

Snail-induced EMT promotes cancer stem cell-like properties in head and neck cancer cells

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Abstract. Epithelial-mesenchymal transition (EMT) is a key process involved in the invasion and metastasis of cancer cells. Furthermore, EMT can induce a cancer stem cell (CSC)-like phenotype in a number of tumor types. We demonstrated that Snail is one of the master regulators that promotes EMT and mediates cancer cell migration and invasion in many types of malignancies including head and neck squamous cell carcinoma (HNSCC). In the present study, we investigated the role of Snail in inducing and maintaining CSC-like properties through EMT in HNSCC. We established HNSCC cell lines transfected with Snail. Stem cell markers were evaluated with real-time RT-PCR and western blot analysis. CSC properties were assessed using sphere formation and WST-8 assays as well as chemosensitivity and chick chorioallantoic membrane *in vivo* invasion assays. Introduction of Snail induced EMT properties in HNSCC cells. Moreover, Snail-induced EMT maintained the CSC-like phenotype, and enhanced sphere formation capability, chemoresistance and invasive ability. These data suggest that Snail could be one of the critical molecular targets for the development of therapeutic strategies for HNSCC.

Introduction

Epithelial-mesenchymal transition (EMT) is an important process in tumor progression that causes epithelial cells to

acquire a migratory mesenchymal phenotype (1,2). EMT is thought to be a critical step in the induction of cell invasion and tumor metastasis (1). Furthermore, it has been shown that cells with an EMT phenotype are more resistant to chemoradiotherapy in regards to head and neck squamous cell carcinoma (HNSCC) (3-6).

It has been indicated in recent studies that EMT induces cancer stem cell (CSC)-like properties in many different types of malignant tumors and that both EMT and a CSC-like phenotype are associated with treatment resistance (7-9). Prince *et al* demonstrated that the purified CD44⁺ population of HNSCC cells possesses the self-renewing properties of CSCs (10). Aldehyde dehydrogenase 1 (ALDH1) has also been shown to be a putative marker of CSCs in HNSCC (11). Furthermore, Chen *et al* showed that CD44⁺/ALDH1⁺ cells resist radiotherapy and may serve as a reservoir for developing tumors and metastasis (6). Additionally, the CD44^{high}/EGFR^{low} subpopulation exhibited the EMT phenotype and resistance to treatment in HNSCC (12). Moreover, a high level of expression of variant isoforms of CD44 (CD44v) could suppress the clustering of EGFR at the surface of HNSCC cells and thereby negatively regulate EGFR signaling in the absence of a differentiation stimulus, suggesting that CD44v-negative HNSCC cells rely on EGFR activity for survival (13).

However, the source and the mechanism of development in regards to CSCs have not yet been fully elucidated. Moreover, it has been shown that it is difficult to extract pure CSC populations with only CSC markers such as CD44 and ALDH1. The same result is shown with side populations (14). Therefore, it is necessary not only to identify novel CSC markers but also to clarify the mechanism of CSC development.

Snail, a member of the zinc-finger transcription factor family, plays an important role in EMT by directly repressing epithelial markers such as E-cadherin and by upregulating mesenchymal markers (7,15-19). Several studies have shown that Snail-related transcription factors play a transcriptional and regulatory role in the invasion, metastasis and poor outcome in different type of malignancies, including HNSCC (20,21). These findings suggest that Snail expression may regulate CSC-like properties via EMT in HNSCC.

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We previously demonstrated that Snail overexpression induced EMT, including cancer cell migration and invasion, and promotes CSC-like phenotype such as CD44⁺/ALDH1⁺ in head and neck cancer cells (22). However, the key role of Snail on the stemness of CSC in HNSCC has not been fully elucidated. In the present study, we first demonstrated that Snail-induced EMT gains CSC-like phenotype such as upregulation of stem cell markers, including CD44 and ALDH1, and enhanced CSC-like properties such as sphere formation capability, chemoresistance and *in vivo* cancer invasion and metastasis in HNSCC.

