Glutathione S-transferase A1 mediates nicotine-induced lung cancer cell metastasis by promoting epithelial-mesenchymal transition

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Abstract. The present study aimed to investigate the effect of glutathione S-transferase A1 (GSTA1) on lung cancer cell viability, invasion and adhesion in the presence of nicotine *in vitro*. Furthermore, the effect of GSTA1 on the epithelial-mesenchymal transition (EMT), a process strongly associated with lung cancer metastasis, was examined. Human lung carcinoma A549 cells were treated with various concentrations of nicotine (0.01, 0.1, 1 and 10 μM) and levels of GSTA1 mRNA and protein were measured by reverse transcription-quantitative polymerase chain reaction and western blot analysis, respectively. To knock down GSTA1 expression, GSTA1-small interfering RNA was transfected into A549 cells. Cell viability, invasion and adhesion abilities were determined by MTT, Transwell-Matrigel invasion and cell adhesion assays, respectively. The expression of the epithelial cell markers E-cadherin and keratin, and the mesenchymal cell markers vimentin and N-cadherin in A549 cells were examined by western blot analysis. The current study indicated that the expression of GSTA1 was increased in A549 cells following nicotine treatment. GSTA1 suppression inhibited the viability, invasion and adhesion of lung cancer cells. In addition, the increase in lung cancer cell viability, invasion and adhesion by nicotine was suppressed following GSTA1 knockdown. Furthermore, GSTA1 affected the expression of EMT markers in nicotine-treated or untreated lung cancer cells. Thus the present study demonstrates that GSTA1 promotes lung cancer cell invasion and adhesion and mediates

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the effect of nicotine on lung cancer cell metastasis *in vitro*. Furthermore, the results demonstrated that GSTA1 exerts its effect on lung cancer cell metastasis by promoting the EMT.

Introduction

Globally, lung cancer is the leading cause of cancer-associated mortality in men and the second leading cause of cancer-associated mortality in women (1). Recurrence and metastasis are the biggest obstacles to effective lung cancer treatment. Although treatments for lung cancer have improved over the past few decades, the 5-year survival rate is only ~16% (2). Thus, it is important to identify biomarkers associated with lung cancer metastasis in order to improve the therapeutic strategies available.

Tumor metastasis is a complex process consisting of multiple biological steps (3) including increased motility, invasion into surrounding tissue, intravasation, entry and survival in the circulation, extravasation and eventual colonization of a distant site (4-6). The epithelial-mesenchymal transition (EMT) is a well-coordinated process that induces metastasis in epithelial cancer (7,8). Glutathione S-transferase A 1 (GSTA1) is an isoform of GST primarily involved in the detoxification of electrophilic compounds by undergoing conjugation with glutathione (9,10). Previous studies have demonstrated that the altered expression of GST genes increases the risk of prostate cancer and hepatocellular carcinoma (11-14). Recently, Pan *et al* (15) identified the potential of GSTA1 in the early diagnosis and treatment of lung cancer. It was determined that the expression of GSTA1 in lung cancer tissues and cells was higher than in healthy tissues and cells, and that GSTA1 suppression inhibits the proliferation of lung cancer cells. However, to the best of our knowledge, the potential role of GSTA1 in lung cancer metastasis has not yet been investigated.

It is estimated that ~85% of total lung cancer cases occur in tobacco smokers (16). Nicotine is the major addictive component of tobacco smoke and many of the carcinogens in tobacco smoke are derivatives of nicotine (17). Numerous studies have demonstrated that nicotine promotes the growth and metastasis of lung tumors (18-20).

In the present study, GSTA1-small interfering RNA was transfected into A549 cells to knock down GSTA1 expression, and the effect of GSTA1 on the viability, invasion and adhesion of lung cancer cells was investigated in the presence of nicotine *in vitro*. Furthermore, the effect of GSTA1 on EMT, a process strongly associated with lung cancer metastasis, was examined by western blot analysis.

