Abstract. Epithelial-mesenchymal transition (EMT) is a critical cellular process in cancer metastasis, during which epithelial polarized cells become motile mesenchymal cells. Since transforming growth factor-β (TGF-β) is a potent inducer of EMT, blocking of TGF-β/Smad signaling has become a promising cancer therapy. Nobiletin, a polymethoxy flavonoid from Citrus depressa, has been shown to be valuable for cancer treatment, yet the mechanism remains unclear. In the present study, lung adenocarcinoma A549 and H1299 cells were used to evaluate the effect of nobiletin on EMT induced by TGF-β1. Nobiletin successfully inhibited TGF-β1-induced EMT, migration, invasion and adhesion in vitro, accompanied by attenuation of MMP-2, MMP-9, p-Src, p-FAK, p-paxillin, Snail, Slug, Twist and ZEB1 expression. Nobiletin inhibited the transcriptional activity of Smads without changing the phosphorylation status or translocation of Smads induced by TGF-β1. Moreover, Smad3 is requisite in TGF-β1-stimulated EMT. Smad3 overexpression meaningfully impaired the ability of nobiletin to reverse TGF-β1-induced EMT. In vivo, nobiletin prohibited the growth of metastatic nodules in the lungs of nude mice. Moreover, nobiletin inhibited tumor growth and reversed EMT in mice bearing A549-Luc xenografts, as revealed by IVIS imaging and immunohistochemical analysis. Collectively, the data suggest that nobiletin prevents EMT by inactivating TGF-β1/Smad3 signaling.
Nobiletin can inhibit TGF-β, which can drive EMT and metastasis in lung adenocarcinoma. To investigate the potential of nobiletin in the treatment of lung adenocarcinoma, experiments were carried out for each experimental condition. Nobiletin (10 and 20 µM) was added to A549 cells for 24 h. Three independent experiments were performed and showed that nobiletin can inhibit TGF-β1-induced EMT by blocking Smad3 transcription, signifying a potential use of nobiletin in the treatment of lung adenocarcinoma.

Materials and methods

Cell culture and viability assay. Human adenocarcinoma A549 and H1299 cells (Sigma-Aldrich, St. Louis, MO, USA) were grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 100 IU/ml penicillin and 100 µg/ml streptomycin. Cellular viability was measured by MTT reduction assay. MTT solution (5 mg/ml) was added to dissolve the formed formazan crystals. The absorbance was measured at 570 nm against a 690 nm reference with a microplate reader.

Adhesion and Transwell migration assays. For adhesion assays, A549 and H1299 cells were treated with TGF-β1 (5 ng/ml) or/and nobiletin for 24 h. Cells were suspended in DMEM and plated at a density of 1x10⁴ cells/well onto 96-well plates coated with collagen IV. Cells were then fixed with methanol and stained with 0.1% crystal violet and absorbance was read at 595 nm spectrophotometrically.

For the scratch assays, A549 cells were plated in 24-well tissue culture plates and allowed to reach confluency. At this point a scratch was introduced into the monolayer using a sterile 200-µl pipette tip. The cells were then washed in warm phosphate-buffered saline (PBS) to remove the debris and the medium was replaced with either fresh medium (control samples) or medium containing TGF-β1 (5 ng/ml) or/and nobiletin (10 and 20 µM) for 24 h. Three independent experiments were carried out for each experimental condition.

The migration assay was performed using 24-well 8-µm pore Transwell inserts (Corning), with the upper face of polycarbonate filters coated with Matrigel. Cells (1x10⁵) were added to the upper chamber. The lower chambers were filled with 20% FBS/DMEM media to serve as a chemoattractant. The non-migrated cells in the upper chamber were removed gently by a cotton swab at 24 h following reseeding, whereas, the migrated cells in the lower side were stained with 0.1% crystal violet. The mean of six fields from three separate trials was used to calculate the average number of migratory cells.

Immunofluorescence assay. Immunofluorescence assay was performed as previously described (16). In brief, cultured cells grown on plastic chamber slides were fixed in 4% formaldehyde for 30 min at room temperature. Then the cells were permeabilized in 0.2% Triton X-100 and blocked with 10% goat serum, and then incubated overnight at 4°C with the primary antibodies (diluted 1:100 in blocking buffer) against E-cadherin and N-cadherin. Afterwards, the sample was washed with PBS containing blocking buffer and incubated with the fluorophore-conjugated secondary antibody (Abcam). Cover glasses were counterstained with 4,6-diamidino-2-phenylindole dihydrochloride (DAPI) (5 mg/ml; Invitrogen Life Technologies). Fluorescence images were captured using a fluorescence microscope (Carl Zeiss, Göttingen, Germany).