Materials and methods

Cell lines and culture. Human HNSCC cells, SAS and HSC-4, were employed in the present study. SAS and HSC-4 cells, obtained from the Japanese Cancer Research Resource Bank (Tokyo, Japan), were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% heat-inactivated fetal bovine serum (FBS) (both from Invitrogen, Carlsbad, CA, USA), 100 U/ml penicillin and 100 mg/ml streptomycin (Gibco, Grand Island, NY, USA) at 37°C in 5% CO₂.

Transient transfection. The cDNA fragment encoding human Snail (NM_005985.2) was inserted into the pCR 3.1 mammalian expression vector (Invitrogen). For transient transfection, SAS and HSC-4 cells (1.5x10⁵ cells) were plated into 6-well culture plates and allowed to adhere for 12 h. Then, the cells were transfected with 2 µg of either pCR 3.1-Snail or pCR 3.1-vector (without insert DNA) with Lipofectamine 2000 reagent (Invitrogen) according to the manufacturer's instructions. We established SAS-Snail and HSC-4-Snail as Snail-expressing cell lines and their respective control cell lines.

Immunoblot analysis and antibodies. Total protein extracts were prepared according to the freeze-thawing lysis method, and protein concentrations were measured with a bovine serum albumin (BSA) protein assay. Samples of extract containing 20 µg of protein were then separated by SDS-PAGE and transferred to polyvinylidene difluoride membranes. After washing with phosphate-buffered saline with Tween-20 (PBST), the membranes were incubated first with rabbit anti-Nanog, rabbit anti-Oct4, rabbit anti-ABCG2, rabbit anti-EGFR and rabbit anti-pEGFR (Cell Signaling Technology, Danvers, MA, USA; diluted 1:1,000), rabbit anti-Sox2 and rabbit anti-Bmi1 (Abcam, Cambridge, MA, USA; diluted 1:1,000) antibodies at 4°C overnight and then with peroxidase-conjugated secondary anti-rabbit antibody or goat immunoglobulin G (IgG) (diluted 1:1,000; Cell Signaling Technology) for 1 h. After rinsing in PBST, immunodetection was accomplished using an ECL western blot analysis detection reagent and analysis system. The membranes were subsequently exposed to X-ray film as previously described (23).

Real-time RT-PCR. Total RNA was isolated from each cell line using the RNeasy Mini kit (Qiagen, Hilden, Germany). cDNA was then reverse transcribed using ReverTra Ace[®] qPCR RT Master Mix with gDNA Remover (Toyobo Life Science, Tokyo, Japan) according to the manufacturer's instructions. Real-time RT-PCR was then carried out using the

Table I. Primers for real-time RT-PCR.

Gene	Primer sequence
<i>ABCG2</i> (NM_004827)	S 5'-CATGTACTGGCGAAGAATATTTGGT-3' A 5'-CACGTGATTCTTCCACAAGCC-3'
<i>Bmi1</i> (NM_005180)	S 5'-AAATGCTGGGAACTGGAAAG-3' A 5'-CTGTGGATGAGGAGACTGC-3'
<i>Nanog</i> (NM_024865)	S 5'-ATTCAGGACAGCCCTGATTCTTC-3' A 5'-TTTTTGCACACTCTTCTCTGC-3'
<i>Oct4</i> (NM_002701)	S 5'-GTGGAGAGCAACTCCGATG-3' A 5'-TGCTCCAGCTTCTCCTTCTC-3'
<i>Sox2</i> (NM_003106)	S 5'-CGAGTGGAAACTTTTGTCGGA-3' A 5'-TGTGCAGCGCTCGCAG-3'
<i>GAPDH</i> (NM_002046)	S 5'-CATCATCCCTGCCTCTACTG-3' A 5'-GCCTGCTTACCACCTTC-3'

S, sense; A, antisense.

primers shown in Table I and EXPRESS SYBR[®]-GreenER[™] qPCR SuperMix with premixed ROX (Invitrogen). PCR was performed with an initial step of 3 min at 95°C followed by 40 cycles of 3 sec at 95°C and 20 sec at 60°C. The level of target mRNA was normalized to the mRNA level of GAPDH as an internal standard.

