

# Metformin Combined with 4SC-202 Inhibited the Migration and Invasion of OSCC via STAT3/TWIST1

This article was published in the following Dove Press journal:  
*OncoTargets and Therapy*

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**Background:** Oral squamous cell carcinoma (OSCC), the most common epithelial malignant neoplasm in the head and neck, characterizes with local infiltration and metastasis of lymph nodes. The five-year survival rate of OSCC remains low despite the advances in clinical methods. Thus, it is necessary to develop a new effective therapeutic scheme for OSCC. Our previous results showed that metformin and 4SC-202 synergistically promoted the intrinsic apoptosis of OSCC in vitro and in vivo, but the effects on invasion and migration remained unclear.

**Methods:** Human OSCC cell lines HSC6 and CAL33 were cultured with metformin (16 mM) or/and 4SC-202 (0.4  $\mu$ M) for 72 h. STAT3 inhibitor S31-201 was applied at concentration of 60  $\mu$ M for 48 h. Wound-healing assays and transwell assays were used to determine the invasion and migration ability of OSCC. qRT-PCR and Western blot were performed to detect mRNA levels and protein levels.

**Results:** Metformin or/and 4SC-202 suppressed the migration and invasion of OSCC cells. Importantly, the expression of TWIST1 was suppressed by metformin and 4SC-202, while the invasion and migration inhibitory effects of metformin and 4SC-202 were countered by the overexpression of TWIST1. In addition, the phosphorylation level of STAT3 decreased after the administration of metformin or/and 4SC-202. Furthermore, inhibition of STAT3 by S31-201 suppressed the expression of TWIST1 and led to a decline in migration and invasion of OSCC, while overexpression of TWIST1 attenuated these effects.

**Conclusion:** Metformin and 4SC-202 suppressed the invasion and migration of OSCC through inhibition of STAT3/TWIST1, and this scheme can serve as a novel therapeutic strategy for OSCC.

**Keywords:** oral squamous cell carcinoma, metformin, 4SC-202, invasion, migration

## Introduction

Oral squamous cell carcinoma (OSCC) is the most common oral cancer, and accounts for more than 90% of all oral tumors.<sup>1</sup> Due to the delayed clinical detection and the high incidence of local invasion and metastasis, the prognosis of OSCC is poor.<sup>2</sup> Despite advances in therapeutic strategies, five-year survival rate of OSCC has not improved significantly in recent years. Therefore, development of new therapeutic methods for OSCC becomes particularly important.

Protein acetylation modification plays a vital role in the tumorigenesis. In OSCC, high expression of HDACs such as HDAC1, HDAC2, HDAC6 were associated with poor prognosis, advanced stage, larger tumor size, and lymph node metastasis,<sup>3-5</sup> suggesting that HDACs were involved in OSCC progress and could serve as potential drug targets. 4SC-202 is a novel selective class I histone deacetylase inhibitor (HDACi). In vitro, 4SC-202 was observed to inhibit survival and proliferation of several types of cancer cells including hepatocellular carcinoma

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cell, colorectal cancer cell, medulloblastoma cell, and urothelial carcinoma cell,<sup>6–9</sup> and 4SC-202 was shown to inhibit TGF- $\beta$ -induced epithelial-to-mesenchymal transition (EMT) of pancreatic cancer cells.<sup>10</sup> Our previous findings revealed that 4SC-202 promoted the apoptosis of OSCC,<sup>11</sup> however, the effects of 4SC-202 on migration and invasion in OSCC remained unclear.

Metformin, a low cost agent widely used for diabetes, received considerable attention for its antitumor effect. It was found that metformin suppressed invasion and migration of various types of cancers including esophageal squamous cell carcinoma, cholangiocarcinoma and prostate cancer.<sup>12–14</sup> In addition, metformin could enhance the efficacy of several chemotherapeutics.<sup>15–17</sup>

Drug combination could achieve the therapeutic efficacy at reduced dosages, thus minimize the toxicity and other side effects of high doses of single drugs. Our previous results showed that metformin and 4SC-202 synergistically promoted the intrinsic apoptosis of OSCC in and in vivo,<sup>11</sup> but the effects of this combination on invasion and migration remained elusive.

In this study, we evaluated the effects and mechanisms of metformin and 4SC-202 on the invasion and migration of OSCC. Here, we found that metformin and 4SC-202 synergistically inhibited the migration and invasion of OSCC. Importantly, our results revealed that the combined treatment of metformin and 4SC-202 inhibited invasion and migration by suppressing STAT3/TWIST1 in OSCC. These findings further emphasized the combination of metformin and 4SC-202 as a promising potential therapeutic strategy for OSCC.

## Materials and Methods

### Cell Culture

Human OSCC cell lines HSC6 and CAL33 were kindly provided by J Silvio Gutkind (NIH, Bethesda, MD, USA), SCC25 (CRL-1628) was purchased from American Type Culture Collection (ATCC, Manassas, VA, USA), and HSC3 was obtained from Professor Qianming Chen (State Key Laboratory of Oral Diseases, Sichuan University, China). The use of these cell lines was approved by our institutional research ethics committee. The HSC3, HSC6 and CAL33 cells were maintained in DMEM (Gibco, Grand Island, NY, USA) supplemented with 10% FBS (Gibco). SCC25 was cultured in DMEM supplemented with 10% FBS and 400 ng/mL hydrocortisone. All cells were maintained at 37°C with 5% CO<sub>2</sub>. The

cells were treated with metformin (16 mM) or/and 4SC-202 (0.4  $\mu$ M) for 72 h, and STAT3 inhibitor (40  $\mu$ M, 60  $\mu$ M) was applied for 48 h.

Information regarding reagents and antibodies were listed in [Supplementary Table 1](#).

