

Phenethyl isothiocyanate suppresses EGF-stimulated SAS human oral squamous carcinoma cell invasion by targeting EGF receptor signaling

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Abstract. Phenethyl isothiocyanate (PEITC) is a natural compound that is involved in chemoprevention as well as inhibition of cell growth and induction of apoptosis in several types of cancer cells. Previous studies have revealed that PEITC suppresses the invasion of AGS gastric and HT-29 colorectal cancer cells. However, the effects of PEITC on the metastasis of SAS oral cancer cells remain to be determined. Our results showed that PEITC treatment inhibited the invasion of EGF-stimulated SAS cells in a concentration-dependent manner, but appeared not to affect the cell viability. The expression and enzymatic activities of matrix metalloproteinase-2 (MMP-2) and matrix metalloproteinase-9 (MMP-9) were suppressed by PEITC. Concomitantly, we observed an increase in the protein expression of both tissue inhibitor of metalloproteinase-1 (TIMP-1) and -2 (TIMP-2) in treated cells. Furthermore, PEITC treatments decreased the protein phosphorylation of epidermal growth factor receptor (EGFR) and downstream signaling proteins including PDK1, PI3K (p85), AKT, phosphorylated IKK and I κ B to inactivate NF- κ B for the suppression of MMP-2 and MMP-9 expression. In addition, PEITC can trigger the MAPK signaling pathway through the increase in phosphorylated p38, JNK and ERK in treated cells. Our data indicate that PEITC is able to inhibit the invasion of EGF-stimulated SAS oral cancer cells by targeting EGFR and its downstream signaling molecules and finally lead to the reduced expression and enzymatic activities of both MMP-2 and MMP-9. These results suggest that PEITC is promising for the therapy of oral cancer metastasis.

Introduction

Head and neck squamous cell carcinoma (HNSCC), including oral cancer, is the sixth most common malignancy in humans worldwide. Oral cancer (OC) is one of the most frequent types of HNSCC. Approximately 95% of OCs are squamous cell carcinomas (OSCC) (1). Histologically, OSCCs are derived from the epithelium lining of the oral cavity and can occur at various sites in the oral cavity, including the lips, hard palate, gum and tongue (2), with a preference on the tongue and floor of the mouth. Each year, ~405,000 new cases of oral cancer (OSCC) are diagnosed and the number is still accumulating in many countries. In Taiwan, OSCC is the sixth leading cause of cancer death. Approximately 5,400 new cases are identified and 2,200 deaths per year and the incidence of OSCC has increased 6-fold during the past decade.

The main causes of oral cancer includes tobacco and alcohol consumption (3), diets poor in vitamin A and carotenoids, indoor air pollution and poor oral hygiene (4,5). The occurrence of oral cancer in Taiwan is closely related to betel quid chewing, cigarette smoking and alcohol consumption (6). The standard treatments for patients with oral cancer include surgery, radiotherapy and chemotherapy (7). Despite the improvement in surgery and chemotherapy during the last 20 years (8), oral cancer remains a disease with poor prognosis and a low survival rate (9). In patients identified with an advanced stage of the disease, there is a high incidence of invasion to adjacent tissues, of metastasis to lymph node and distant areas and of recurrence during the patient's lifetime (10,11). As compare to 90% of patients without metastasis, the 5-year survival rate for patients with lymph node metastasis at presentation is significantly reduced to 25-40% (12). Additionally, lymph node metastasis occurs in ~40% of patients with oral cancer. Therefore, there is an urgent need to identify agents that can inhibit the invasion and metastasis of oral cancers.

The active components in natural products such as polyphenolic and isothiocyanate (ITC)-containing compounds are the intensive target of research for their promising cancer

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preventative and therapeutic properties (13,14) and low toxicity to cells (15). Epidemiological investigations have reported an inverse relation between the dietary intake of fruits and vegetables, including cruciferous vegetables and the risk of various types of malignancies (16-18). The anti-carcinogenic effects of cruciferous vegetables such as broccoli (18) and watercress (19) have been ascribed to certain chemicals with the isothiocyanate ($-N=C=S$) functional group (19,20). Isothiocyanates are produced from the hydrolysis of the inactive precursor glucosinolates by myrosinase in cruciferous vegetables when the plant tissues are crushed or masticated (19-21). These isothiocyanates, which consist of phenethyl ITC (PEITC), allyl ITC (AITC), benzyl ITC (BITC) (22) and sulforaphane (SFN), have been shown to have potential cancer chemopreventive activity in a number of experimental models, including cancer of the esophagus, mammary gland, lung, liver, pancreas, fore-stomach, colon, small intestine and bladder of mice, rats, other rodents and colon cancer in humans (23-25).

PEITC, a member of isothiocyanate, possesses a variety of biological activities such as the induction of phase II detoxification enzymes, the inhibition of cytochrome P450 (CYP) enzymes (26), arrest of cell cycle (27) and stimulation of apoptosis (28-31), inhibition of nuclear factor- κ B (NF- κ B)-regulated gene expression (32) and activation of Atg5-mediated autophagy (33) in different cancer cell lines. PEITC also suppresses the pulmonary neoplasia induced by 4-(methylnitrosamino)-1-(3-pyridyl)-1-butone in the lung of A/J mouse (34,35), prevents the formation of colonic aberrant crypt foci induced by azoxymethane (36) and reduces the number and size of polyps in $Apc^{Min/+}$ mice (23). The incidence and burden (affected area) of poorly differentiated tumors in the dorsolateral prostate of transgenic adenocarcinoma of mouse prostate (TRAMP) model mice were reduced when given 3 μ mole PEITC/kg of diet (37). Further, PEITC is currently in clinical trials for lung cancer (38). These effects suggest a potential role of PEITC in the suppression of tumorigenesis.