Sphere formation assay. The capability of self-renewal was assessed using 96-well NanoCulture plates (Scivax, Tokyo, Japan). Cells (1x10⁴) were seeded and cultured for 1 week in DMEM supplemented with 10% FBS or serum-free medium, and phase-contrast images were obtained.

Chemotherapy of the cultured cells. Chemosensitivity was assessed using the Cell Counting Kit-8 (WST-8 cleavage; Dojindo, Mashikimachi, Japan) as previously described (23). The cells were seeded into 96-well plates at an initial density of 4x10³ cells/well and incubated for 24 h. For chemotherapy, cisplatin (Nihon Kayaku Co., Tokyo, Japan) at a concentration of 1.0 µM was added to each well. Following incubation for an additional 48 h, 10 µl of WST-8 solution was added to each well, and the plate was incubated for a further 2 h. The absorbance of each well at 450 nm (reference wavelength at 620 nm) was measured by a Multiscan FC microplate photometer (Thermo Scientific). The measurement was repeated at least three times for each cell line.

Chick chorioallantoic membrane (CAM) *in vivo* invasion assay. The CAM *in vivo* invasion assay was conducted using 11-day-old chick embryos wherein HSC-4 or the transfected cells (10⁵ cells labeled with green Fluoresbrite carboxylated polystyrene nanospheres of 45-nm diameter; Polysciences) were seeded atop the CAM and incubated for three days as previously described (24,25). The CAM was dropped without damaging the epithelial basement membrane (BM) by applying gentle negative pressure at the air sac, and an opening of ~1 cm² was cut in the shell above the CAM with an electric drill.

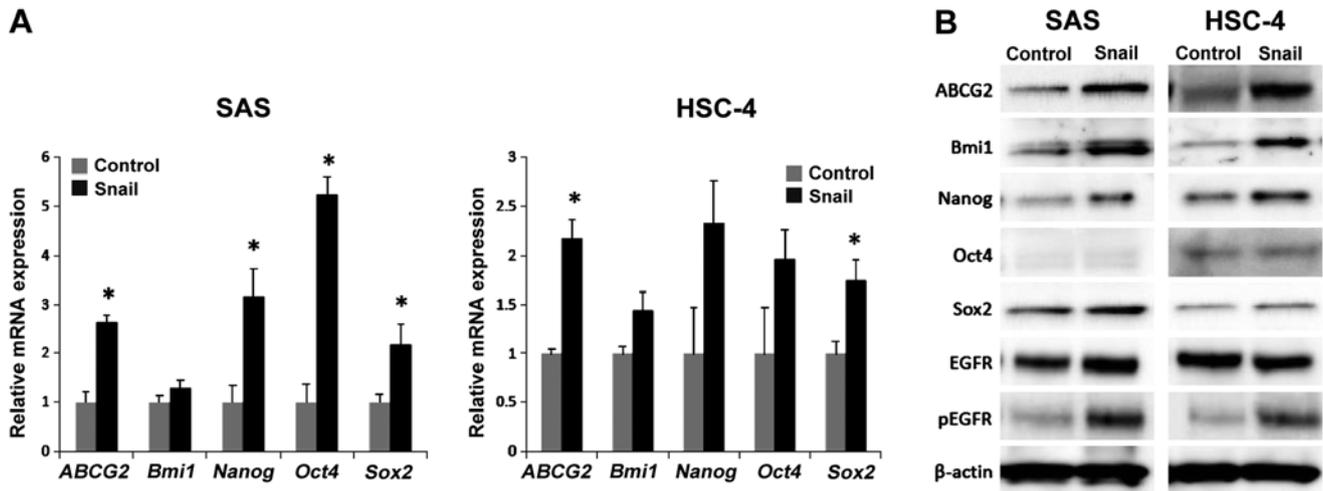


Figure 1. Snail induces expression of several stem cell markers in SAS and HSC-4 cells. (A) Relative expression of mRNA-encoding stem cell genes ABCG2, Bmi1, Nanog, Oct4 and Sox2 normalized to the endogenous control GAPDH. (B) The level of each protein was determined by western blot analysis with β -actin used as the internal control. Data represent the mean \pm SE. * $P < 0.05$.