### Wound-healing Assays

The cells were seeded into six-well plates, and then cells were wounded in a line across the slides with a sterile 200  $\mu$ L plastic pipette tip. All cell debris was removed by washing with PBS. Cell migration, indicating wound-healing effect, was demonstrated as percentage of wounded area, that was the distance migrated by OSCC cells at the indicated point in time relative to the initial length of the wound. Experiments were performed at least three times.

### Transwell Assays

Cell migration and invasion capacities were examined with 24-well transwell chamber (pore size: 8  $\mu$ m) (Corning Costar, Corning, NY, USA). The upper chambers were precoated with or without 0.1 mL (300  $\mu$ g/mL) Matrigel matrix (Corning) for invasion or migration assay, respectively. The prepared cells were seeded in the upper chambers with serum-free DMEM, but the bottom chambers were filled with DMEM containing 10% FBS. Then, cells were incubated for 24 h. The noninvading cells were gently removed by a cotton swab, whereas the invading cells were fixed in 4% paraformaldehyde for 15 min, stained with 0.4% crystal violet for 10 min and photographed under an inverted light microscope (Olympus Corporation, Tokyo, Japan). Finally, the invasive cells in five randomly selected visual fields at 100 $\times$  magnification were counted.

### RT-PCR

Total RNA was isolated with TRIzol (Invitrogen, CA, USA) and cDNA was synthesized by reverse transcription according to the manufacturer's procedure (Takara, Dalian, China). The real-time PCR was carried out utilizing the LightCycler 480 SYBR Green I Master system (Roche, Basel, Switzerland). Relative mRNA expression levels were normalized to *GAPDH* using the  $2^{-\Delta\Delta Ct}$  method. The primers used were listed in [Table 1](#).

### Western Blot (WB) Analysis

Cells were lysed with RIPA buffer supplemented with protease and phosphatase inhibitor cocktail (CW

**Table 1** Primers Used for RT-PCR

Gene	Forward Primer (5'-3')	Reverse Primer (5'-3')
<i>TWIST1</i>	GACCTAGATGTCATTGTTCCAGAG	GACCTAGATGTCATTGTTCCAGAG
<i>SNAIL</i>	ACTGCAACAAGGAATACCTCAG	GCACTGGTACTTCTTGACATCTG
<i>SNAI2</i>	CGAACTGGACACACATACAGTG	CTGAGGATCTCTGGTTGTGGT
<i>ZEB1</i>	CAGCTTGATACCTGTGAATGGG	TATCTGTGGTCGTGTGGGACT
<i>GAPDH</i>	GCACCGTCAAGGCTGACAAC	TGGTGAAGACGCCAGTGGA

**Abbreviation:** qRT-PCR, quantitative reverse transcription-polymerase chain reaction.

Biotech, Beijing, China). 40–60 µg denatured protein of each sample was separated on 10% SDS-PAGE gels and transferred to a PVDF membrane (Millipore, Billerica, MA, USA). Then the membrane was blocked in 5% non-fat milk for one hour at room temperature and incubated with primary antibodies overnight at 4°C. Subsequently, the membrane was washed by 0.1% TBST for three times and incubated with HRP-conjugated secondary antibody for one hour at room temperature. The immunoreactive bands were visualized by a highly sensitive chemiluminescence (ECL) detection system (Millipore) and the immunoreactive bands were analyzed by densitometry using Image J (Bethesda, MD, USA). Similar results were obtained from three independent experiments.

## Cell Transfection

CAL33 and SCC25 cells were transfected with Twist1 siRNA (50 nM, RiboBio, Guangzhou, China) using the lipofectamine RNA iMAX transfection reagent (Invitrogen) following the manufacturer's instructions. The OSCC cells were transfected with lentivirus plasmid sh-TWIST1 or lentivirus plasmid Flag-TWIST1 (Cyagen, CA, USA), then the stable transfected cells were screened with puromycin. The *TWIST1* sequences were as follows: sh-TWIST1, 5'-ATGGCAAGCTGCAGCTATGTG3', si1-TWIST1, 5'-TGAGCAACAGCGAGGAAGA-3', and si2-TWIST1, 5'-GATGGCAAGCTGCAGCTAT-3'.

## Statistics

All statistical analyses were performed using SPSS 20.0 software (SPSS, Chicago, IL, USA). All results were graphically depicted as mean ±SD from triplicate experiments performed in a parallel manner. Two-tailed Student's *t*-test and one-way ANOVA were performed to detect statistically significant differences.  $P < 0.05$  was considered statistically significant.