It has been shown that PEITC inhibited the migration and invasion of human gastric cancer AGS cells (39) and colon cancer HT29 cells (40). In addition, PEITC treatment reduced angiogenesis and cell motility of human umbilical vein endothelial cells and PC-3 prostate cancer cells (41). However, the effects and underlying mechanism of PEITC on the metastasis of oral squamous cell carcinomas are still not clear. In this study, we demonstrated that PEITC acted on the phosphorylation of EGFR and sequentially inactivated the PI3K/AKT kinase cascade, repressed the NF- κ B-mediated signaling and hence reduced expression of matrix metalloproteases (MMPs), finally leading to the inhibition of OSCC invasion.

Materials and methods

Chemicals and reagents. Phenethyl isothiocyanate (PEITC) was purchased from Sigma-Aldrich (St. Louis, MO, USA). Antibodies against AKT, EGFR, ERK, I κ B, JNK, MMP-2, MMP-9, p38, PI3K, TIMP-1, TIMP-2, β -actin and GAPDH were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Antibodies against phospho-AKT (S308), phospho-

AKT (S473), phospho-EGFR (Y845), phospho-EGFR (Y992), phospho-EGFR (Y1068), phospho-ERK (Thr202/Tyr204), phospho-IKK, phospho-I κ B, phospho-JNK (Thr183/Tyr185), phospho-PDK1, phospho-PI3K and phospho-p38 (Thr183/Tyr185) were obtained from Cell Signaling Technology (Danvers, MA, USA). HRP-conjugated secondary antibodies such as rabbit anti-mouse IgG, goat anti-rabbit IgG and donkey anti-goat IgG were purchased from Santa Cruz Biotechnology. MTT (3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide) and epidermal growth factor (EGF) were obtained from Sigma-Aldrich. DMEM medium, fetal bovine serum (FBS), L-glutamine, penicillin-streptomycin and trypsin-EDTA were purchased from Gibco BRL (Invitrogen Life Technologies, Carlsbad, CA).

Cell culture. Human OSCC SAS cell line was cultured in DMEM medium supplemented with 10% of fetal bovine serum, 100 U/ml penicillin, 100 μ g/ml streptomycin and 2 mM glutamine and incubated at 37°C in a humidified chamber with 5% CO₂ (42).

Cell invasion assay. The membrane of each transwell insert was washed with 1X PBS and pre-coated with Matrigel (2 mg/ml, 20 μ l; BD Matrigel™ Invasion chamber). SAS cells (2×10^4) were seeded into the chamber of the insert and incubated with 0.5 ml of complete DMEM medium in each transwell. Cells were treated with EGF (100 ng/ml) and various concentrations of PEITC (0, 0.5, 1 and 2 μ M) for 48 h and then cells inside the chamber were removed by a cotton swab. Invaded cells were fixed with 4% formaldehyde in PBS and stained with 0.1% of hematoxylin, captured and the number of invaded cells was counted (43,44).

Cell viability assay. SAS cells (2×10^4) were seeded into the 96-well plate and treated with EGF (100 ng/ml) and PEITC (0, 0.5, 1 and 2 μ M) for 48 h. Medium was removed and replaced with fresh DMEM medium containing MTT (0.5 mg/ml) and cultured at 37°C incubator for an additional 4 h. Medium was again removed and 200 μ l of DMSO was added into each well to dissolve the formazan crystals and the absorbance of each well was measured at 570 nm with a reference wavelength at 620 nm on an ELISA reader. The data of control sample (0 μ M of PEITC) was set as 100% and the relative cell viability of drug-treated samples was calculated accordingly. Cell morphology was recorded by using a phase-contrast microscope (43,44).

Gelatin zymography assay. SAS cells (1×10^6) were seeded into 12-well plate for 48 h and treated with EGF (100 ng/ml) and various concentrations of PEITC (0, 0.5, 1 and 2 μ M) in serum-free DMEM medium for an additional 48 h. Culture medium was spun at 1000 x g for 10 min at 4°C, supernatant was collected and protein concentration was determined as described below. 5 μ g of total proteins were mixed with 2X sample buffer (0.125 M Tris-HCl, pH 6.8, 4% SDS, 20% glycerol, 0.01% bromophenol blue) and electrophoresed in an 8% SDS-polyacrylamide gel with 1% gelatin. Gel was incubated with 2.5% Triton X-100 at room temperature for 30 min to remove residual SDS and then incubated in Zymogen developing buffer (50 mM Tris, pH 7.5, 200 mM NaCl, 5 mM

CaCl₂, 1 μ M ZnCl₂, 0.02% Brij-35; Bio-Rad Laboratories, Hercules, CA, USA) at 37°C overnight. Gel was then washed extensively with water and stained with 0.5% Coomassie blue G-250 (0.5% Coomassie blue G-250, 50% methanol and 10% acetic acid) for 2 h and de-stained in de-staining solution (50% methanol and 10% acetic acid) until clear zones were evident. The gel was scanned by a scanning digitizing system and digitized by using free Image J software (NIH) (43,44).