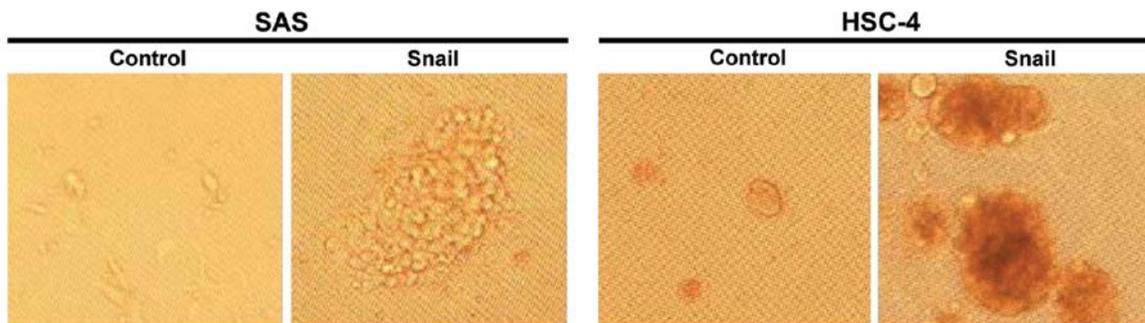


Figure 2. Transfection of Snail increases tumor sphere-forming capability. Tumor sphere-forming capability was assessed by a sphere formation assay. Cells were seeded and cultured for 1 week.

Statistical analysis. Data are presented as the mean \pm standard error (SE). Experimental differences between groups were assessed with a t-test. The differences were considered to indicate a statistically significant result at $P < 0.05$.

Results

Snail expression induces a CSC-like phenotype. To elucidate that EMT by Snail expression induces a stem cell-like phenotype in HNSCC cells, we investigated the expression of CSC surface markers in HNSCC cells. We previously demonstrated that expression of both CD44 and ALDH1 increased in SAS-Snail and HSC-4-Snail cells compared with their control cells (22). RT-PCR analysis revealed that mRNA levels of several stem cell markers, Bmi1, Nanog, Oct4, Sox2 and ABCG2, which are drug-resistant proteins, were high in the Snail-transfected cells compared with the levels in the controls (Fig. 1A). Moreover, at the protein level, Nanog, Bmi1 and ABCG2 were also upregulated in the Snail-transfected cells, whereas the expression of Oct4 and Sox2 showed no difference between the Snail-expressing cells and the controls (Fig. 1B). These data revealed that Snail-induced

EMT elicits a CSC-like phenotypic change as CD44⁺/ALDH⁺, and directly regulates the expression of Nanog, Bmi1 and ABCG2. In addition, there were no differences in the level of EGFR protein between the Snail-transfected cells and the controls, whereas, the levels of phospho-EGFR protein increased in the Snail-transfected cells compared with that in the control (Fig. 1B).

Snail induces CSC properties. The transfection of Snail induced tumor sphere-forming capability in the SAS and HSC-4 cells, but not in the control cells (Fig. 2). Furthermore, the cells transfected with Snail showed significantly low chemosensitivity at 1.0 μ M of cisplatin, as compared with the control cells (Fig. 3). Thus, these data revealed that the acquisition of a CSC-like phenotype caused by Snail-induced EMT results in enhancement of the ability of sphere formation and chemoresistance in the HNSCC cells.

Snail induces EMT as a CSC property in vivo. We previously demonstrated that Snail-regulated EMT promotes CSC properties, including cell migration and invasion as well as E-cadherin suppression in HNSCC cells *in vitro* (22). Snail was