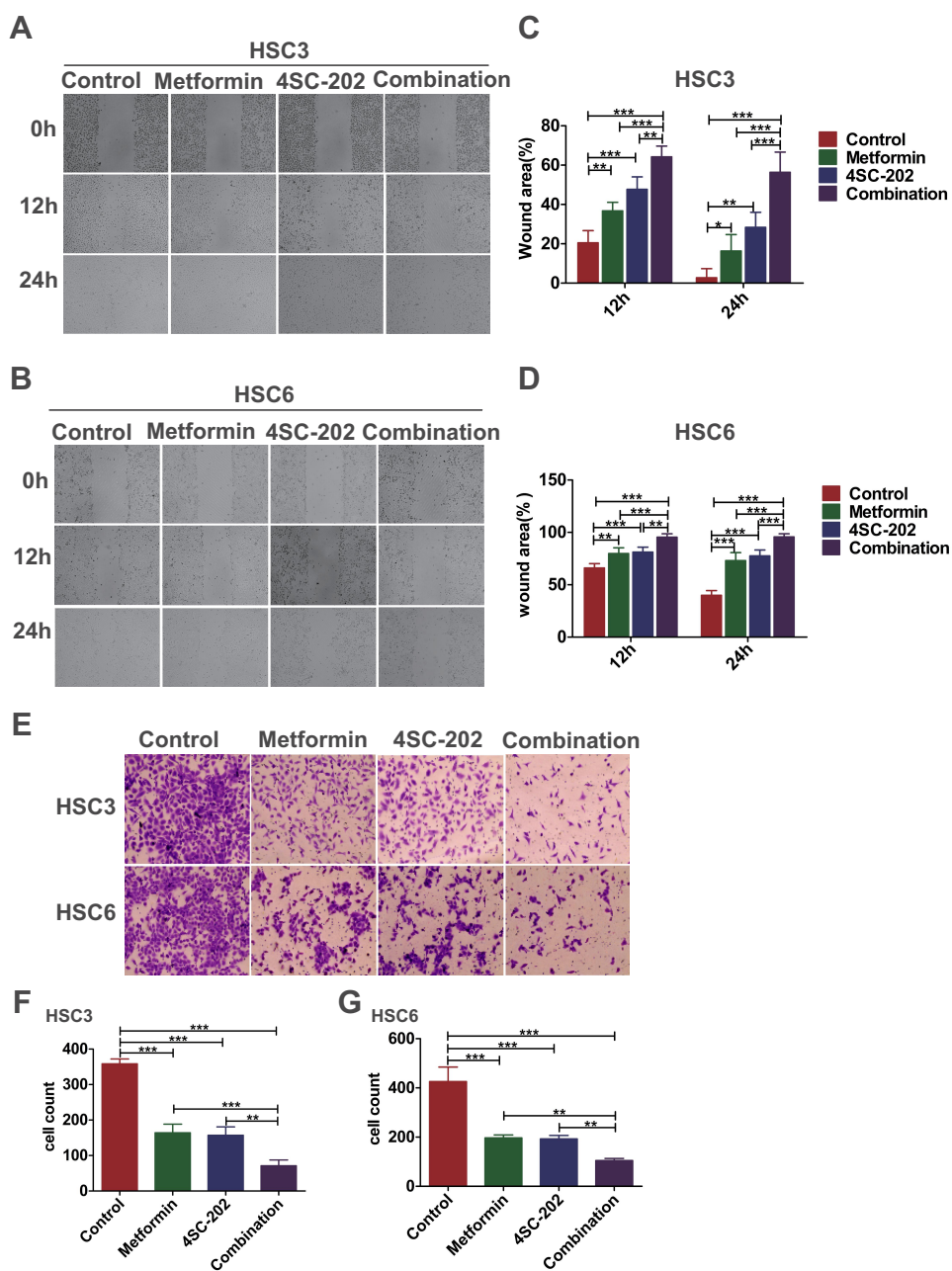
## Results

### Metformin and 4SC-202 Inhibited Invasion and Migration of OSCC

To investigate the effects of metformin and 4SC-202 on migration of OSCC, a wound-healing assay was performed. The results showed that metformin or/and 4SC-202 inhibited the migration of OSCC (Figure 1A and B), and the combination had the most significant inhibition. With metformin or/and 4SC-202 treatment, the wound area of HSC3 was increased from 20.53±5.33% to 36.74±3.79% (metformin), 47.68±5.49% (4SC-202) and 64.22±4.74% (combination) at 12 h and from 2.80±3.97% to 16.37±7.24% (metformin), 28.35±6.64% (4SC-202) and 56.39±8.86% (combination) at 24 h vs control ( $P < 0.05$ ), and the wound area of HSC6 was increased from 66.05±3.59% to 79.78±4.76% (metformin), 81.21±4.06% (4SC-202) and 95.731±2.80% (combination) at 12 h and from 39.92±3.88% to 73.10±6.58% (metformin), 77.52±4.86 (4SC-202) and 95.41±2.55% (combination) at 24 h vs control ( $P < 0.05$ ) (Figure 1C and D). Subsequently, transwell results revealed that metformin or/and 4SC-202 reduced the counts of the invading cells and the combination had the most dramatic reduction (Figure 1E). The counts of HSC3 cells invading to lower chamber were 164.33±19.67 with metformin alone, 157.33±19.29 with 4SC-202 alone, 71.00±13.37 with metformin and 4SC-202 treatment, and 358.33±11.32 in control group ( $P < 0.01$ ) (Figure 1F). Like HSC3, the counts of HSC6 cells invading the lower chamber were 196.67±9.98 with metformin, 192.67±11.6 with 4SC-202 and 104.67±7.04 with combination of metformin and 4SC-202, while the control group was 425.67±48.24 ( $P < 0.01$ ) (Figure 1G).

