-1 shRNA or negative control shRNA. Meanwhile, NCI-H520 and SK-MES-1 cell lines were also transfected with human galectin-1 plasmid pCMV3-SP-N-His-Gal-1 or empty vector plasmid pCMV3-SP-N-His to increase the expression of galectin-1. Real-time PCR and western blotting were applied to confirm galectin-1 expression at mRNA and protein levels (Fig. 5).
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Figure 4. Deguelin down-regulated the expression of galectin-1 in lung SCC cells. (A, B) NCI-H520 and SK-MES-1 cells were treated with various concentrations of deguelin for 24 hours. Total cellular RNA was extracted and analysed using real-time PCR. The data from three independent experiments are presented as mean±SD. **P<0.01 vs Control. (C, D) NCI-H520 and SK-MES-1 cells were incubated with different concentrations of deguelin for 24 hours, cell lysates were harvested and the indicated proteins were determined by western blotting.

Figure 5. Construction of stable galectin-1 overexpression and shGal-1-knockdown lung SCC cells. (A) pCMV3-SP-N-His-Gal-1 mediated overexpression and shRNA mediated knockdown of galectin-1 were confirmed by real-time PCR (A, B, D, E) and western blotting (C, F), respectively. The data are representatives of three independent experiments and presented as mean±SD. **P<0.01.

To investigate the role of galectin-1 in deguelin treatment, we performed experiments to determine the proliferation, colony formation and apoptosis of cells exposed to 25μM deguelin. We found that galectin-1 overexpression cells displayed significantly increased proliferation (Fig. 6A and 6B) and colony formation (Fig. 6C and 6D) after deguelin treatment. Meanwhile, galectin-1 knockdown inhibited the cell proliferation and colony forming efficiency compared with the negative control (NC) group. Additionally, flow cytometry analysis showed that shGal-1-knockdown NCI-H520 cells had a higher apoptotic rate with deguelin exposure than NC group while the galectin-1 overexpression had lower apoptotic rate than the equivalent control group (Fig. 6E). The similar results of flow cytometry and colony formation were also observed in SK-MES-1 cell line (data not shown). Next, we performed western blotting to determine the effects of galectin-1 knockdown on apoptosis-related proteins after
deguelin treatment. We observed that both in shGal-1-knockdown NCI-H520 and SK-MES-1 cell lines, suppression of galectin-1 was coupled with decreased Bcl-2 and increased Bax, cleaved caspase-3 and cleaved caspase-9 activities. Furthermore, compared with galectin-1 knockdown, we observed that galectin-1 overexpression had contrary results as expected (Fig. 6F), which suggested that deguelin induced lung SCC cells apoptosis via galectin-1 regulation.

Suppression of galectin-1 resulted in deactivation of Ras/Raf/ERK pathway

Previously we had demonstrated that galectin-1 was involved in deguelin-induced apoptosis process, and we further explored the galectin-1-related cellular mechanisms. It has been reported that galectin-1 is a selective binding partner of H-Ras[19] and mediates a variety of biological functions through direct interaction with H-Ras. To confirm this interaction, co-immunoprecipitation assay was performed and the result indicated that galectin-1 interacted with H-Ras directly in lung SCC cells (Fig. 7A).

Figure 6. Deguelin induced apoptosis of lung SCC cells in a galectin-1 dependent manner. (A, B) Both galectin-1 overexpression and knockdown cells were incubated with 25μM deguelin for 72 hours and then measured cell viability using WST-8 assays. (C, D) Galectin-1 overexpression or knockdown cells were seeded in six-well plates. After 24 hours of incubation, the cells were treated with 10μM deguelin for 14 days. Then the colony formation efficiencies were summarized. (E) Galectin-1 overexpression or knockdown cells were incubated with 25μM deguelin for 24 hours. Flow cytometry was used for cell apoptosis analysis. (F) Galectin-1 overexpression or knockdown cells were incubated with 25μM deguelin for 24 hours. Cell lysates were harvested and the indicated proteins were determined by western blotting. All data are from three independent experiments and presented as mean±SD. *P<0.05, **P<0.01.
Figure 7. Suppression of galectin-1 resulted in deactivation of the Ras/Raf/ERK pathway with deguelin treatment. (A) Mock- or deguelin-treated cells were lysed for co-immunoprecipitation and then analyzed by western blotting. (B) Galectin-1 overexpression and shGal-1-knockdown cells were incubated with 25μM deguelin for 24 hours, respectively. Cell lysates were harvested and the indicated proteins were determined by western blotting. (C) Galectin-1 overexpression cells with or without L779450 (10μM) were incubated with 25μM deguelin for 24 hours. Flow cytometry was used for cell apoptosis analysis. The data are representatives of three independent experiments and presented as mean±SD. **P<0.01. (D) Galectin-1 overexpression cells with or without L779450 (10μM) were incubated with 25μM deguelin for 24 hours. Then cell lysates were harvested for western blotting.