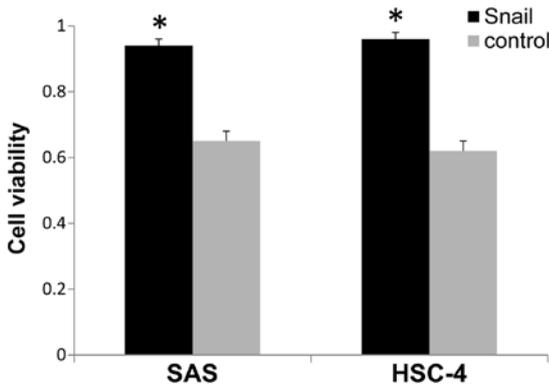


Figure 3. Snail expression enhances the chemoresistance to cisplatin in SAS and HSC-4 cells. Cell viability was assessed using WST-8 assay for chemosensitivity. Cells were incubated for 24 h. For chemotherapy, the cells were treated with 1.0 μ M of cisplatin. Following an additional 48 h, cell viability was determined by WST-8 assay. Data are shown as the mean \pm SE. *P<0.05.

also functional in inducing *in vivo* cancer invasive progression of non-invasive SAS and HSC-4 cells, as noted in the crossing of cells into the BM in the CAM assay (Fig. 4).

Discussion

The identification of cell surface-specific markers of CSCs is critical for the possible establishment of target-specific cancer therapies. CD44 (10) and ALDH1 (11) have been reported to represent candidate markers of CSCs in HNSCC. Recently, CD271 (26) and CD10 (27) have been identified as additional

markers of CSCs in HNSCC. However, it is difficult to extract CSCs selectively with such markers only. Several studies have indicated that there could be key markers of CSCs that regulate stemness genes, such as Bmi1, Nanog, Oct4, Sox2 and ABCG2 as follows.

Bmi1 has been demonstrated to play a role in the tumorigenesis of HNSCC (28,29). Expression of Bmi1, ALDH1 and Snail could be associated with the maintenance of stemness in CSCs and correlate with poor overall survival in HNSCC patients (30). Additionally, Bmi1 can regulate Snail and ALDH1 in inducing EMT and CSC properties (30). In the present study, we obtained results consistent with those in our previous study (22).

Nanog maintains the pluripotency of embryonic stem cells and functionally blocks differentiation (31,32). Recent studies have demonstrated that Nanog promotes CSC properties, and the downregulation of Nanog inhibits sphere formation and tumor development (33-36). Another study suggested that Nanog was upregulated by TGF- β through Smad signaling and that Snail directly regulates Nanog promoter activity (37). Nanog expression was also upregulated in the present study, whereas there were no significantly differences in the expression of Oct4 and Sox2 between Snail-transfected cells and the controls. A previous study showed that Oct4 and Sox2 regulate Nanog expression (38). Therefore, direct regulation against the Nanog promoter by Snail may result in the suppression of Oct4 and Sox2.

Yoshikawa *et al* showed that the expression of EGFR was low in CSCs, but high in non-CSCs. They suggested that a high level of expression of variant isoforms of CD44 (CD44v) could suppress the clustering of EGFR at the surface