Transient transfection. PcDNA3.1-Flag-Smad3 plasmid was prepared using Qiagen plasmid DNA preparation kit. Plasmid or Smad2/3 small interfering RNA (siRNA) (Santa Cruz Biotechnology, Santa Cruz, CA, USA) transient transfection of A549 cells was performed according to the protocol of the Lipofectamine 2000™ reagent (Invitrogen, Carlsbad, CA, USA).

Western blot analysis. A549 and H1299 cells were lysed at 4°C (30 min) in Tris-buffered saline. Whole lysates were analyzed by SDS-PAGE, and transferred to polyvinylidene difluoride (PVDF) membranes. The membranes were blocked with 5% skim milk with Tween-20 and probed with primary antibodies against E-cadherin, N-cadherin, vimentin, Snail, Slug, ZEB1, Twist, Src, FAK and β-actin (Thermo Fisher Scientific). Bound antibodies were conjugated with horseradish peroxidase secondary antibodies, and the membranes were developed with chemoluminescence reagents. Protein expression was semi-quantified by densitometric analysis.

RNA extraction and RT-PCR. Total RNA was isolated from A549 cells using the total RNA extraction kit as described by the manufacturer (Life Technologies, Inc., Grand Island, NY, USA). RT-PCR was performed using the Qiagen RT-PCR kit. The appropriate primers (MMP-2, 5'-GGCCCTGTCACTCTCGAGAT-3' and 5'-GGCATCCAGGTTATCGGGGA-3'; MMP-9, 5'-AGGCCTCTACAGAGTCTTTG-3' and 5'-GGCATCCAGGTTATCGGGGA-3'; β-actin, 5'-AGGCCTCTACAGAGTCTTTG-3' and 5'-GGCATCCAGGTTATCGGGGA-3') were used for PCR amplifications under the following conditions: 30 cycles at 95°C for 1 min, 60°C (MMP-2 and GAPDH) or 59°C (MMP-9) for 1 min, 72°C for 1 min and 72°C for 10 min. The PCR products were separated by 1.5% agarose gel and visualized by etidium bromide staining.
In vivo xenograft metastasis and tumor growth experiments. All mice were obtained from the SLAC Laboratory Animal Center of Shanghai. Ethical approval was obtained for the use of the animals, and all experiments were performed in accordance with the Guidelines for Animal Care of the Institutional Animal Care and Use Committee of the Tumor Hospital of Xinjiang Medical University. i) Five-week-old nude mice were injected with A549 cells via the tail vein (4x10^5 cells in 0.2 ml PBS). Nobiletin (20 and 40 mg/kg) was orally administered once daily. Cyclophosphamide (20 mg/kg) was used as the positive control, injected intraperitoneally every three days. Saline was administered orally every day for the control group (n=8 mice/group). After 42 days, the mice were sacrificed. After fixation in Bouin’s solution, the number of lung metastases were confirmed under a dissecting microscope. The lungs were further subjected to hematoxylin and eosin (H&E) staining. ii) C57BL/6 mice were injected with Lewis cells via the tail vein (8x10^5 cells in PBS) to establish a metastatic model. Administration was the same as the first model. After 42 days, the mice were sacrificed. After fixation in Bouin’s solution, the number of lung metastases were confirmed under a dissecting microscope. The lungs were further subjected to hematoxylin and eosin (H&E) staining. iii) Nude mice were implanted subcutaneously with A549-Luc cells. Administration was the same as the first model. Thirty days after cell implantation, imaging was performed with an IVIS imaging system (Caliper Life Sciences, Inc., Hopkinton, MA, USA), and nude mice were sacrificed and tumors were excised and weighed.

Immunohistochemistry. Immunohistochemistry of lung tissue was performed as previously reported (17). Formalin-fixed, paraffin-embedded tissue sections were cut at a thickness of 4 µm. Antigen recovery was performed by heating in 0.01 M sodium citrate buffer at 80˚C for 30 min using a microwave oven. Endogenous peroxidase activity and non-specific staining were blocked with the kit reagents. The tissues were incubated overnight with the primary E-cadherin and N-cadherin antibodies. Tissues were then incubated with peroxidase-bound IgG. The slides were counterstained with hematoxylin. Mounted tissues were imaged on an Olympus microscope. The numbers of E-cadherin- or N-cadherin-positive cells were determined at a magnification of x400 in three segments/slide.