### Metformin and 4SC-202 Suppressed the Expression of TWIST1

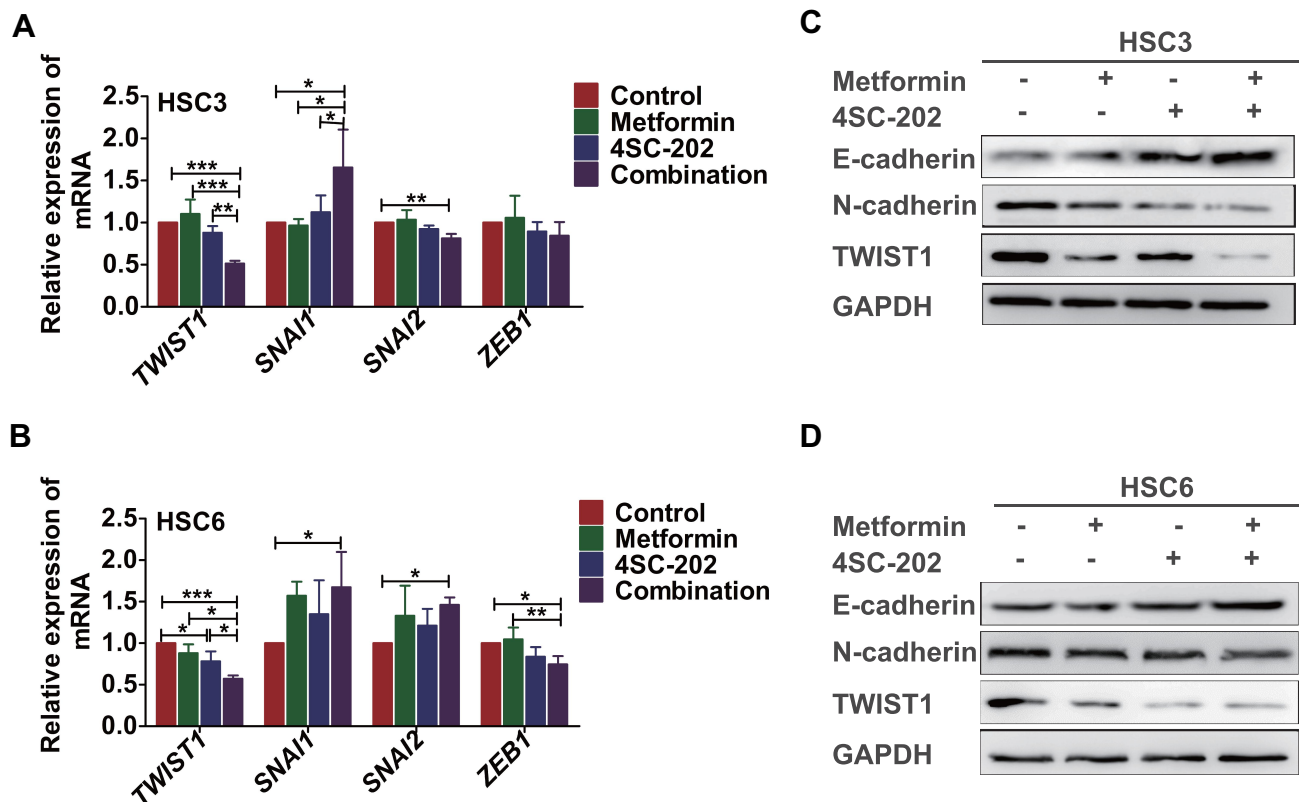
EMT plays vital roles in the invasion and migration of cancer cells,<sup>18</sup> so we used qRT-PCR to determine the expression of EMT-related transcription factors including *TWIST1*, *SNAIL*, *SNAI2*, and *ZEB1*. The results showed that under the



**Figure 1** Metformin and 4SC-202 combination inhibited the migration and invasion of OSCC. OSCCs were treated with 16 mM metformin or/and 0.4  $\mu$ M 4SC-202 for 72 h. **(A, B)** Wound-healing assay was performed to investigate the migration ability of OSCC (original magnification  $\times 50$ ). **(C, D)** Quantification of wound area in HSC3 or HSC6 cells (one-way ANOVA). **(E)** Transwell assay was utilized to investigate the invasion ability of OSCC (original magnification  $\times 100$ ). **(F, G)** Quantification of invasion cells in HSC3 or HSC6 cells (one-way ANOVA). Data are shown as mean  $\pm$ SD ( $n=3$ ). \* $P<0.05$ , \*\* $P<0.01$ , \*\*\* $P<0.001$ .

treatment of metformin and 4SC-202, the relative expression level of *TWIST1* decreased to  $0.51 \pm 0.01$  ( $P<0.001$ ) of control in HSC3 and  $0.57 \pm 0.03$  ( $P<0.001$ ) of control in HSC6. However, the expression of *SNAIL*, *SNAI2*, and *ZEB1* remained unperturbed or even increased (Figure 2A and B). Subsequently, the protein level of mesenchymal protein N-cadherin, epithelial protein E-cadherin and TWIST1 was

evaluated by WB. These results revealed that metformin or/and 4SC-202 inhibited the expression of N-cadherin and TWIST1, but promoted the expression of E-cadherin in HSC3 and HSC6 cells, and the combination had the most significant effects (Figure 2C and D). Therefore, we speculated that TWIST1 may be the main target of metformin and 4SC-202.



**Figure 2** Metformin and 4SC-202 combination inhibited the expression of TWIST1 in OSCC cells. OSCCs were treated with 16 mM metformin or/and 0.4  $\mu$ M 4SC-202 for 72 h. (A, B) The mRNA levels of EMT transcriptional factors *TWIST1*, *SNAI1*, *SNAI2* and *ZEB1* were determined by qRT-PCR (one-way ANOVA). (C, D) The protein levels of E-cadherin, N-cadherin and TWIST1 were determined by WB. GAPDH was used as internal control. Data are shown as mean  $\pm$ SD (n=3). \* $P$ <0.05, \*\* $P$ <0.01, \*\*\* $P$ <0.001. The "+" symbol indicated the treatment of the given drug.