Preparation of whole cell lysate. SAS cells were challenged with EGF (100 ng/ml) and treated with various concentrations of PEITC for the specified time and cells were collected for the preparation of whole cell lysate using iced-cold RIPA buffer (50 mM Tris-base, 150 mM NaCl, 0.1% SDS, 1% sodium deoxycholate, 1% NP-40, pH 7.5) supplemented with protease inhibitors including leupeptin (17 mg/ml), sodium orthovanadate (10 mg/ml), phenylmethanesulfonyl fluoride (10 mg/ml). Cells were completely re-suspended in extraction buffer and kept in ice for 30 min with occasional mixing and cell lysate were collected by a spin at 12,000 x g for 10 min at 4°C. The protein concentrations present in the samples were measured by using Bio-Rad Protein Assay Dye Reagent Concentrate (Bio-Rad) (43,45).

Western blotting. The obtained whole cell lysate was resolved in sodium dodecyl sulfate-polyacrylamide gel and transferred onto polyvinylidene fluoride (PVDF) membrane (Millipore) by using the iBlot Dry Blotting Transfer System (Invitrogen/Life Technologies). The transferred membranes were blocked in 5% non-fat milk (prepared in Tris-buffered saline supplemented with 0.1% Tween-20; TBST) at ambient temperature for 1 h and incubated with primary antibody at 4°C overnight. Membranes were washed with TBST three times for 10 min before incubated with HRP-coupled secondary antibody for 1 h. Protein signals were visualized by enhanced chemiluminescence (ECL) and exposed to Bio-MAX MR X-ray film (Eastman Kodak, Rochester, NY, USA) (43,46,47).

Quantitative real-time PCR analyses. SAS cells were treated with 0, 1 and 2 μ M of PEITC and EGF (100 ng/ml) for 24 h and cells were collected. Total RNAs were isolated using the Qiagen RNeasy mini Kit. cDNAs were synthesized using the High Capacity cDNA Reverse Transcription kit according to the supplier's brochure (Applied Biosystems). For the quantitative PCR reaction, 1 μ l of cDNAs were mixed with 2X SYBR Green PCR Master Mix (Applied Biosystems) and 200 nM of forward and reverse primers (see below for detailed sequences). PCR reaction was performed on an Applied Biosystems 7300 Real-Time PCR system in triplicate according to the following conditions: 2 min at 50°C, 10 min at 95°C and 40 cycles of 15 sec at 95°C, 1 min at 60°C. Fold changes of the gene expression were derived using the comparative C_T method (40,48). The used primer pairs were: human MMP-2-forward, 5'-CCCCAGACAGGTGATCTTGAC-3'; human MMP-2-reverse, 5'-GCTTGCGAGGGAAGAAGT TG-3'; human MMP-9-forward, 5'-CGCTGGGCTTAGAT CATTCC-3'; human MMP-9-reverse, 5'-AGGTTGGATACAT CACTGCATTAGG-3'; human GAPDH-forward, 5'-ACACC CACTCCTCCACCTTT-3'; human GAPDH-reverse, 5'-TAGC CAAATTCGTTGTCATACC-3' (49).

Immunofluorescence staining. SAS cells were seeded onto slides overnight and incubated with EGF (100 ng/ml) and PEITC (0, 1 and 2 μ M) for 6 h. Cells were fixed in 4% formaldehyde at room temperature for 15 min, washed with PBS and permeabilized with 0.1% Triton X-100 in PBS for 15 min, then incubated with primary antibodies at 4°C overnight. After extensive washes, cells were incubated with FITC-conjugated secondary antibodies for 2 h. Cells were sealed and images were acquired with a fluorescence microscope (Nikon) and processed in Photoshop 7.0 software (50).

Statistical analysis. One-way ANOVA followed by Student's t-test was used to evaluate the differences between treated and experimental groups. p<0.05 was considered to define a statistically significant difference (49,51).

Results

PEITC inhibits EGF-stimulated invasion of SAS cells. We therefore determined the effects of PEITC on EGF-stimulated SAS cells. EGF-treatment increased the invasion of SAS cells, as revealed by Matrigel invasion assay (Fig. 1A). Treatment of EGF-stimulated SAS cells with PEITC decreased the invasion of cells in a concentration-dependent manner (Fig. 1A). From Fig. 1A, we have known that PEITC inhibited the invasion of EGF-stimulated SAS cells, a result that could be due to the inhibition of PEITC on the viability of EGF-stimulated cells. To test this, we treated EGF-stimulated SAS cells with different concentrations (0, 0.5, 1 and 2 μ M) of PEITC and performed MTT cell viability assay. The result showed that PEITC at 0.5-2 μ M of concentrations did not inhibit the viability of EGF-stimulated SAS cells, as compared to cells without drug treatment (Fig. 1B). The cell morphology was comparable between treatments with 0 and 2 μ M of PEITC (see inserts in Fig. 1B).