Materials and methods

Cell culture and transfection. The A549 cell line was obtained from the American Type Culture Collection (Manassas, VA, USA) and cultured in Dulbecco's modified Eagle's medium (DMEM; Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA) containing 10% newborn calf serum (NBCS; Invitrogen; Thermo Fisher Scientific, Inc.). Cells were maintained at 37˚C in a humidified atmosphere containing 5% CO_2. A549 cells were seeded ($1x10^5$) in DMEM and Nicotine (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) was used to treat A549 cells at the concentrations of 0.01, 0.1, 1 and 10 μM at 37˚C for 24 h. The concentration of nicotine was selected by assessing which nicotine concentration exhibited the maximum effect on GSTA1 expression for subsequent experiments. Cells were treated with nicotine for 6, 12, 24 and 48 h in the preliminary experiments, and 24 h was selected as the duration following treatment with 10 μM nicotine, as together they had the maximum effect on GSTA1 expression for the subsequent experiments. GSTA1-small interfering RNA (siRNA) was synthesized by Sangon Biotech Co., Ltd. (Shanghai, China). A scramble siRNA (Sangon Biotech Co., Ltd.) was used as the control. A549 cells were transfected with siRNA (1 μM) using Lipofectamine® 2000 (Invitrogen; Thermo Fisher Scientific, Inc.) following the manufacturer's instructions. The cells used in this study were divided into four groups: PBS + Scr group (cells were transfected with scramble siRNA and treated with 1 μl PBS), PBS + Si group (cells were transfected with GSTA1-siRNA and treated with 1 μl PBS), Nicotine + Scr group (cells were transfected with scramble siRNA and treated with 10 μM nicotine) and Nicotine + Si group (cells were transfected with GSTA1-siRNA and treated with 10 μM nicotine). The PBS + Scr group was used as the control.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Total RNA was isolated from A549 cells using TRIzol reagent (Invitrogen; Thermo Fisher Scientific, Inc.). Total RNA (1 μg) was converted into cDNA using a First Strand cDNA Synthesis kit (Fermentas; Thermo Fisher Scientific, Inc.) following the manufacturer's instructions. The following primers were used in the present study: GSTA1, forward, 5'-GGCTGCAGCTGGAGTAGAGT-3' and reverse 5'-GCAAGCTTGGCATCTTTTTC-3' and β-actin, forward, 5'-AGAGCTACGAGCTGCCTGAC-3' and reverse 5'-AGCACTGTGTTGGCGTACAG-3'. qPCR was performed using a SYBR Green PCR kit (Applied Biosystems; Thermo Fisher Scientific, Inc.) in a 7300 Sequence Detection system (Applied Biosystems; Thermo Fisher Scientific, Inc.). The reaction was performed over 40 cycles at 95˚C for 30 sec, 59˚C for 30 sec and 72˚C for 30 sec. All reactions were performed in triplicate. Levels of GSTA1 mRNA were

normalized to those of β-actin, as an internal control using the $2^{-\Delta\Delta Cq}$ method (21).

Western blot analysis. Total protein was extracted using radioimmunoprecipitation assay lysis buffer (Beyotime Institute of Biotechnology, Shanghai, China) according to the manufacture's protocol. Briefly, the cells were incubated with the lysis buffer at room temperature for 5 min. Then cell lysates were centrifuged at 13,000 x g for 5 min at room temperature and supernatants were harvested. Equal amounts of total protein (20 μg) were separated using 12% SDS-PAGE and subsequently transferred onto nitrocellulose membranes (EMD Millipore, Billerica, MA, USA). Following blocking with 3% bovine serum albumin (Sigma-Aldrich; Merck KGaA) at 4˚C overnight, membranes were incubated with keratin rabbit polyclonal antibody (1:400; cat. no. 41723; Signalway Antibody Inc., College Park, MD, USA), GSTA1 monoclonal antibody (1:500; cat. no. sc-100546), E-cadherin rabbit polyclonal antibody (1:500; cat. no. sc-7870), vimentin monoclonal antibody (1:800; cat. no. sc-373717), N-cadherin mouse monoclonal antibody (1:500; cat. no. sc-393933) and β-actin monoclonal antibody (1:1,000; cat. no. sc-130301; all Santa Cruz Biotechnology, Inc., Dallas, TX, USA) at 4˚C overnight. Membranes were subsequently incubated with horseradish peroxidase-labeled goat anti-mouse (1:2,000; cat. no. sc-2005) or goat anti-rabbit antibodies (1:2,000; cat. no. sc-2004; both Santa Cruz Biotechnology, Inc.) at 37˚C for 1 h. Immunoreactive bands were detected using an enhanced chemiluminescence detection kit (Pierce; Thermo Fisher Scientific, Inc.). ImageJ version 1.37 software (National Institutes of Health, Bethesda, MD, USA) was used for quantification, and the experiments were independently performed three times.