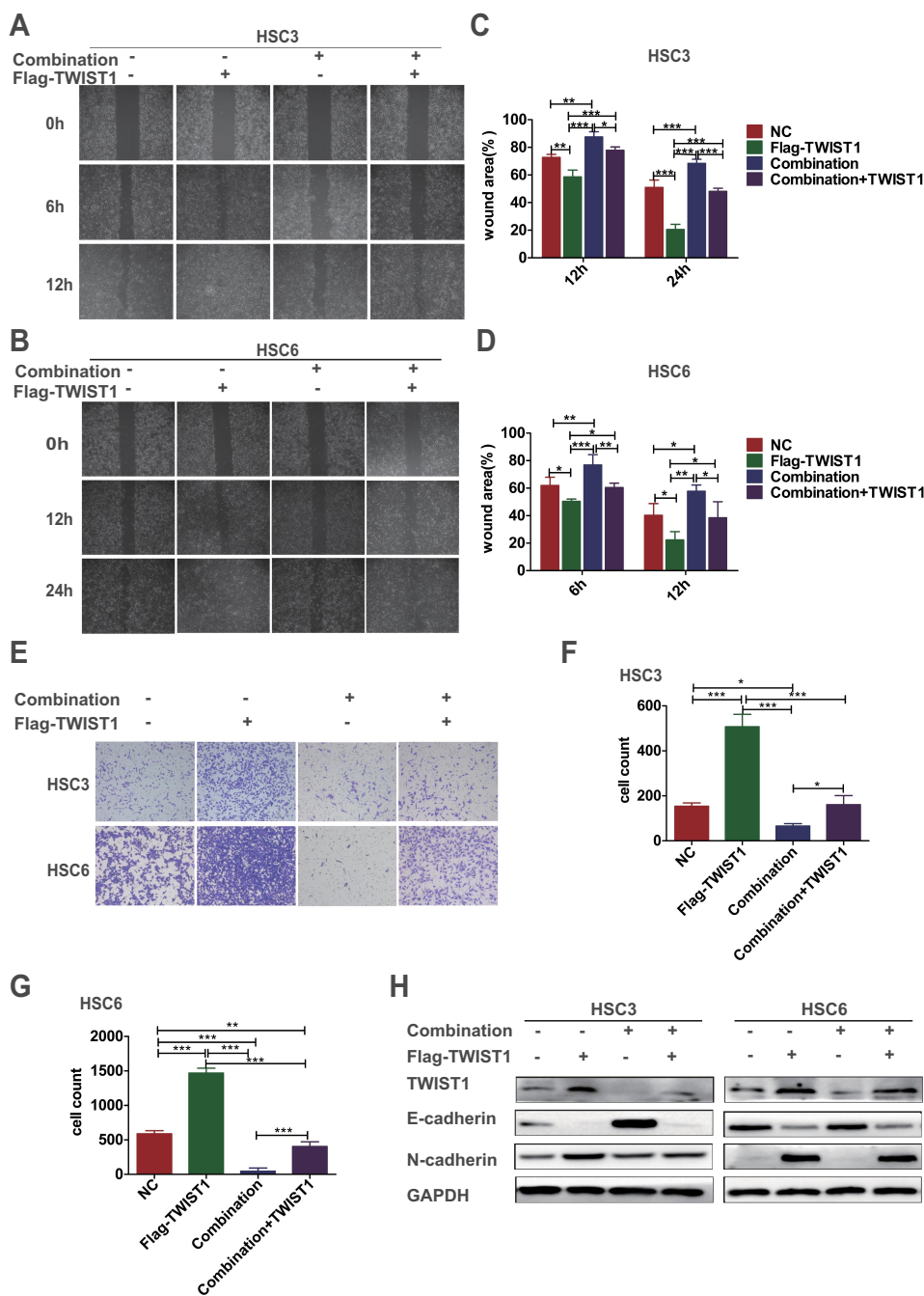
## Metformin and 4SC-202 Suppressed the Invasion and Migration by Inhibiting TWIST1

The efficiency of overexpression and knockdown of TWIST1 in OSCC was evaluated (Supplementary Figure 1). In line with previous reports, overexpression of TWIST1 promoted the invasion, migration and EMT of OSCC cells, and knockdown of TWIST1 had the reverse effects (Supplementary Figure 2). TWIST1 overexpression attenuated the invasion, migration and EMT-inhibiting effects of 4SC-202 and metformin, as the wound area was decreased from  $57.74 \pm 3.68\%$  to  $38.49 \pm 9.50\%$  (12 h) ( $P$ <0.05) in HSC3 cells (Figure 3A and C), and from  $68.37 \pm 2.48\%$  to  $48.15 \pm 1.84\%$  (24 h) ( $P$ <0.001) in HSC6 cells (Figure 3B and D), the counts of cells invading the lower chamber were increased from  $66.00 \pm 8.84$  to  $161.00 \pm 33.22$  ( $P$ <0.05) in HSC3 cells (Figure 3E and F), and from  $46.00 \pm 38.68$  to  $404.00 \pm 56.70$  in HSC6 cells (Figure 3G). Subsequently, the expression of E-cadherin and N-cadherin was determined by WB. The results revealed that TWIST1 overexpression partly rescued the expression of N-cadherin and impeded the downregulation of E-cadherin (Figure 3H).

Overall, our results indicated that TWIST1 was the main target of metformin and 4SC-202.

## Metformin and 4SC-202 Suppressed STAT3 Signaling

As the mRNA changed resembled to those of the protein level, TWIST1 could be controlled at the transcriptional level. STAT3 signal plays vital roles in the proliferation, invasion and migration of HNSCC,<sup>19–22</sup> and STAT3 could regulate TWIST1 expression directly by binding to its promoter.<sup>23</sup> The phosphorylation level of STAT3 was determined by WB, as the results showed, p-STAT3 was reduced after metformin or/and 4SC-202 treatment, and the combination had more remarkable inhibition compare to each alone (Figure 4A and B). Subsequently, one inhibitor of STAT3, S31-201, was applied at different concentrations to investigate regulatory relations between STAT3 and TWIST1. Our results showed that S31-201 inhibited the expression of TWIST1 and N-cadherin, but promoted the expression of E-cadherin (Figure 4C and D), indicating that TWIST1 and EMT was modulated by STAT3.