PEITC inhibits the enzymatic activities and gene expression of MMP-2 and MMP-9 in EGF-stimulated SAS cells. Studies have shown that matrix metalloproteinases including MMP-2 (gelatinase A) and MMP-9 (gelatinase B) are expressed in oral cancers. These two MMPs are closely linked to the malignant potential of tumor cells and are also important for tumor invasion and metastasis (44,52). To evaluate the effects of PEITC on the enzymatic activities of MMP-2 and MMP-9, we treated EGF-stimulated SAS cells with different concentrations (0, 0.5, 1 and 2 μ M) of PEITC and assessed the enzymatic activities of MMP-2 and MMP-9 by gelatin zymography. As shown in Fig. 2A, treatments of cells with PEITC suppressed the enzymatic activities of both MMP-2 and MMP-9 in a concentration-dependent manner. Pronounced inhibition of the activities was observed at concentrations >1 μ M of PEITC (see columns 3 and 4 and inserts of Fig. 2A). To address whether the inhibition of MMP-2 and MMP-9 was at the transcriptional level, we treated EGF-stimulated cells with PEITC (0, 1 and 2 μ M) and the effects of PEITC were analyzed by quantitative RT-PCR. As shown in Fig. 2B, treatment of PEITC significantly decreased the gene expression of both MMP-2 and MMP-9 in a concentration-dependent fashion. Taken together, these data suggested that both gelatinases (MMP-2 and MMP-9) are involved in the EGF-induced invasion of SAS cells.

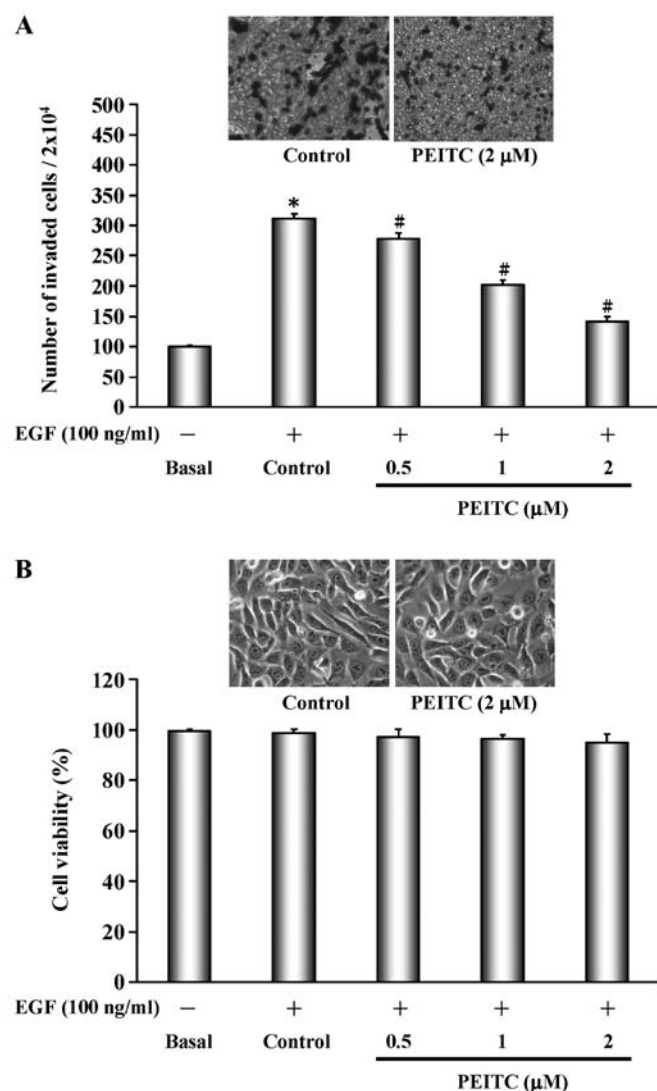


Figure 1. The effects of PEITC on invasion and viability of EGF-stimulated SAS cells. (A) PEITC inhibited the cell invasion of EGF-stimulated SAS cells. Cells were incubated in drug-free medium (basal) or treated with PEITC (0, 0.5, 1 and 2 μ M) in medium containing EGF (100 ng/ml) for 48 h and cells that invaded through Matrigel into the lower surface of the filter were stained and counted. Inserts, images display the results of treatments with 0 (control) and 2 μ M of PEITC. * P <0.05, as compared to the treatment in basal medium. # P <0.05, as compared to the treatment with 0 μ M of PEITC (control) in EGF-containing medium. (B) PEITC did not affect the cell viability of EGF-stimulated SAS cells. Cells were treated with PEITC (0, 0.5, 1 and 2 μ M) in the presence of EGF (100 ng/ml) for 48 h and cell viability was determined by MTT assay as described in Materials and methods. The cell viability is expressed as percentage by setting control treatment as 100%. Each point is the mean \pm SD of three repeats. Inserts, images display the cell morphology of treatments with 0 and 2 μ M of PEITC.