MTT assay. An MTT assay was performed to determine cell viability. A549 cells were plated into 96-well plates at the density of $1x10^4$/well, and allowed to grow at 37˚C for 24, 48, 72 and 96 h. Subsequently, they were incubated with MTT solution (1 mg/ml; Sigma-Aldrich; Merck KGaA) at 37˚C for 4 h. Dimethyl sulfoxide (Sigma-Aldrich; Merck KGaA) was added to each well to solubilize the formazan product. Control cells were incubated with the dimethyl sulfoxide alone. The absorbance of each sample at 570 nm was measured using a microplate reader (Multiskan Ascent, Thermo Fisher Scientific, Inc.).

Cell invasion assay. The invasion of A549 cells was measured using a Transwell-Matrigel assay. Transwell inserts (Corning, Inc., Corning, NY, USA) were precoated with Matrigel matrix (BD Biosciences, Franklin Lakes, NJ, USA) at 37˚C for 30 min. Cells were trypsinized and plated into the upper chambers at a concentration of $3x10^4$ cells/ml in serum-free DMEM. Cell medium containing 10% NBCS was added to the lower chambers. The chambers were incubated at 37˚C overnight and a cotton swab was used to remove non-invasive cells. The invasive cells were then fixed in 95% ethanol at room temperature for 15 min followed by staining in hematoxylin for 10 min. The invaded cells were observed under an inverted microscope (XDS-200D; Caikon Optical Instrument Co., Ltd, Shanghai, China) at a magnification of

x400, and counted in 10 randomly selected fields in three independent experiments.

Cell adhesion assay. The 96-well plates were precoated with fibronectin (Sigma-Aldrich; Merck KGaA) and then blocked with 1% bovine serum albumin at 37°C for 2 h. A total of 3×10^4 A549 cells in serum-free DMEM were seeded into each well and incubated at 37°C for 2 h. Following washing with phosphate-buffered saline, cells were fixed in 4% para-formaldehyde (Shanghai Solarbio Bioscience & Technology Co., Ltd., Shanghai, China) at room temperature for 30 min. Subsequently, cells were stained with 0.5% crystal violet (Beyotime Institute of Biotechnology) followed by incubation with 2% SDS at room temperature for 1 min. The absorbance at 570 nm was measured using a microplate reader.

Statistical analysis. All the data are presented as the mean ± standard deviation. Statistical analysis was performed using SPSS 19.0 statistical software (SPSS, Inc., Chicago, IL, USA). One-way analysis of variance followed by the least significant difference test was used for comparison of multiple groups and $P < 0.05$ was considered to indicate a statistically significant difference.

Results

mRNA and protein expression of GSTA1 is induced by nicotine in A549 cells. A549 cells were treated with 0, 0.01, 0.1, 1 or 10 μM nicotine for 24 h. Subsequently, levels of GSTA1 mRNA and protein were examined by RT-qPCR and western blot analysis, respectively. Results from RT-qPCR demonstrated that levels of GSTA1 mRNA were elevated in A549 cells following nicotine stimulation and the maximum level was reached at 10 μM. Compared with controls, levels of GSTA1 mRNA were significantly increased following treatment with 0.1 ($P < 0.05$), 1 ($P < 0.01$) and 10 μM ($P < 0.01$) nicotine (Fig. 1A). Similar results were obtained following western blot analysis (Fig. 1B). We treated cells with nicotine for 6, 12, 24 and 48 h in the preliminary experiments, and 6 and 12 h of incubation was demonstrated to exhibit no effect on GSTA1 expression. Following preliminary assessments, 10 μM nicotine was selected for use in subsequent experiments, as the concentration of nicotine that exhibited the maximum effect on GSTA1 expression to treat cells, following 24 h of incubation, for the subsequent experiments.