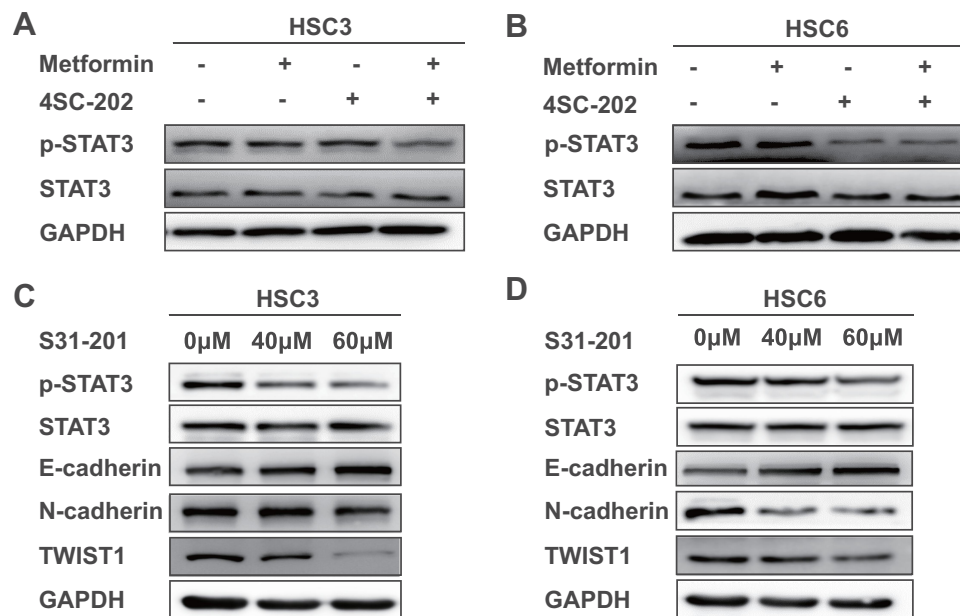


**Figure 3** TWIST1 overexpression attenuated the inhibition of migration and invasion induced by metformin and 4SC-202. **(A, B)** Wound-healing assays were used to determine the migration of OSCC under metformin (16 mM) and 4SC-202 (0.4  $\mu$ M) treatment with TWIST1 overexpression (original magnification  $\times 50$ ). **(C, D)** Quantification of wound areas (one-way ANOVA). **(E)** Transwell invasion assays were performed to investigate the invasion ability of OSCC under metformin (16 mM) and 4SC-202 (0.4  $\mu$ M) treatment with TWIST1 overexpression (original magnification  $\times 100$ ). **(F, G)** Quantification of invasion cells in OSCC (one-way ANOVA). **(H)** WB was utilized to determine the expression levels of E-cadherin, N-cadherin and TWIST1 under metformin (16 mM) and 4SC-202 (0.4  $\mu$ M) treatment with TWIST1 overexpression. GAPDH was used as internal control. Data are shown as mean  $\pm$  SD (n=3). \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ . The "+" symbol indicated the application of the given approach such as drug treatments or lentivirus infection.

## STAT3 Inhibited the Migration and Invasion of OSCC by Regulating TWIST1

Next, the effects of STAT3/TWIST1 on migration and invasion of OSCC cells were examined. The results showed that the migration was decreased significantly after treatment of S31-

201, while overexpression of TWIST1 reversed this inhibitory effect (Figure 5A and B). With S31-201 administration, the wound area of HSC3 cells was increased from  $51.15 \pm 2.90\%$  to  $67.34 \pm 7.30\%$  ( $P < 0.01$ ) at 12 h, and the wound area of HSC6 cells was increased from  $47.74 \pm 2.50\%$  to  $74.38 \pm 5.97\%$



**Figure 4** Metformin and 4SC-202 inhibited the STAT3 signaling. **(A, B)** The phosphorylation level of STAT3 under 16 mM metformin or/and 0.4 μM 4SC-202 treatment for 72 h was determined by WB in HSC3 or HSC6 cells, respectively. **(C, D)** WB was used to determine the phosphorylation level of STAT3 and the expression levels of E-cadherin, N-cadherin and TWIST1 under different concentration (40 μM and 60 μM) of STAT3 inhibitor S31-201 for 48 h in HSC3 or HSC6 cells, respectively. GAPDH was used as internal control. The “+” symbol indicated the treatment of the given drug.