PEITC inhibits the protein expression of MMP-1, MMP-2 and increased the protein expression of TIMP-1 and TIMP-2 in EGF-stimulated SAS cells. Since PEITC inhibited the enzymatic activities and gene expression of MMP-2 and MMP-9 in EGF-stimulated cells, we inferred that PEITC could inhibit the protein expression of both metalloproteinases. As shown in Fig. 3A, treatment of EGF-stimulated SAS cells with PEITC (1 and 2 μ M) reduced the protein expression of MMP-2 and MMP-9. It is well known that tissue inhibitor of metalloproteinases (TIMPs), TIMP-1 and TIMP-2, can bind and inhibit the enzymatic activities of MMP-2 and MMP-9 (53). We there-

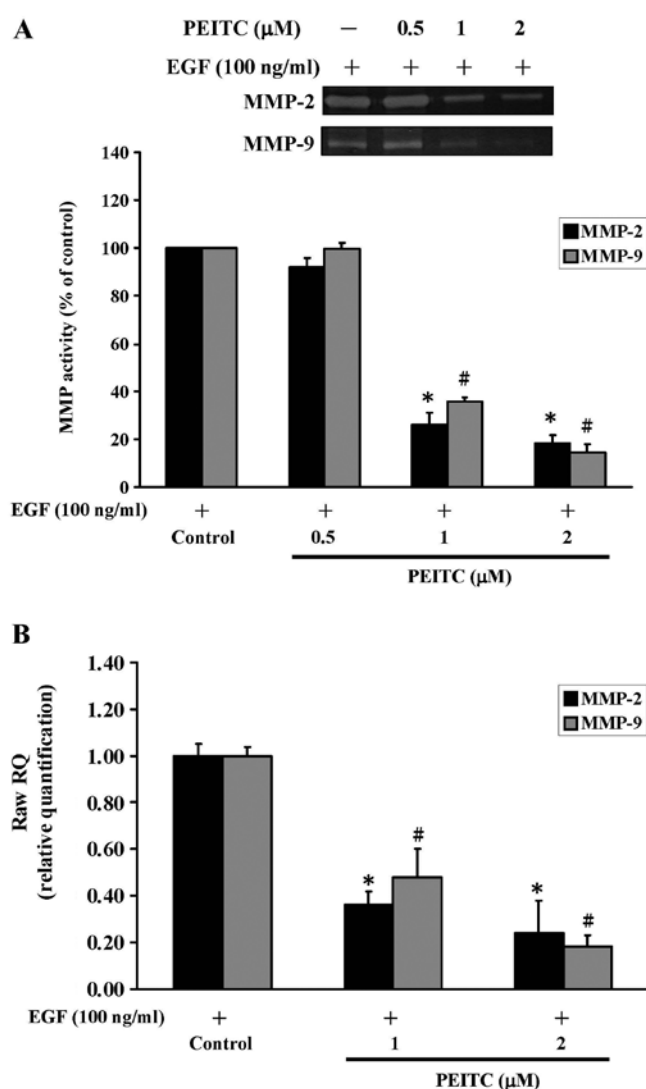


Figure 2. PEITC inhibits the EGF-stimulated enzymatic activities and gene expression of MMP-2 and MMP-9 of SAS cells. (A) PEITC inhibited the EGF-induced enzymatic activities of MMP-2 and MMP-9. Cells were treated with different concentrations of PEITC (0, 0.5, 1 and 2 μ M) in the presence of EGF (100 ng/ml) for 48 h. The conditioned media were collected for gelatin zymography assay to determine the MMP-2 and MMP-9 activities. Shown here are the densitometric data expressed as the mean \pm SD of three independent experiments. Enzymatic activity from control treatment (0 μ M) was set as 100% and the percentage of higher drug concentrations was calculated consequently. * P <0.05, a significant statistical difference compared to the enzymatic activity of MMP-2 without PEITC treatment (control). # P <0.05, a significant statistical difference compared to the enzymatic activity of MMP-9 without PEITC treatment (control). (B) PEITC decreased the gene expression of MMP-2 and MMP-9. Cells were treated with different concentrations of PEITC (0, 1 and 2 μ M) in the presence of EGF (100 ng/ml) for 24 h and gene expression of MMP-2 and MMP-9 was analyzed by quantitative RT-PCR. Gene expression of both control treatments were set as 1.0. Each point is the mean \pm SD of three experiments. * P <0.05, a significant statistical difference compared to the gene expression of MMP-2 without PEITC treatment (control). # P <0.05, a significant statistical difference compared to the gene expression of MMP-9 without PEITC treatment (control).

fore examined the protein expression of TIMP-1 and TIMP-2 in EGF-stimulated cells treated with PEITC (0, 1 and 2 μ M). The result showed that treatment with PEITC significantly increased the protein expression of TIMP-1 and TIMP-2 (Fig. 3A). These data suggested that PEITC can suppress the gene expression and protein expression of MMP-2 and MMP-9