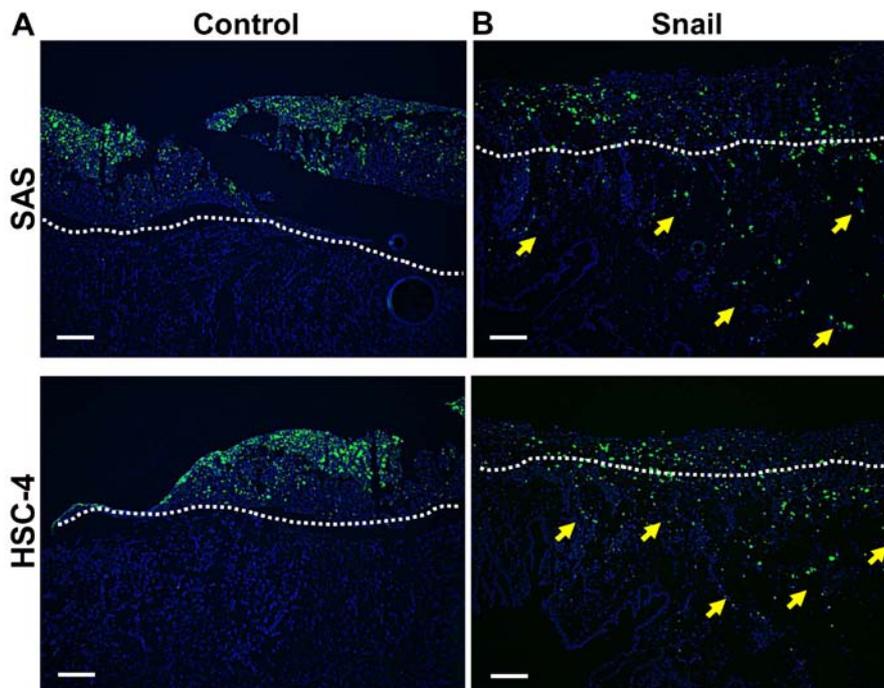


Figure 4. Snail expression increases *in vivo* cell invasion in head and neck cancer cells. The non-invasive SAS and HSC-4 cells were transiently co-transfected with either a control (A) or Snail (B) labeled with green fluorescent nanospheres, respectively. The cells were cultured atop the embryonic chick chorioallantoic membrane (CAM) for 3 days and the fixed tissue sections then were examined by fluorescence microscopy. The upper face of the CAM is indicated by dashed white lines; invasive cancer cells (green) are denoted by yellow arrows. Scale bars, 200 μ m.

of HNSCC cells and thereby negatively regulate EGFR signaling in the absence of a differentiation stimulus, and that CD44v-negative HNSCC cells rely on EGFR activity for survival (13). It has been suggested that the suppression of differentiation and treatment resistance in CSCs could be attributed to CD44v-positive and EGFR-negative expression. In the present study, Snail overexpression induced CSC-like properties, while there was no significant difference in the expression of EGFR between the Snail-transfected cells and the controls. Furthermore, the expression of phospho-EGFR was enhanced in the Snail-transfected cells compared with the controls. However, this result is considered to be compatible with the findings in the previous study. It was predicted from the present study that the CSC-like properties acquired by Snail occurred not in all cells, but in a portion of them. This raises the possibility that upregulation of phospho-EGFR occurred in non-CSCs around CSCs. Wang *et al* showed that reduction in E-cadherin resulted in upregulation of EGFR transcriptionally and activation of EGFR resulted in overexpression of Snail (39). It was suggested that Snail could not only induce CSC-like properties, but also phosphorylates EGFR in non-CSCs, and it contributed to the maintenance of the microenvironment by interaction between CSCs and non-CSCs. Although the precise involvement of CD44 and EGFR expression by Snail in the regulation of CSCs remains to be elucidated, they may be involved in a latent effect. The critical mechanisms remain to be further investigated. The majority of studies on CSCs have been performed *in vitro* to evaluate clinical cancer therapies. Therefore, it should be noted that the results of *in vitro* studies on CSCs cannot be translated to the same cells *in vivo*. Although it is still too early to discuss its clinical efficiency, these data could support the hypothesis that the interaction between CSCs and non-CSCs contributes to cell proliferation and treatment resistance.

In addition, the upregulation of ABCG2 expression was also observed in the Snail-transfected cells. However, higher expression of ABCG2 was observed not in ALDH1⁺ cells, but in ALDH cells around CSCs (40). This suggests that ABCG2 itself is not a CSC marker, but the interaction between CSCs and the cells around CSCs results in upregulation of ABCG2. The upregulation of ABCG2 in the present study could similarly support the possibility that Snail contributes to the interaction between CSCs and non-CSCs.

Furthermore, we first showed that Snail regulates cancer cell invasion through EMT as a CSC-like property in HNSCC cells *in vivo* as well as *in vitro* (22). These data suggest that Snail can promote cancer invasion and metastasis as well as maintain the stemness similar to CSCs.

In summary, we demonstrated the possibility that Snail induces CSC-like properties through EMT and maintains stemness by upregulating various CSC markers. However, other transcription factors of EMT as well as Snail could be involved in the maintenance of CSCs and the microenvironment. These interactions between various CSC- and EMT-relating genes make it difficult to select a pure CSC population. Snail-induced EMT is also considered to play an essential role in tumor progression, such as cancer invasion and metastasis *in vivo*. Although the critical mechanisms remain to be further investigated, Snail could prove to be one of the valid targets in the treatment for HNSCC.

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