NCI-H520 and SK-MES-1 cells transfected with Galectin-1 shRNA, negative control shRNA, galectin-1 plasmid vector and empty plasmid vector, respectively, were treated with 25μM deguelin for 24 hours, then cells were harvested and analysed by western blotting. We found that galectin-1 knockdown led to decreased expression of H-Ras, p-Raf-1 and p-ERK1/2, while galectin-1 overexpression resulted in the activation of Ras/Raf/ERK pathway (Fig. 7B).

In order to explore the role of Ras/Raf/ERK pathway in deguelin-induced tumor apoptosis, we then apply a Raf kinase inhibitor L779450[30] to block the Ras/Raf/ERK pathway in galectin-1 overexpression cells before deguelin administration. As shown in Fig. 7C, flow cytometry was applied and we observed a significant cell apoptosis in L779450 treatment group. Next, we performed western blotting to detect the expression of apoptosis-related proteins. Bax, cleaved caspase-3 and cleaved caspase-9 expression were up-regulated, while Bcl-2 activity was decreased in deguelin-treated galectin-1 overexpression cells when Ras/Raf/ERK pathway was blocked (Fig. 7D). These results indicated that the Ras/Raf/ERK pathway was involved in the deguelin-induced cells apoptosis.

Anti-tumor effect of deguelin in NCI-H520 xenograft nude mice model

To determine whether the anti-tumor effect of deguelin observed in vitro was consistent with that in vivo, we constructed NCI-H520 xenograft models as mentioned in materials and methods. The NCI-H520 xenograft nude mice group receiving oral deguelin treatment showed a markedly inhibition in tumor growth, by performance of the reduced tumor size compared with control group receiving physiological saline (Fig. 8A and 8B). Meanwhile, tumors of the group which had been injected with galectin-1 overexpression cells, grew significantly faster than the equivalent group with control plasmid (Fig. 8A and 8B). We then analyzed whether deguelin inhibited the expression of galectin-1 in vivo by using immunoblotting. We found that in the deguelin-treated NCI-H520 xenograft group, galectin-1 was suppressed (Fig. 8C), indicating that deguelin had a significant anti-tumor ability and could reduce the expression of galectin-1 in vivo.

Discussion

Lung cancer, with approximately 24% of cancer-related mortality, remains the single deadliest cancer worldwide. Despite continuous efforts devoted to improving lung cancer treatment, there is no significant improvement in the overall five-year survival rate. Moreover, effective clinical therapies targeted EGFR and EML4-AKL which have been widely applied in adenocarcinoma are rarely associated with patients with lung SCC, another main subtype of NSCLC[31]. So it is meaningful to find more effective therapeutic agents against lung SCC.
Deguelin is a nature compound of the flavonoid family products extracted from plants including *Derris trifoliata* Lour. (Leguminosae), *Mundulea sericea* (Leguminosae) and *Tephrosia vogelii* Hook.f. (Leguminosae)[13]. This plant-derived rotenoid has been reported to be an effective cancer chemo-preventive agent suppressing the growth of various types of cancers[12, 32-34]. Previous studies have shown that the possible mechanisms of anti-tumor effect of deguelin may include DNA damage, reducing DNA repair genes, inhibiting vasculogenic function, and blocking anti-apoptotic pathways[13, 14, 35]. Although the application of deguelin on tumors has been paid more attention, researches focused on deguelin treatment for lung SCC are still limited.