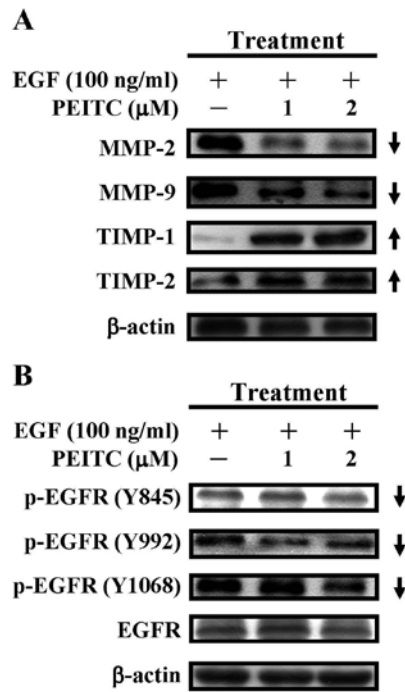


Figure 3. PEITC inhibits the expression of metalloproteases MMP-2, MMP-9 and induces the expression of tissue inhibitor of metalloproteinases TIMP-1 and TIMP-2 through inactivation of the epidermal growth factor receptor (EGFR) in EGF-stimulated SAS cells. (A) The effects of PEITC on the expression of MMP-2, MMP-9, TIMP-1 and TIMP-2 proteins in EGF-stimulated SAS cells. Cells were treated with EGF (100 ng/ml) and PEITC (0, 1 and 2 μ M) for 48 h and cells were harvested for western blot analyses with MMP-2, MMP-9, TIMP-1 and TIMP-2 antibodies, respectively. The β -actin served as the loading control. (B) The effects of PEITC on the activation of the epidermal growth factor receptor (EGFR) in EGF-stimulated SAS cells. Cells were treated with EGF (100 ng/ml) and PEITC (0, 1 and 2 μ M) for 6 h and cell lysates were subjected to western blot analyses and detected with p-EGFR (Y845), p-EGFR (Y992), p-EGFR (Y1068) and EGFR antibodies, respectively. The β -actin served as the loading control.

and increase the protein expression of TIMP-1 and TIMP-2, leading to the decrease in the enzymatic activities of MMP-2 and MMP-9 in EGF-stimulated SAS cells.

PEITC suppressed the activation of EGFR. Binding of EGF to its cognate receptor EGFR results in the activation of EGFR that involves the autophosphorylation of EGFR and activation of intracellular signaling pathways, such as activation of PI3K/AKT, mitogen-activated protein kinases (MAPKs) and the signal transducer and activators of transcription (STATs) pathways, leading to cell proliferation and survival, invasion, metastasis and angiogenesis (54-56). Thus, we determined the effects of PEITC on the activation of EGFR proteins by examining the tyrosine phosphorylation of EGFR in EGF-challenged SAS cells. As shown in Fig. 3B, PEITC treatment (1 and 2 μ M) inhibited the tyrosine phosphorylation of EGFR at Y845, Y992 and Y1068 concentration-dependently, while the protein levels of total EGFR remained largely unchanged. Immuno-fluorescent staining with anti-p-EGFR (Y1068) and anti-p-EGFR (Y845) also showed that tyrosine-phosphorylated EGFRs at Y1068 and Y845 were dramatically reduced in EGF-challenged cells treated with PEITC (Fig. 4). These data suggested that PEITC treatment suppressed the activation of EGFR in EGF-challenged SAS cells.

PEITC decreases the protein phosphorylation of PI3K, AKT, IKK and I κ B α in EGF-stimulated SAS cells. Since treatment with PEITC suppressed the activation of EGFR in EGF-challenged cells (Figs. 3B and 4), we next examined the effects of PEITC on downstream PI3K/AKT and NF- κ B signaling pathways. PEITC treatment (1 and 2 μ M) reduced the protein phosphorylation of PDK1, PI3K (p85) and profoundly reduced the protein phosphorylation of AKT (S308) and AKT (S473) in EGF-challenged cells (Fig. 5A, panels 1, 2, 4 and 5). Total proteins of PI3K (p85) was not changed during drug treatment, although total proteins of AKT were slightly reduced at 2 μ M of PEITC. In addition, PEITC treatment (1 and 2 μ M) decreased the protein phosphorylation of IKK and of I κ B α , while increased the protein stability of I κ B α . These data indicated that PI3K/AKT and NF- κ B signaling pathway downstream of EGFR signaling is downregulated by PEITC treatment.

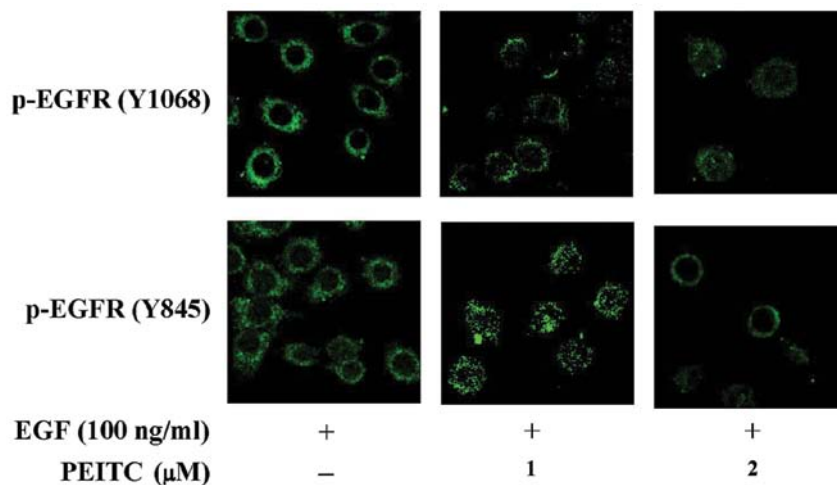


Figure 4. PEITC suppresses the phosphorylation of EGFR in EGF-stimulated SAS cells by immuno-fluorescent staining. Cells were treated with EGF (100 ng/ml) and PEITC (0, 1 and 2 μ M) for 6 h, fixed and stained with anti-p-EGFR (Y845) and anti-p-EGFR (Y1068), followed by staining with FITC-coupled goat anti-mouse antibodies.