Statistical analysis. Statistical analysis was performed using ANOVA, χ^2 or Student’s t-test using SPSS 16.0 software. Statistical significance was assigned at P<0.05.

Results

Nobiletin rebalances EMT-related genes. An MTT assay was used to determine the non-cytotoxic concentrations of

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Figure 1. Nobiletin inhibits TGF-β1-induced EMT in A549 and H1299 cells. (A) Chemical structure of nobiletin. (B) A549 and H1299 cells were treated with TGF-β1 (5 ng/ml) or/and nobiletin (Nob) for 24 h, and then cell viability was determined with the MTT assay. Values are expressed as the mean ± SD from triplicates. *P<0.05, compared with the control. †P<0.05, compared with the cells treated with TGF-β1 only. (C) The expression levels of epithelial marker, E-cadherin, and mesenchymal marker, N-cadherin, were evaluated by immunofluorescence. (D) EMT-related proteins were assessed by western blotting.
nobiletin (0-100 µM) in the presence of TGF-β1 (5 ng/ml) for 24 h. According to the MTT assays (Fig. 1B), concentrations of 10 and 20 µM were selected for further study, which exhibited no significant toxicity on cell viability. EMT is a crucial step in tumor metastasis, characterized by polarity and epithelial markers, as well as motile mesenchymal cells. Specifically, E-cadherin is a negative EMT biomarker, while N-cadherin is a mesenchymal biomarker. As shown in the immunofluorescence assay, TGF-β1-stimulated A549 and H1299 cells exhibited characteristic N-cadherin increase and E-cadherin decrease (Fig. 1C). However, nobiletin rebalances EMT-related genes disturbed by TGF-β1. In addition, similar results were obtained from the protein assays (Fig. 1D). The most prominent E-cadherin regulators are Snail-related zinc-finger transcription factors (Snail and Slug), ZEB1 and Twist. TGF-β1 treatment increased the expression of Snail, Slug, ZEB1 and Twist, and the increase was reversed by nobiletin treatment (Fig. 1D).

**Nobiletin inhibits TGF-β1-induced adhesion, migration and invasion in lung adenocarcinoma cells.** (A) The effect of nobiletin (Nob) on TGF-β1-induced migration, invasion and adhesion. The invasive ability of A549 and H1299 cells was quantified by counting the number of cells that invaded to the underside of the porous membrane. Images were captured under phase contrast microscope. *P<0.05, compared with the control. *P<0.05, compared with the cells treated with TGF-β1 only. (B) Effects of nobiletin on TGF-β1-induced expression of MMP-2 and MMP-9 were measured by RT-PCR and western blotting, respectively. (C) p-FAK, p-Src, p-paxillin levels were assessed by western blotting.

**Nobiletin inhibits the metastatic capacity of lung adenocarcinoma cells along with the downregulation of MMP-2 and MMP-9 and the FAK/Src pathway.** Nobiletin suppressed both A549 and H1299 cell invasion in the transmembrane migration assay induced by TGF-β1 (Fig. 2A). Moreover, nobiletin
inhibited the capacity of A549 and H1299 cells to adhere to wells coated with human collagen type IV. After TGF-β stimulation, A549 cells displayed enhanced wound closure activity and almost filled the scratched area, whereas the wound still displayed an obvious gap in the nobiletin-pretreated group.

RT-PCR and western blot assays were applied to examine the mRNA and protein levels of MMP-2/-9. An increase in the mRNA and protein levels of MMP-2/-9 was observed following TGF-β incubation, while nobiletin treatment abrogated the increase (Fig. 2B). Activation of FAK is normally induced by various growth factors, including TGF-β. Therefore, we further explored the effect of nobiletin on the activation of focal adhesion-related signaling molecules, FAK, Src and paxillin. As shown in Fig. 2C, TGF-β induced the phosphorylation of FAK, Src and paxillin and these phosphorylation events were inhibited by nobiletin.