GSTA1 suppression inhibits the nicotine-induced increased viability of A549 cells. To knock down GSTA1 expression, A549 cells were transfected with GSTA1-siRNA; scramble siRNA was used as a control. It was demonstrated by western blot analysis that levels of GSTA1 protein were significantly decreased in A549 cells following GSTA1-siRNA transfection ($P < 0.05$). Treatment with nicotine significantly increased GSTA1 expression compared with controls ($P < 0.01$), however, this increase in GSTA1 expression by nicotine was abrogated following transfection with GSTA1-siRNA ($P < 0.01$; Fig. 2).

To investigate the role of GSTA1 on cell viability, A549 cells transfected with GSTA1-siRNA were treated with 10 μM nicotine for 24 h and an MTT assay was performed. As
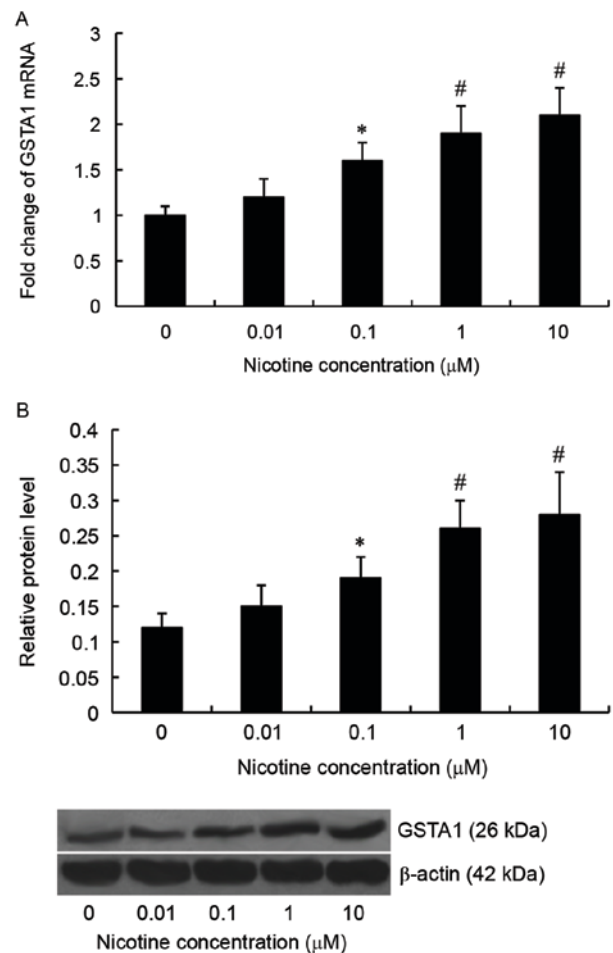

Figure 1. Levels of GSTA1 mRNA and protein in A549 cells following treatment with 0, 0.01, 0.1, 1 or 10 μM nicotine. (A) Relative mRNA expression of GSTA1. (B) Relative protein expression of GSTA1. *$P < 0.05$ and #$P < 0.01$ vs. 0 μM nicotine. GSTA1, glutathione S-transferase alpha 1.
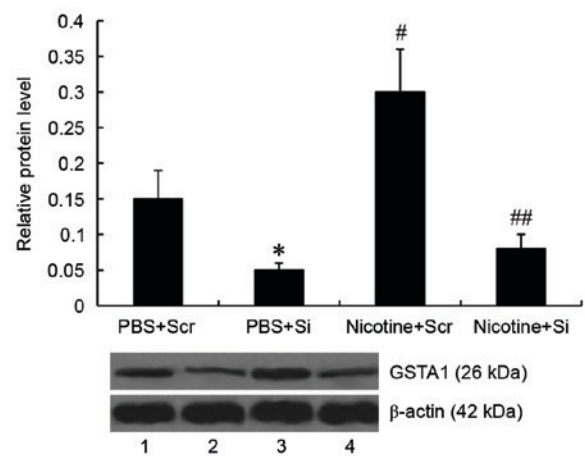

