Cryptotanshinone inhibits lung cancer invasion via microRNA-133a/matrix metalloproteinase 14 regulation

HUIJUAN WANG1, YANSHAN ZHANG2, YINGGUO ZHANG2, WENLING LIU2 and JIHONG WANG2

Departments of 1Tumor Chemotherapy and 2Thoracic Surgery, Tumor Hospital of Wuwei, Wuwei, Gansu 733000, P.R. China

Received April 10, 2018; Accepted April 4, 2019

DOI: 10.3892/ol.2019.10580

Abstract. Cryptotanshinone inhibits the proliferative and colony formation abilities of human non-small cell lung cancer cells (NSCLCs). The present study reported that signal transducer and activator of transcription 3 is not the only target of cryptotanshinone during the inhibition of human NSCLCs. It was identified that cryptotanshinone upregulates the expression levels of microRNA (miR)-30d-5p, miR-126-3p, miR-133a, miR-338-3p and miR-451a, and downregulates miR-21-5p, miR-96-5p, miR-182-5p and miR-205-5p. Among these, miR-133a was the most significantly upregulated. miR-133a targets and downregulates the expression of matrix metalloproteinase (MMP)14; however, MMP15, MMP16 and MMP24 were determined to be unaffected. This process was identified to be independent of tissue inhibitor of metallopro-

Introduction

Lung cancer is a malignant lung tumor characterized by uncontrolled lung cell proliferation (1). It is the most commonly diagnosed cancer and the leading cause of cancer-associated mortality (2). Therefore, discovering an effective drug with low toxicity for the treatment of lung cancer is necessary. Human non-small cell lung cancer (NSCLC) cell line, including human A549 lung adenocarcinoma epithelial cell line is widely used as a lung cancer study model (3).

Cryptotanshinone is a natural quinoid diterpene isolated from the roots of a traditional herb Salvia miltiorrhiza, which is considered a potent anticancer agent with anticancer activities (4). In our previous study, it was reported that cryptotanshinone could inhibit the growth of human lung cancer cells and induce apoptosis (5). Signal transducer and activator of transcription 3 (STAT3) is a key transcription activator for cancer cell proliferation. As a STAT3 inhibitor, cryptotanshinone responds to STAT3-induced cytokines and growth factors (6,7); however, to the best of our knowledge, whether STAT3 is the only target of cryptotanshinone remains unclear.

MicroRNAs (miRNAs/miRs) are small non-coding RNA molecules that are 22 nucleotides long and function in RNA silencing and post-transcriptional gene regulation (8). miRNAs serve an important role in lung cancer (9,10). The principal miRNAs involved in lung cancer are members of the let-7 family, including miR-34, miR133, miR-17 and miR-124 (11-13). The present study first investigated if STAT3 was the only target of cryptotanshinone in lung cancer cells. It was identified that even without regulation of STAT3 signaling, cryptotanshinone could inhibit the proliferation and colony formation of lung cancer cells. Based on this result, the effects of cryptotanshinone on miRNAs were investigated using reverse transcription-quantitative polymerase chain reaction (RT-qPCR). In addition, the expression levels of matrix metalloproteinases (MMPs) were studied. Finally, the effects of cryptotanshinone on lung cancer cell invasion were also detected. In summary, the findings of the present study provides further insight into the anticancer effects of cryptotanshinone.