Nobiletin prevents the transcriptional activity of Smads induced by TGF-β. In the TGF-β/Smad signaling pathway, TGF-βR2 induces recruitment and phosphorylation of TGF-βR1 after binding to TGF-β1 (18). Both TGFβR1 and TGFβR2 can be detected in A549 cells. As expected, the expression of TGFβR1 and TGFβR2 did not change following either TGF-β1 or nobiletin treatment. TGF-β1-triggered EMT is Smad-dependent. The activated TGFβR1 subsequently induces phosphorylation of Smad2/3. Therefore, western blot analysis was carried out to examine whether nobiletin influences Smad phosphorylation. TGF-β1-treated cells showed aggravated phosphorylation of Smad2/3 after a 60-min incubation, and the increase in phosphorylation was not affected after nobiletin exposure (Fig. 3A). Phosphorylated Smad2 and Smad3 form heteromeric complexes with the mediator Smad4. These Smad2/3/4 complexes translocate to the nucleus, and bind to the Smad binding elements (SBEs). Nobiletin failed to prevent Smads from translocating into the nucleus, as induced by TGF-β (Fig. 3B). Transcriptional activity of Smad was also examined. A SBE-luciferase reporter plasmid was stably transfected into A549 cells to evaluate the luciferase activity. Consistent with previous studies, TGF-β considerably augmented the luciferase-reporter activity of Smad, which was greatly abolished by nobiletin treatment (Fig. 3C). These data infer that nobiletin prohibited EMT by eliminating the transcriptional activity of Smads, rather than influencing the translocation or phosphorylation.

Nobiletin suppresses EMT via the TGF-β1/Smad3 signaling pathway. In order to explore the detailed mechanism of nobiletin in regards to EMT, we transfected Smad2 siRNA and Smad3 siRNA into A549 cells, respectively. After the A549 cells were treated with TGF-β1 and nobiletin (Nob) for 24 h, E-cadherin, N-cadherin, Snail, ZEB1 were detected by western blotting. The promoter expression of E-cadherin and N-cadherin after nobiletin treatment was significantly downregulated due to Smad3 knockdown. In contrast, Smad2 siRNA had little or negligible effect on the expression of EMT-related proteins. In the nobiletin-treated group, these
protein expression levels exhibited a similar tendency compared with the Smad3 siRNA-transfected A549 cells (Fig. 4A). To further confirm the interaction between nobiletin and Smad3, A549 cells were transfected with PcDNA3.1-Flag-Smad3. Accordingly, the expression of E-cadherin was decreased, while N-cadherin, Snail and ZEB1 were increased. Smad3 overexpression considerably reduced the ability of nobiletin to inhibit the TGF-β1-induced increase in N-cadherin and to reverse the TGF-β-stimulated decrease in E-cadherin. These data reveal the essential role of Smad3 in EMT progression and propose that nobiletin obstructs EMT by blocking Smad3-dependent transcriptional activity.

Nobiletin decreases metastasis and tumor growth in vivo. To further explore the antemetastatic potential of nobiletin in vivo, two experimental metastatic models of lung cancer were applied. One model was established by injecting A549 cells in nude mice via the tail vein. Nobiletin (20 and 40 mg/kg) was administered orally once daily. Cyclophosphamide was used as the positive control. As shown in Fig. 5A, the number of lung metastatic nodules was 65±15 in the control group on day 42, while there were only 43±11 and 29±9 nodules in the
in vivo EM 

EMT and the growth of lung cancer cells

protein. All these results suggest that nobiletin suppressed E-cadherin protein expression, and inhibited N-cadherin group) (Fig. 5C). In the IHC analysis, nobiletin facilitated E-cadherin protein expression, and inhibited N-cadherin protein. All these results suggest that nobiletin suppressed EMT and the growth of lung cancer cells in vivo.

Discussion

Despite advancements in the diagnosis of NSCLC along with revolutionary progress in targeted therapies, treatment of NSCLC after distant organ metastasis remains a great challenge. Various plant or fruit-derived agents with few side-effects have been chosen as potential alternatives for the therapy of lung cancer. Citrus species is a natural product containing phytochemicals with promise in cancer therapy. Numerous studies have demonstrated the citrus potential as a chemopreventive agent. As well, nobiletin has anticancer activity in various malignant tumors such as gastric, lung, colon and nasopharyngeal cancers. TGF-β is a multifunctional secreted polypeptide that signals through receptor serine/threonine kinases. Accumulating evidence has revealed that TGF-β stimulates metastasis and invasion in various tumors. TGF-β has received much attention as a characterized inducer of EMT during cancer progression and metastasis (19). Upon activation of the TGF-β pathway, cancer cells undergo evident and irreversible EMT, finally resulting in more aggressive and metastatic tumors. In the present study, we demonstrated that nobiletin suppresses adhesion, EMT, metastasis and invasion of lung cancer via antagonizing the TGF-β1/Smad3 signaling pathway.