Figure 2. Expression of GSTA1 in A549 cells following treatment with nicotine and transfection with siRNA. *$P < 0.05$ and #$P < 0.01$ vs. PBS + Scr; ##$P < 0.01$ vs. Nicotine + Scr. Lane 1, PBS + Scr; lane 2, PBS + Si; lane 3, Nicotine + Scr; lane 4, Nicotine + Si. GSTA1, glutathione S-transferase alpha 1; si, siRNA; siRNA, small interfering RNA; Scr, scramble RNA.

presented in Fig. 3, nicotine stimulation significantly increased the viability of A549 cells transfected with scramble siRNA
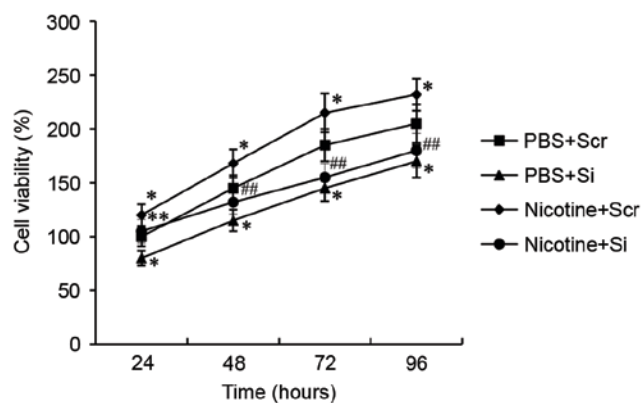
Figure 3. Viability of A549 cells following treatment with 10 μM nicotine and transfection with siRNA. *P<0.05 vs. PBS + Scr; **P<0.05 and ##P<0.01 vs. Nicotine + Scr. GSTA1, glutathione S-transferase alpha 1; Si, siRNA; siRNA, small interfering RNA; Scr, scramble RNA.
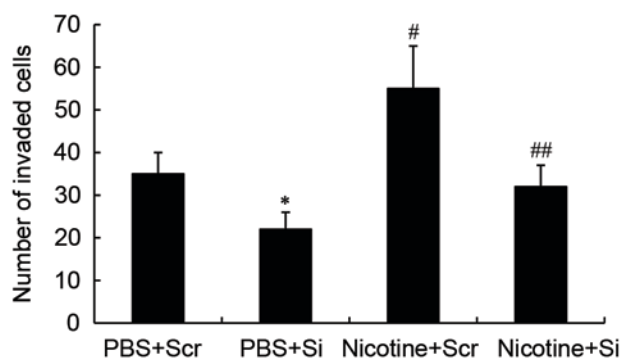

Figure 4. Invasion of A549 cells following treatment with 10 μM nicotine and transfection with siRNA. *P<0.05 and #P<0.01 vs. PBS + Scr; ##P<0.01 vs. Nicotine + Scr. GSTA1, glutathione S-transferase alpha 1; Si, siRNA; siRNA, small interfering RNA; Scr, scramble RNA.