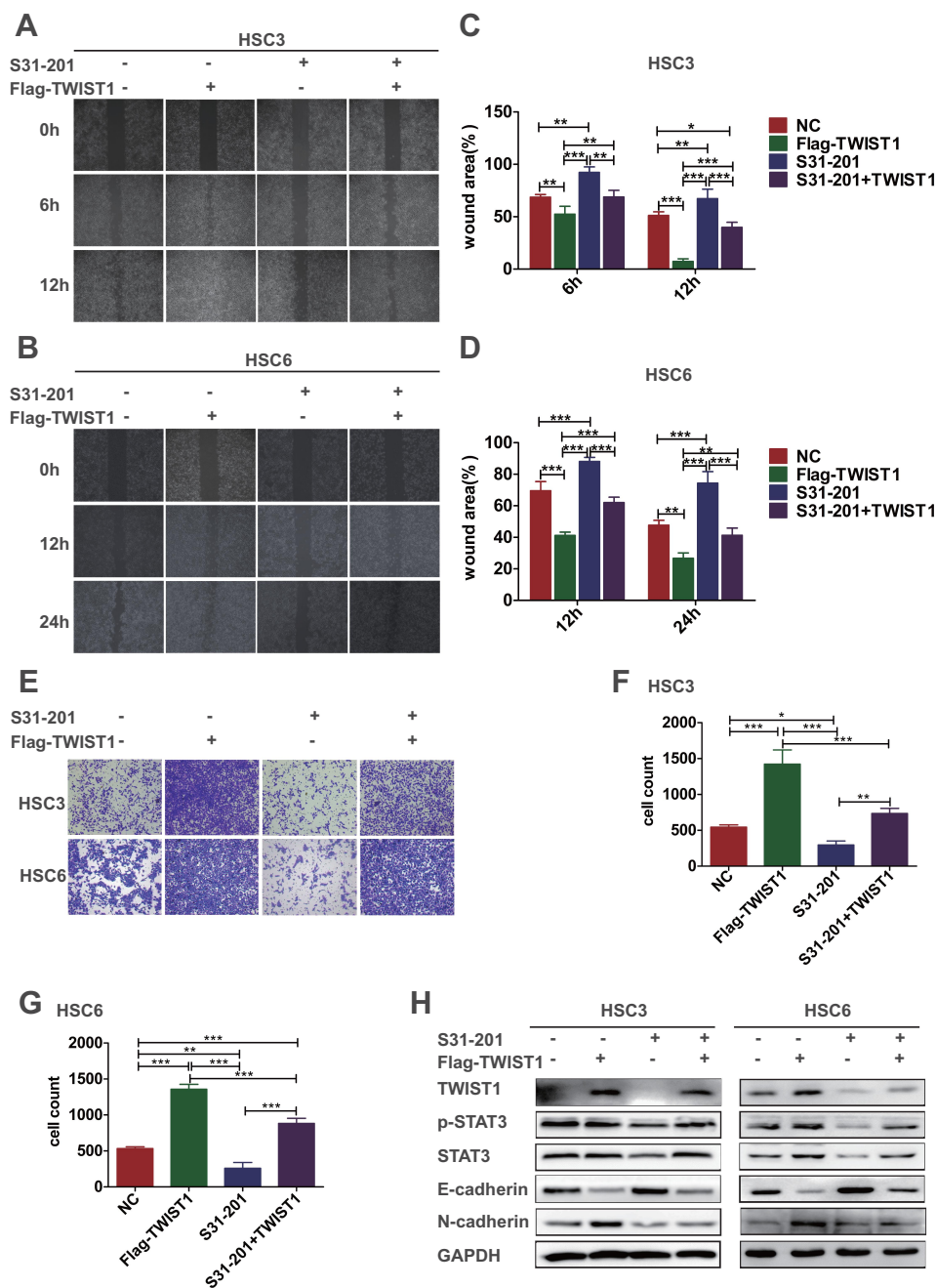
( $P < 0.01$ ) at 24 h. While TWIST1 could reverse this inhibition, as with TWIST1 overexpression, the wound area was decreased to  $39.84 \pm 3.97\%$  ( $P < 0.001$ ) in HSC3 at 12 h and  $41.32 \pm 3.75\%$  ( $P < 0.001$ ) in HSC6 at 24 h (Figure 5C and D). Subsequently, transwell assays were performed to investigate the invasion of OSCC. The results demonstrated that S31-201 inhibited the invasion ability of HSC3 and HSC6, and overexpression of TWIST1 eliminated this inhibition (Figure 5E). The quantitative analysis demonstrated that S31-201 decreased the counts of HSC3 cells invading to lower chamber from  $544.00 \pm 25.66$  to  $295.00 \pm 45.90$  ( $P < 0.05$ ), while TWIST1 overexpression restored it to  $734 \pm 58.70$  ( $P < 0.01$ ), and a decrease of HSC6 cells from  $533.00 \pm 19.38$  to  $259.00 \pm 65.01$  was observed ( $P < 0.01$ ), while TWIST1 overexpression reinstated it to  $883 \pm 58.35$  ( $P < 0.001$ ) (Figure 5F and G). Furthermore, S31-201 was found to reduce the protein level of N-cadherin while it increased the protein level of E-cadherin in HSC3 and HSC6, and TWIST1 overexpression reversed these effects (Figure 5H). According to these data, we could infer that metformin and 4SC-202 inhibited the invasion and migration of OSCC by regulating STAT3/TWIST1.

## Discussion

In present study, our results indicated that combination of metformin and 4SC-202 suppressed the invasion and migration of OSCC. Importantly, the expression of TWIST1 was

inhibited by metformin and 4SC-202, and overexpression of TWIST1 attenuated the invasion and migration inhibiting effects of the drugs. In addition, phosphorylation of STAT3 was reduced by metformin and 4SC-202, and overexpression of TWIST1 reversed the invasion and migration inhibitory effects of STAT3 inhibitor S31-201. Our results revealed that metformin and 4SC-202 synergistically inhibited the invasion and migration of OSCC via STAT3/TWIST1, and could be a promising therapeutic scheme for OSCC.

As mentioned in the literature, HDACi could inhibit the growth of multiple cancer cells. However, the impacts of HDACi on cell invasion and migration seemed to be contradictory because the same agent had a contradictory effect on different cell types or the same cell type had a different response to different HDACi.<sup>24</sup> In hepatocellular carcinoma, class II HDACi TSA administration was found to enhance migration and invasion,<sup>25</sup> while pan HDACi panobinostat had an opposite effect.<sup>26</sup> For head and neck cancer cells, prior studies have noted that SAHA enhanced migration and metastasis in human nasopharyngeal carcinoma cell line,<sup>26</sup> while migration and invasion of nasopharyngeal carcinoma cells were inhibited by TSA.<sup>27</sup> Different HDAC targets, different gene expression profile or side effects may account for these contradictions. In this study, we found that the invasion and migration of OSCC were inhibited by 4SC-202.



**Figure 5** STAT3 inhibited the migration and invasion of OSCC by regulating TWIST1. OSCC cells were divided into four groups with different treatments: NC, with TWIST1 overexpression, with S31-201 (60  $\mu$ M) for 48 h, with TWIST1 overexpression and S31-201 (60  $\mu$ M) for 48 h. (A, B) Wound-healing assays were performed to determine the migration ability of cells under different conditions (original magnification  $\times 50$ ). (C, D) Quantification of wound area (one-way ANOVA). (E) Transwell assays were utilized to investigate the invasion ability of cells under different conditions (original magnification  $\times 100$ ). (F, G) Quantification of invasion cells (one-way ANOVA). (H) WB was used to determine the phosphorylation level of STAT3 and the expression levels of E-cadherin, N-cadherin and TWIST1 under different conditions. GAPDH was used as internal control. Data are shown as mean  $\pm$ SD (n=3). \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ . The “+” symbol indicated the application of the given approach such as S31-201 or lentivirus infection.