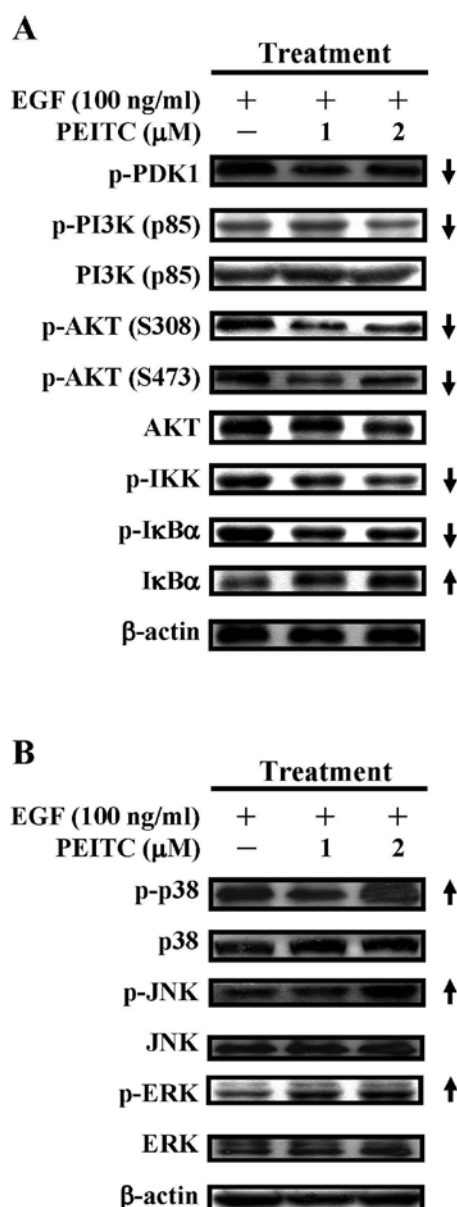


Figure 5. The effects of PEITC on the PI3K/AKT and MAPK signaling pathways in EGF-stimulated SAS cells. Cells were treated with EGF (100 ng/ml) as well as PEITC (0, 1 and 2 μ M) for 6 h and cell lysates were prepared for the western blot analyses of (A) phosphorylated PDK1, PI3K (p85), phosphorylated PI3K (p85), AKT, phosphorylated AKT (S308 and S473), phosphorylated IKK, I κ B α and phosphorylated I κ B α and (B) p38, phosphorylated p38, JNK, phosphorylated JNK, ERK and phosphorylated ERK respectively. The β -actin served as the loading control.

The effects of PEITC on the MAPK signaling pathway in EGF-stimulated SAS cells. We also determined the effects of PEITC on the MAPK signaling pathways-p38, JNK and ERK signaling pathways downstream of EGFR activation. PEITC treatment (1 and 2 μ M) increased the protein phosphorylation of p38 at higher drug concentration (2 μ M) (Fig. 5B, top panel). PEITC treatment (1 and 2 μ M) also increased the protein phosphorylation of JNK and ERK in a concentration-dependent manner. The protein expression of p38, JNK and ERK appeared not to be affected (Fig. 5B, panels 2, 4 and 6), suggesting that the MAPK signaling pathway can be activated after PEITC treatment.

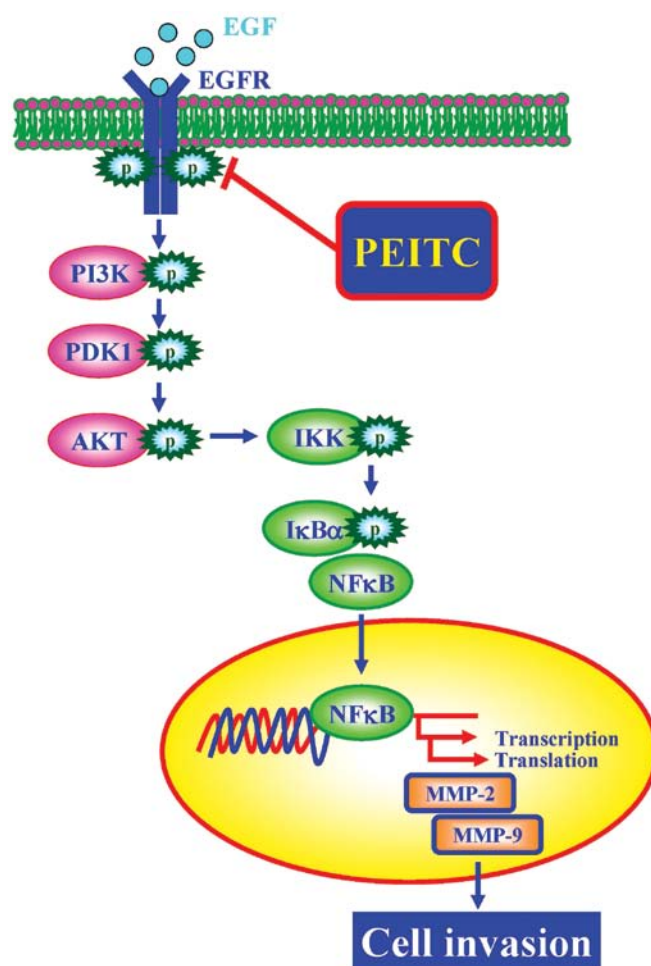


Figure 6. A model depicting the molecular mechanisms of PEITC on the inhibition of EGF-stimulated SAS cell invasion.