Materials and methods

Chemical treatments and cytotoxic effects assessment. The human A549 lung adenocarcinoma epithelial cell line, one of the NSCLC cell lines, is the most common human lung cancer cell line used for studies regarding lung cancer (1,3). Cell culture was performed following our previous method (5). Briefly, A549 was propagated in RPMI-1640 cell culture medium (Gibco; Thermo Fisher Scientific, Inc.) with L-glutamine, 100 U/ml penicillin, 100 µg/ml streptomycin, and 10% (v/v) fetal bovine serum (FBS; Gibco; Thermo Fisher Scientific, Inc.) at 37°C with 5% CO2. A549 cells (5x103 cells/well) were seeded in a 96-well plate (Corning, Inc.). Cells were treated with cryptotanshinone (LKT Laboratories, Inc.) dissolved in 0.1% dissolved in dimethyl sulfoxide (DMSO), defined as the CT group, at a final concentration of 20 µM (4) for 24,
36 and 48 h. The control group was treated with 0.1% DMSO, defined as the NC group. Colivelin, a STAT3 activator, was purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). The final concentration of colivelin (dissolved in DMSO) was 100 µM and cells treated, at 37°C, with colivelin were termed the CV group, while cells treated with both 20 µM cryptotanshinone and 100 µM colivelin were termed the CT+CV group. Activity of STAT3 in the NC, CT, CV and CT+CV groups was detected by enzyme-linked immunosorbent assay (ELISA) using a STAT3 (Phospho) [pY705] Multispecies ELISA kit (cat. no. KHO0481; Thermo Fisher Scientific Inc.), according to the manufacturer's protocol. Following treatment at 37°C for 24, 48 and 72 h the cytotoxic effects of cryptotanshinone and colivelin in the NC, CT and CT+CV groups were detected using an MTT assay. The optical density (OD) was measured using iMark™ Microplate Absorbance Reader (Bio-Rad Laboratories, Inc.) at 450 nm, representing the activity of STAT3. Colony formation was detected as described previously (4). Briefly, A549 cells (n=250) were seeded in 6-well Corning™ Costar™ Flat Bottom Cell Culture Plates (D=35 mm, Corning, Inc.) and allowed to attach overnight prior to treatment with 20 µM cryptotanshinone dissolved in DMSO or 0.1% DMSO (control), and were incubated for 10 days. The colonies were stained with 0.5% crystal violet in methanol/acetic acid (3:1) and those composed of >50 cells were counted. Experiments were performed three times in duplicate.

RT-qPCR and western blot analysis. A549 cells were treated with 20 µM cryptotanshinone or with 0.1% DMSO at 37°C for 6 h, followed by miRNA detection. miRNA was isolated from the NC, CT and CT+CV groups using an mirVana™ miRNA Isolation kit with phenol (Thermo Fisher Scientific Inc.) and converted to complementary DNA using a TaqMan™ MicroRNA Reverse Transcription kit (cat. no. 4366597; Thermo Fisher Scientific Inc.). miRNA analysis was conducted using a miScript miRNA PCR array (cat. no. MIHS-114Z; Qiagen GmbH, Hilden, Germany) following its instruction. RT-qPCR was performed on Applied Biosystems™ 7500 Fast Real-time PCR. PCR thermocycling conditions were as follows: 50°C 2 min; 95°C 2 min; 95°C 15 sec; 60°C 1 min, 40 cycles. Melt curve stage: 95°C 15 sec; 95°C 2 min; 95°C 15 sec.

Western blot analysis. For western blot analysis, 20 µg protein was extracted using Cell lysis buffer for Western and IP (cat. no. P0013; Beyotime Institute of Biotechnology), determined using BCA kits (cat. no. P0012; Beyotime Institute of Biotechnology) and boiled with NuPAGE™ LDS Sample Buffer (4X) (Invitrogen; Thermo Fisher Scientific, Inc.; cat. no. NP0007) at 100°C for 5 min prior to injection in a 12.5% SDS-PAGE. Following electrophoresis and transfer, the polyvinylidene fluoride (PVDF) membrane (Invitrogen; Thermo Fisher Scientific Inc.) was blocked in 5% non-fat milk for 1 h, incubated overnight at 4°C with primary antibodies as follows: MMP14 (dilution, 1:800; cat. no. ab53712), MMP15 (dilution, 1:500; cat. no. ab15475), MMP16 (dilution, 1:800; cat. no. ab53145), MMP24 (dilution, 1:400; cat. no. ab135564) and tissue inhibitor of metalloproteinases 2 (TIMP2; dilution, 1:1,000; cat. no. ab99707). All primary antibodies were purchased from Abcam (Cambridge, UK) and incubated subsequently with secondary antibody Goat Anti-Rabbit IgG H&L (HRP) (dilution, 1:5,000; cat. no. ab205718) for 1 h at room temperature, washed and exposed. GAPDH (dilution, 1:10,000; cat. no. ab181602) was used as loading control. Membranes were incubated in Pierce™ ECL Western Blotting Substrate for 1 min at room temperature (solution A:solution B=1:1; Thermo Scientific Inc.; cat. no. 32106) and images were captured using LAS-3000 Imaging System (Fuji, Japan). The expression value (target protein/ GAPDH) in the control group was set as 1, while the relative values for the cryptotanshinone group were calculated in association with the control group using ImageJ (National Institutes of Health, Bethesda). All experiments were performed three times with a duplicate each time. Data are compared between the NC and CT groups.