Transformation of a non-motile polarized epithelial phenotype to a motile and invasive mesenchymal phenotype is frequently correlated with a high risk of metastasis and poor prognosis (20). The cadherin switch during EMT in NSCLC is regarded as a hallmark of invasiveness and metastasis. E-cadherin is a cell-cell adhesion molecule and EMT is characterized by E-cadherin loss. Reduced E-cadherin expression promotes cell mobility and invasion due to several transcriptional repressors, including ZEB1, Twist, Slug and Snail. EMT-related protein alteration is a significant prognostic marker to predict overall survival in patients with lung adenocarcinoma (21). Snail and Slug, which are zinc finger factors, are overexpressed in epithelial cell lines. Slug is a direct repressor of E-cadherin through E-box interaction and promotes cell invasion and metastasis. ZEB1 is enhanced by Slug (22). Our finding presents compelling evidence that nobiletin effectively reverses the acquisition of mesenchymal characteristics and suppresses the expression of transcription factors in TGF-β-stimulated lung cancer cells. The key mechanism that we elucidated to account for this important novel function of nobiletin is that it inhibits the transcriptional activity of Smad3.

Ligand binding to TGF-β receptors initiates Smad2/3/4 complex formation. Smads are mainly localized in the cytoplasm. After the binding of TGF-β to the receptor and subsequent TGF-βR1 phosphorylation, trimers consisting of phosphorylated-Smad2/3 and Smad4 translocate to the nucleus, where they cooperate with transcription factors such as Snail and Twist to repress the expression of epithelial markers and activate the expression of mesenchymal markers at the mRNA level (23). This signaling is referred to as TGF-β-activated Smad signaling in EMT. Smad3 is regarded as a positive regulator of EMT while Smad2 is known as a negative mediator (24). In line with previous research, the primary role of Smad3 was confirmed in the present study. Mesenchymal markers such as N-cadherin, Snail and ZEB1 were downregulated due to Smad3 siRNA transfection of A549 cells. However, Smad2 siRNA had little or negligible effect on the expression of EMT-related proteins. Nobiletin influenced neither the expression of phospho-Smad nor Smad translocation into the nucleus. However, the transcriptional activity of Smad3 was almost abolished following nobiletin treatment. Moreover, the imbalanced promoter expression of E-cadherin and N-cadherin were brought to a normal level by nobiletin treatment. Collectively, these data suggest that nobiletin targets Smad3 to suppress EMT.

MMP-2 and MMP-9 have been shown to promote cell invasion and metastasis of NSCLC. MMPs have also been identified as another class of inducers that activate developmental EMT. MMPs remodel the extracellular matrix to facilitate EMT and to promote cell specification during embryonic development. Furthermore, MMP-2 was reported to be involved in the suppressive effect of nobiletin on tumor formation and metastasis in a nasopharyngeal cancer xenograft model in vivo (25). Moreover, nobiletin inhibited tumor cell invasive activity not only by suppressing MMP-9 but also augmenting TIMP-1 production in human fibrosarcoma HT-1080 cells (26). EMT transcription factors stimulate MMP expression, leading to an increase in MMPs, which further activates EMT transcription factors in a feed-forward loop (27). TGF-β signaling regulates the balance of MMP/TIMP to control matrix homeostasis (28). Recent evidence indicates that the recruitment of MMP-9 to the tumor cell surface triggers TGF-β activation and enhances tumor growth and invasion (29). TGF-β incubation resulted in an obvious surge in both the mRNA and protein levels of MMP-2 and MMP-9, while a significant reduction in MMP-2 and MMP-9 levels was observed due to nobiletin treatment.

High expression of FAK protein is reported in many cancers (30). A novel function for paxillin, Src and FAK has been discovered in mediating EMT and aggressiveness regulated by TGF-β (30-32). FAK can be activated by a reduction in E-cadherin to facilitate focal adhesion assembly (33). FAK correlates with integrin clustering at the site of focal adhesions. As an oncogene involved in cell migration and in lung cancer tissues, paxillin is also highly expressed, amplified, and associated with EMT (34). In addition, Src-mediated regulation of E-cadherin and vimentin has been described in several cancers (35). Moreover, EMT is associated with acquired resistance to the EGFR inhibitor. Inhibition of the Src/FAK pathway
contributed to overcome the EMT-associated resistance to erlotinib (36). In line with results revealing TGF-β1-mediated upregulation of Src, FAK and paxillin, TGF-β1 in A549 cells strongly induced the phosphorylation of Src, FAK and paxillin, and nobiletin markedly inhibited the induction.

In summary, abrogation of TGF-β1-induced EMT may be an appropriate therapy to block metastasis. The present study provides compelling evidence that nobiletin exerts modulatory effects on TGF-β1 signaling and its downstream responses, not only in vitro but also in metastatic experimental models in vivo. Nobiletin may be advanced as a new therapy to treat NSCLC metastasis and EMT-related disorders.

References