Figure 5. Adhesion of A549 cells following treatment with 10 μM nicotine and transfection with siRNA. *P<0.05 and #P<0.01 vs. PBS + Scr; ##P<0.01 vs. Nicotine + Scr. GSTA1, glutathione S-transferase alpha 1; Si, siRNA; siRNA, small interfering RNA; Scr, scramble.
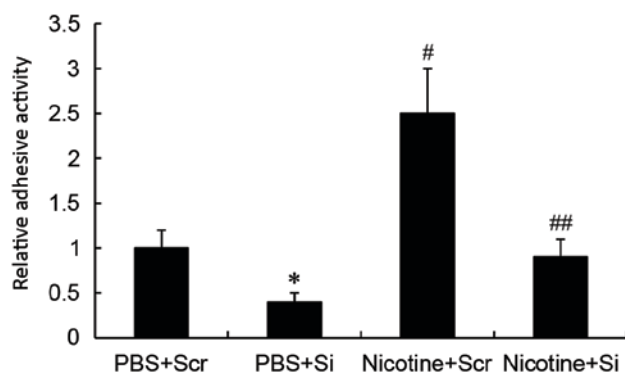
compared with untreated A549 cells (P<0.05). However, viability was significantly reduced following transfection with GSTA1-siRNA in nicotine-treated cells compared with the nicotine-treated scramble-siRNA cells (P<0.01) and in untreated cells transfected with GSTA1-siRNA compared with untreated cells transfected with scramble siRNA (P<0.05).
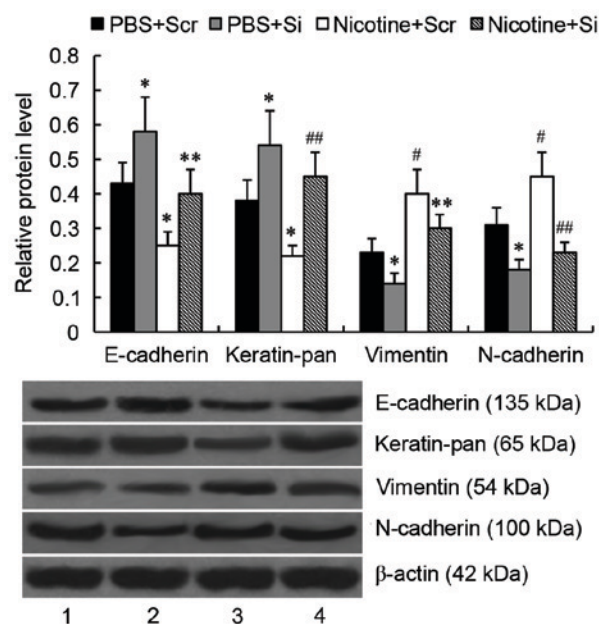

Figure 6. Levels of epithelial-mesenchymal transition markers in A549 cells following treatment with 10 μM nicotine and transfection with siRNA. *P<0.05 and #P<0.01 vs. PBS + Scr; **P<0.05 and ##P<0.01 vs. Nicotine + Scr. Lane 1, PBS + Scr; lane 2, PBS + Si; lane 3, Nicotine + Scr; lane 4, Nicotine + Si. GSTA1, glutathione S-transferase alpha 1; Si, siRNA; siRNA, small interfering RNA; Scr, scramble RNA.

GSTA1 suppression inhibits the nicotine-induced invasion abilities of A549 cells. A Transwell-migration assay was performed to determine whether GSTA1 affects cell invasion. In PBS-treated cells, GSTA1 knockdown significantly decreased the cell invasion ability (P<0.05) compared with cells transfected with scramble siRNA. The invasion ability of A549 cells significantly increased following nicotine treatment compared with PBS-treated A549 cells (P<0.01). However, nicotine-treated cells transfected with GSTA1-siRNA exhibited significantly decreased invasive ability compared with nicotine-treated cells transfected with scramble siRNA (P<0.01; Fig. 4).

GSTA1 suppression inhibits the nicotine-induced adhesion abilities of A549 cells. Cell adhesion assays demonstrated that significantly increased adhesion activity was observed in nicotine-stimulated A549 cells compared with PBS-treated cells transfected with scramble siRNA (P<0.01). However, GSTA1-siRNA knockdown significantly inhibited the adhesion activity of nicotine treated (P<0.01) and untreated (P<0.05) A549 cells compared with scramble siRNA nicotine-treated and untreated cells, respectively (Fig. 5).