Recently, metformin was shown to impede the invasion and migration of several types of cancers including prostate adenocarcinoma, colorectal cancer, gastric cancer, cervical cancer.<sup>28</sup> Several studies have shown that metformin could increase oral cancer cell sensitivity to

chemotherapeutic drugs (such as 5-FU, gefitinib), improve the therapeutic efficacy, and reduce the dose and toxicity.<sup>29,30</sup> Therefore, combination of metformin with other chemotherapeutics could be a potential candidate for the development of new treatment strategies for



OSCC. In line with previous reports, our finding demonstrated that metformin inhibited invasion and migration of OSCC. Importantly, combination of metformin and 4SC-202 synergistically suppressed the migration and invasion of OSCC.

EMT plays a key role in tumor progression and is associated with stemness, invasiveness, drug-resistance, and the ability to form metastases in distant organs.<sup>22</sup> HDACi exerted different effects on EMT in head and neck cancer cells, the EMT progression was induced by TSA,<sup>31</sup> but class I, II HDACi valproic acid suppressed EMT.<sup>32</sup> Previous study of pancreatic cancer indicated that 4SC-202 had negative effects on EMT,<sup>10</sup> and metformin suppressed EMT in various cancer types.<sup>12,14,33,34</sup> In addition, metformin sensitized HNSCC to gefitinib and inhibited EMT.<sup>30</sup> In our study, both 4SC-202 and metformin had negative impact on EMT of OSCC. Interestingly, metformin combined with 4SC-202 resulted in a more pronounced inhibition of tumor EMT compared with either drug alone.

EMT is modulated by TWIST1, a conserved helix-loop-helix (bHLH) transcription factor, which plays pivotal roles in multiple stages of embryonic development<sup>35</sup> and significantly contributes to tumor metastasis.<sup>36</sup> In OSCC, there is positive association between TWIST1 and poor prognosis, and TWIST1 promotes invasion and migration.<sup>37</sup> Our results showed that metformin and 4SC-202 induced the decrease of TWIST1, and when applied together, a more dramatic suppression was observed, while the expression of other EMT transcription factors such as *SNAI2*, *SNAI1*, *ZEB1* remained unperturbed or even increased. Overexpression of TWIST1 attenuated the invasion, migration and EMT inhibitory effects of metformin and 4SC-202, indicating that TWIST1 was the main target of metformin and 4SC-202.

TWIST1 is regulated by multiple signaling pathways including ERK, STAT3 and NF- $\kappa$ B.<sup>38</sup> As a novel prognostic factor in HNSCC, STAT3 contributes to tumor growth,<sup>19</sup> and induces invasion and migration.<sup>20–22</sup> STAT3 was clarified to activate TWIST1 expression by directly binding to its promoter,<sup>23</sup> and STAT3/TWIST1 regulatory axis was shown to be a critical regulator that drove the EMT progress in multiple types of cancer.<sup>39–44</sup> Therefore, STAT3 could be the upstream regulator of TWIST1. In the present study, the inhibition of STAT3 led to a dramatic decrease of TWIST1 level in OSCC, supporting that TWIST1 was a downstream gene of STAT3. We speculated that the expression of TWIST1 was induced by STAT3 via direct transcriptional activation.

Metformin was reported to suppress invasion and migration of several cancers including bladder cancer, prostate cancer and pancreatic cancer through STAT3 inhibition.<sup>12,45,46</sup> HDACi also exerts its tumor inhibitory impact through STAT3, as the class II HDAC inhibitor mercaptoacetamide suppresses migration and invasion in human glioma cells by inhibiting FAK/STAT3,<sup>47</sup> and class I HDAC inhibitor chidamide and BG45 inhibits the viability of MDS and AML cells by suppressing JAK2/STAT3 signaling.<sup>48,49</sup> Thus, STAT3 could be the target of anticancer drugs. Our results revealed that phosphorylation of STAT3 decreased under metformin or/and 4SC-202 treatment, suggesting metformin or/and 4SC-202 may suppress TWIST1 through STAT3, and intriguingly, combination of metformin and 4SC-202 had a more dramatic inhibition than either alone. Metformin activates AMPK, which in turn deactivates mTOR,<sup>28</sup> and mTOR could phosphorylate STAT3 directly to activate it.<sup>50–52</sup> Hence, we speculated that metformin suppressed STAT3 via AMPK activation in OSCC. Class I HDAC HDAC3 forms a complex with STAT3, and HDAC3 inhibition by HDACi represses STAT3 phosphorylation.<sup>53,54</sup> Therefore, a possible explanation for STAT3 inhibition by 4SC-202 was that as selective class I HDAC inhibitor, 4SC-202 suppressed the enzymatic activity of HDAC3, and inhibition of HDAC3 deactivated STAT3. Thus, it could be assumed that metformin and 4SC-202 exerts its STAT3-suppressing effects via a different approach, and further work is required to determine the exact mechanisms.

A more dramatic decrease of TWIST1 was observed under metformin and 4SC-202 treatment, together with STAT3 inhibitor administration, and overexpression of TWIST1 rescued the invasion, migration, and the expression of EMT genes repressed by STAT3 inhibitor. These results demonstrated STAT3 was the key component that linked the anticancer effects of the drugs and TWIST1.

In summary, this study showed that metformin and 4SC-202 synergistically inhibited the invasion and migration of OSCC through STAT3/TWIST1. Our results complemented our previous studies, corroborating metformin and 4SC-202 as a promising therapeutic scheme for OSCC.

## Abbreviations

OSCC, oral squamous cell carcinoma; HDAC, histone deacetylase; HDACi, histone deacetylase inhibitor; EMT, epithelial-to-mesenchymal transition; qRT-PCR, quantitative reverse transcription-polymerase chain reaction; WB, Western blot; ECL, chemiluminescence.

## Acknowledgments

This work was supported by grants from the National Natural Science Foundation of China (no. 81671000, 81870769) and the Science and Technology Program of Guangzhou, China (no. 201803010019, 201607010354).

## Disclosure

The authors report no conflicts of interest in this work.

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