Cancer cell invasion assay. A cancer cell invasion assay was conducted using a Corning® BioCoat™ Matrigel® Invasion Chamber (Corning Inc., Corning, NY, USA), according to the manufacturer's protocol at 37°C. The culture medium was Gibco DMEM, High Glucose, Pyruvate (cat. no. 11-995-040; Gibco; Thermo Fisher Scientific Inc.) with 10% FBS (cat. no. 10437010; Gibco; Thermo Fisher Scientific Inc.) for lower chamber and 0.1% BSA (Gibco; Thermo Fisher Scientific Inc.; cat. no. 11021029) for upper chamber. Briefly, the medium was subsequently rehydrated for 2 h, and fresh culture medium containing 5x10^4 A549 cells/ml was plated in the upper 24-well chambers with 0.1% DMSO (NC group) or 20 µM cryptotanshinone (CT group). The cells were then incubated for 36 h at 37°C with 5% CO₂. Subsequently, the inserts were transferred to 100% methanol for 2 min and then to 1% Toluidine Blue in 1% borax for 2 min at room temperature. Three fields/view were selected randomly and then analyzed. Images were obtained under Nikon TE2000 light microscope (magnification, x100; Nikon Corporation, Japan).

Statistical analysis. Data are presented as the mean ± standard deviation. A Mann Whitney U test was used for comparisons between two groups. One-way analysis of variance followed by a post-hoc Tukey's test was used to compare multiple groups. All statistical analysis was performed using SPSS 19.0 (IBM Corp., Armonk, NY, USA). P<0.05 was considered to indicate a statistically significant difference.

Results

Inhibition of STAT3 activity. As demonstrated in Fig. 1A, the OD value representing the STAT3 activity, was significantly decreased following treatment with cryptotanshinone, while treatment with colivelin significantly increased the OD 450 value compared with the control group. In the combined treatment group, the OD value was significantly increased compared with NC and CT groups, but remained lower than the CV group. These results indicated that cryptotanshinone could serve as a STAT3 inhibitor and colivelin as a STAT3 activator.

Cytotoxic effects of cryptotanshinone and colivelin. As demonstrated in Fig. 1B, following treatment with cryptotanshinone for 24, 48 and 72 h the cellular proliferation of A549 cells
significantly decreased compared with the control. Combined treatment with the STAT3 activator colivelin, was revealed to reduce the cytotoxic effects of cryptotanshinone, however, the cellular proliferation rate remained lower compared with control group. As demonstrated in Fig. 2, compared with NC group, the number and size of colony formation was significantly decreased in the CT group. Combined treatment markedly increased the number and size of colonies. (B) Data analysis for tumor colony formation in the CT, NC and CT+CV groups. *P<0.05, **P<0.01. CT, cryptotanshinone-treated; CT+CV, cryptotanshinone and colivelin-treated; NC, negative control.

Upregulation of miR-133a following treatment with cryptotanshinone. As demonstrated in Fig. 3, the following miRNAs were identified to be upregulated in A549 cells following treatment with cryptotanshinone: miR-30d-5p, miR-126-3p, miR-133a, miR-338-3p and miR-451a. By contrast, the following miRNAs were revealed to be downregulated following treatment with cryptotanshinone: miR-21-5p, miR-96-5p, miR-182-5p and miR-205-5p. These miRNAs target anti-apoptotic or pro-anti-apoptotic genes (14). Among cells. The colony number in NC group was set as 1 and relative values in other groups were calculated by comparison. Data are presented in Fig. 2B.
the microRNAs identified, miR-133a was the most significantly upregulated.

**Downregulation of MMP14 following treatment with cryptotanshinone.** As demonstrated in Fig. 4, following treatment with cryptotanshinone, the expression levels of MMP14 were significantly downregulated compared with the control. By contrast, no significant differences were identified in the expression levels of MMP15, MMP16 and MMP24. This suggested the potential involvement of MMP14 in cryptotanshinone-mediated cellular apoptosis. In addition, no significant difference was revealed in the expression of TIMP2 following treatment with cryptotanshinone. Based on these findings, MMP14 may be considered as a potential target of miR-133a, which acts independently of TIMP2 regulation.

**Inhibition of cancer cell invasion.** As demonstrated in Fig. 5, the number of invasive A549 cells was significantly reduced
following treatment with cryptotanshinone. This indicated that the invasive capability of human lung adenocarcinoma epithelial A549 cells was significantly inhibited following treatment with cryptotanshinone.