In the present study, we found that deguelin could induce the apoptosis of lung SCC cells *in vitro*, which is consistent with the effect of deguelin performed in other cancer cells[36, 37]. In nude mice xenograft models, we also showed a significant anti-tumor ability of deguelin *in vivo*. The tumor sizes of mice receiving oral deguelin administration were significantly reduced compared with those receiving physiological saline treatment. These results indicate that deguelin represents as an effective chemopreventive agent against lung SCC. Importantly, our further study shows that deguelin-induced apoptosis is mediated through the regulation of galectin-1 expression.

Galectin-1, a member of the β-galactoside binding protein family, is encoded by the LGALS1 gene located on chromosome 22q12[38]. It has been reported that the expression of galectin-1 is increased in a wide variety of tumors including breast[39], colon[40], ovarian[41], pancreatic[23], prostate[42] and lung[20, 24] cancers. It is well documented that galectin-1 is involved in multiple processes such as tumor cell proliferation, differentiation, invasiveness and metastasis[38, 43, 44]. Carlini et al.[45] recently reported that NSCLC patients showing high galectin-1 expression were evidenced to have a poorer clinical outcome. Similarly, studies in gastric, colon, breast and epithelial ovarian cancers also showed a positive correlation between galectin-1 expression and poor prognosis[25, 39, 40, 46, 47] underscoring the relevance of galectin-1 as a possible biomarker or therapeutic target in cancers.

Here, we explore the effect of deguelin on galectin-1 and put forward that deguelin can reduce the galectin-1 expression level both *in vivo* and *in vitro*. These results imply deguelin as a potential inhibitor of galectin-1. To confirm the role of galectin-1 in deguelin-induced apoptosis, we silenced and
over-expressed galectin-1 in NCI-H520 and SK-MES-1 cell lines. After exposure to deguelin, we found that galectin-1 knockdown sensitized lung cancer cells to deguelin treatment, while galectin-1 overexpression cells were insensitive to deguelin compared with control cells in vitro. Besides, a similar result was also observed in nude mice xenograft models that the mice group injected with galectin-1 overexpression NCI-H520 cells showed significantly increased tumor volumes compared with mice injected with NCI-H520 control cells. These results suggest that the anti-tumor effect of deguelin on lung SCC cells is performed by down-regulating galectin-1 expression.

It has been reported that galectin-1 can interact with H-Ras and strengthen its membrane anchorage[19]. In agreement with previous studies, we performed co-immunoprecipitation assay and demonstrated a direct interaction between H-Ras and galectin-1 in NCI-H520 cells. Besides, silencing of galectin-1 resulted in down-regulation of H-Ras, p-Raf-1 and p-ERK1/2 in both deguelin-treated NCI-H520 and SK-MES-1 cells, while galectin-1 overexpression could up-regulate H-Ras, p-Raf-1 and p-ERK1/2, indicating that galectin-1 can regulate the Ras/Raf/ERK pathway through the interaction with H-Ras. For further study, we inhibited the activity of Raf-1 to block the Ras/Raf/ERK pathway. We found the apoptosis ratio of galectin-1 overexpression NCI-H520 cells after deguelin administration was significantly increased when the Ras/Raf/ERK pathway was blocked. Thus, we demonstrate that deguelin can induce the lung SCC cells apoptosis by inhibiting the Ras/Raf/ERK pathway through suppressing galectin-1 expression.

Previous studies had shown that the activation of Ras could also stimulate the PI3K signaling pathways, which played an important role in Ras-mediated cellular physiological processes like cell survival and proliferation[48, 49], and deguelin is well known as a PI3K/AKT inhibitor[11, 50]. In the present study, we demonstrated that deguelin could down-regulate H-Ras by inhibiting galectin-1 as mentioned previously. Therefore, we infer that deguelin may also regulate the PI3K/AKT pathway through inhibiting galectin-1 expression, which merits further investigation.

Conclusion

In summary, we have shown evidence that deguelin can inhibit the growth of lung SCC by inducing cell apoptosis. Additionally, we found that overexpression of galectin-1 could reduce the apoptotic effects of deguelin through the up-regulation of Ras/Raf/ERK pathway, while suppression of galectin-1 sensitized lung cancer cells to deguelin treatment, suggesting that deguelin induces the apoptosis of lung SCC cells by inhibiting the Ras/Raf/ERK pathway through suppressing galectin-1 expression. This study proposes that deguelin may represent as a novel and effective agent against lung squamous cell carcinoma.

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Competing Interests

The authors have declared that no competing interest exists.

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