Discussion

Oral squamous cell carcinoma (OSCC) is a leading cause of cancer deaths in the world, characterized by poor prognosis and a low survival rate in spite of advances in treatment with surgery and radiotherapy. The main cause of death in OSCC is metastasis which primarily occurs through the lymphatic system. Once it has metastasized to the lymph nodes, the overall mortality rate of the disease is high and the 5-year overall survival rate does not exceed 50%, which is one of the lowest rates for all major cancers (10,57). Therefore, identification of new drugs for the chemotherapy of OSCC metastasis is highly desirable.

The epidermal growth factor receptor (EGFR) belongs to the HER/ErbB protein family of receptor tyrosine kinases. The EGFR gene encodes a 170-kDa transmembrane glycoprotein with its tyrosine kinase domain located within the cytoplasmic region. Ligand binding induces the activation of tyrosine kinase activity that triggers intracellular signaling cascades, including the Ras-Raf-mitogen-activated protein kinase pathway, the phosphatidylinositol 3-kinase-AKT pathway and the signal transducer and activators of transcription pathway, which contribute to cell proliferation and survival (58). Dysregulation in the signaling of EGFR and

other members of the tyrosine kinase receptor family has been linked to cell transformation, autonomous cell growth, angiogenesis, invasion and metastases in a number of cancers (59). Up to 90% of head and neck squamous cell carcinoma (HNSCC) patients are identified with EGFR overexpression, which is considered to be involved in tumorigenesis and metastasis (60). Overexpression of EGFR in HNSCC is often associated with the simultaneous increase in its ligands such as the transforming growth factor α (54,61), which will lead to excessive activation of EGFR signaling either in an autocrine- or paracrine-dependent manner. Thus, EGFR appears to be a promising therapeutic target for oral cancer metastasis (44). Our data indicated that PEITC can inhibit the EGF-induced invasion of SAS cells through the inactivation of EGFR and downstream signaling, including the suppression in the phosphorylation cascade of PI3K, PDK1 and AKT and hence the reduction of phosphorylated IKK, the decrease in the phosphorylation of I κ B and the simultaneous increase in the stability of I κ B and the block in the release and activation of NF- κ B. These results are consistent with previous studies using prostate cancer PC-3 cells (32,62).

Matrix metalloproteases (MMPs) are responsible for the degradation of the extracellular matrix and facilitating spreading and metastasis of tumor cells. They are strongly blocked by the endogenous tissue inhibitors of metalloproteinases (TIMP-1, -2, -3 and -4). The expression of MMP-2 and MMP-9 was shown to be associated with tumor invasion and lymph node metastasis of oral cancer (63). The expression can also be regulated by NF- κ B as their promoters possess NF- κ B binding sites. As expected, the activities and expression of MMP-2 and MMP-9 were downregulated after PEITC treatment in EGF-stimulated SAS cells (Figs. 2 and 3A). These results could be due to the inactivation of NF- κ B caused by the disruption in the EGFR signaling after PEITC treatment, leading to the failure in the expression of MMP-2 and MMP-9. Concomitantly, the increase in the expression of MMP-2 and -9 inhibitors, TIMP-1 and -2 proteins, was observed (Fig. 3A). These data suggested that PEITC suppressed the invasion and metastasis of EGFR-overexpressed oral cancers by reducing the expression and activities of MMP-2 and MMP-9 through the interference with the phosphorylation of EGFR and downstream signaling.

The other major downstream pathway regulated by EGFR is MAPK. Our finding that the phosphorylation of p38, JNK and ERK was increased after PEITC treatment in SAS cells is contradictory to the previous reports that PEITC can suppress the MAPK activation to inhibit the invasion and metastasis of HT-29 colon cancer cells and AGS gastric cancer cells (39,40). These discrepancies could be due to cell type-specific effects. Alternatively, it has been reported that PEITC can induce cell apoptosis through the activation of MAPK (29,64-68) and this raises the possibility that PETIC may trigger apoptosis in addition to the suppression of invasion and metastasis in our system. However, our data showed that the cell viability was not affected after PEITC treatment (1-2 μ M) (Fig. 1B). This suggests that the activation of MAPK by PEITC may elicit the downstream signaling and drive specific genes expression to repress the invasion and metastasis of OSCC, but not to cause apoptosis. Further detailed mechanism needs to be elucidated.

In conclusion, the signaling pathway underlying the effects of PEITC on the invasion of EGF-stimulated SAS cells is shown in Fig. 6. PEITC suppressed the phosphorylation and activation of EGFR and inhibited the sequential phosphorylation and activation of PI3K, PDK1 and AKT, resulting in the reduced expression of phosphorylated IKK and hence the reduced protein phosphorylation and increased protein stability of I κ B α , which in turn suppressed the expression and enzymatic activities of MMP-2 and MMP-9, contributing to the inhibition of invasion in EGF-challenged SAS cells. Our data suggested that PEITC will be a promising therapeutic agent for the treatment of oral cancer metastasis.

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