Discussion

As a transcription activator that is translocated to the cell nucleus, STAT3 is activated following phosphorylation by receptor-associated Janus kinases (15). STAT3 serves a key role in the growth and apoptosis of cancer cells, including lung cancer cells (16). Cryptotanshinone can regulate mitochondrial function (17). As a STAT3 inhibitor, cryptotanshinone was demonstrated to inhibit the proliferation of lung cancer cells in the present study. Colivelin is a neuroprotective peptide and activator of STAT3 that suppresses neuronal death by activating STAT3 in vitro (18). The current study identified that even following activation of STAT3 with colivelin, cryptotanshinone could inhibit cancer cell growth and colony formation, and promote cancer cell apoptosis. This suggests that cryptotanshinone may serve a role independent of the STAT3 pathway.

miRNAs are small non-coding RNA molecules that are 22 nucleotides long, which exhibit functions in RNA silencing and post-transcriptional regulation of gene expression by targeting mRNAs. The human genome encodes >1,000 miRNAs in many cell types, which target >60% of all genes (19). The current study investigated the effects of cryptotanshinone on a number of miRNAs. miRNA expression analysis was performed following 6 h of treatment, while protein expression analysis was performed following 24 h of treatment as alterations in miRNA expression levels have been reported to occur at a faster rate compared with alterations in protein expression levels (20). It was identified that the expression levels of the following miRNAs were significantly altered following treatment with cryptotanshinone: miR-30d-5p, miR-126-3p, miR-133a, miR-338-3p, miR-451a, miR-21-5p, miR-96-5p, miR-182-5p and miR-205-5p. Among these miRNAs, the expression levels of miR-133a were the most significantly affected following treatment with cryptotanshinone.

MMPs, also termed matrixins, are calcium-dependent zinc-containing endopeptidases (21). MMPs are capable of degrading numerous extracellular matrix proteins and are involved in the cleavage of cell surface receptors, the release of apoptotic ligands and chemokine/cytokine inactivation (22). In addition, MMPs serve key roles in cell proliferation, migration, differentiation, angiogenesis, apoptosis and host defense (23,24). The present study identified that the expression levels of MMP14 were significantly downregulated following treatment with cryptotanshinone. Conversely, the expression levels of other type-I transmembrane proteins, MMP15, MMP16 and MMP24, were not significantly affected by cryptotanshinone treatment. Therefore, it can be suggested that MMP14 is a potential target of miR-133a. MMP14 can interact with TIMP2, an inhibitor of MMPs that is critical for the maintenance of tissue homeostasis (25,26). The current study investigated the expression of TIMP2 following treatment with cryptotanshinone, however, no significant change in expression level was detected compared with untreated control cells. This suggests that miR-133a, instead of TIMP2, regulates MMP14. A previous study has also demonstrated that miR-133a may target and regulate the expression of MMP14 (27).

The tumor microenvironment is important in the process of cancer metastasis and can be regulated by many natural products of herbs (28). Therefore, in future studies it may be beneficial to investigate the tumor microenvironment following treatment with cryptotanshinone. Furthermore, the metabolism of numerous chemicals is dependent on cytochrome P450, therefore, it may be beneficial to study the effects of cryptotanshinone on cytochrome P450 reductase, which serves a key role in cellular proliferation, and astrocytosis in particular (29).

Cancer metastasis is the leading cause of cancer-associated mortality, and is defined as the spread of cancer cells to neighboring tissues and organs beyond the initial tumor location (30). Cell invasion is the first and most important step of metastasis, which is a complex process that involves invasive cancer cells infiltrating nearby tissues and disseminating to secondary sites through the extracellular matrix (31,32). To the best of our knowledge, it remains unclear whether miR-133a directly induces a decrease in MMP14 expression or indirectly regulates this process via other factors. Providing that miR-133 is upregulated during this process, it may be beneficial to use an miR-133 inhibitor in the presence or absence of cryptotanshinone to determine the mechanism of regulation in future studies.

In conclusion, the present study identified that following treatment with cryptotanshinone, the metastatic capabilities of lung cancer cells were significantly inhibited. This indicates that cryptotanshinone may serve as a potential therapeutic agent for the treatment of lung cancer. Further research regarding this topic is currently being performed in our lab.

Acknowledgements

Not applicable.

Funding

This study was supported by Science and Technology Major Project of Gansu Province (grant no. 143FKDH002).

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

YSZ made substantial contributions to the design of the study. HW, YSZ, YGZ and WL performed the experiments. YGZ and JW performed the data analysis. HW and YSZ wrote 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.

References