GSTA1 suppression inhibits nicotine-induced EMT of A549 cells. To investigate the effect of GSTA1 on EMT in A549 cells, the expression of the epithelial markers E-cadherin and keratin, and the mesenchymal markers vimentin and N-cadherin, were determined by western blot analysis. As presented in Fig. 6, compared with untreated cells transfected with scramble siRNA, levels of E-cadherin and keratin were significantly downregulated (P<0.05), whereas levels of vimentin and N-cadherin were significantly upregulated (P<0.01) in nicotine-treated cells transfected with scramble siRNA. Following GSTA1

knockdown, levels of E-cadherin and keratin were significantly increased (P<0.05), whereas levels of vimentin and N-cadherin were significantly reduced (P<0.05) compared with untreated cells transfected with scramble siRNA. In nicotine-treated A549 cells, levels of E-cadherin (P<0.05) and keratin (P<0.01) were significantly increased and levels of vimentin (P<0.05) and N-cadherin (P<0.01) were significantly reduced compared with nicotine-treated cells transfected with scramble siRNA.

Discussion

Glutathione S-transferases are a multigene family of phase II isoenzymes that catalyze the conjugation of a variety of carcinogens, environmental toxins and endogenous compounds with glutathione (22). Polymorphisms associated with the altered expression of these genes may affect the metabolism of carcinogens and chemotherapeutic agents (23), resulting in an increased risk of various malignances (24-30). Previous studies have demonstrated that GSTA1 overexpression protects tumor cells from doxorubicin and reactive oxygen species-induced apoptosis (31-34). GSTA1 also serves a role in regulating the proliferation of Caco-2 cells (35). Recently, it has been demonstrated that GSTA1 promotes the proliferation of lung cancer cells (15). In the present study, it was demonstrated that the results from the MTT assay were consistent with those from a previous study (15). Furthermore, the present study demonstrated that GSTA1 suppression inhibited lung cancer cell invasion and adhesion, indicating that GSTA1 is associated with lung cancer metastasis.

Although there is insufficient evidence to classify nicotine as a carcinogen, nicotine has tumor-promoting properties. It has been demonstrated that nicotine promotes tumor growth and metastasis by inducing EMT, cell migration and invasion, angiogenesis, as well as suppressing apoptosis induced by drugs or radiation (36-38). In the present study, nicotine was used to treat A549 cells, and it was determined that the expression of GSTA1 was induced by nicotine. To determine whether GSTA1 was involved in mediating the effect of nicotine on lung cancer cells, GSTA1 was knocked down in A549 cells by siRNA. The results indicated that nicotine-induced lung cancer cell viability, invasion and adhesion were suppressed following GSTA1 knockdown.

It has been suggested that the EMT is the driving force of metastasis (7). During EMT, epithelial cells lose their apical and basolateral polarity, break cell-cell attachment and transform into mesenchymal cells (39). Notably, the EMT is reversible and the mesenchymal-epithelial transition is a reversion back towards the epithelial phenotype (40). E-cadherin and keratin are key markers of the epithelial phenotype and the loss of these proteins leads to the loss of cell junctions and the promotion of metastasis (41,42). Vimentin and N-cadherin are mesenchymal cell markers and have crucial roles in cellular migration (43,44). In the present study, GSTA1 affected the expression of EMT markers of A549 lung cancer cells. In addition, the present findings indicated that the nicotine-induced EMT was reversed following GSTA1 knockdown. These results suggest that the effect of GSTA1 on EMT may explain its effect on lung cancer metastasis.

In conclusion, the present study demonstrated that GSTA1 promotes lung cancer cell invasion and adhesion. In addition,

the present results suggest that nicotine increases the viability, invasion and adhesion abilities of A549 lung cancer cells and that this effect is mediated by GSTA1. GSTA1 exerts its effect on lung cancer cell metastasis by promoting the EMT. The present study suggests that GSTA1 serves a potential role GSTA1 in lung cancer metastasis and therefore may be a novel target for lung cancer treatment.

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