

# SDF-1-CXCR4 axis: Cell trafficking in the cancer stem cell niche of head and neck squamous cell carcinoma

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**Abstract.** Stromal cell-derived factor-1 $\alpha$  (SDF-1 $\alpha$ ), also known as CXCL12, has variable effects on a plurality of cells. CXCR4 has been identified as its corresponding receptor. The SDF-1-CXCR4 axis is postulated to be a crucial key pathway in the interaction between (cancer) stem cells and their surrounding supportive cells in the cancer stem cell niche. We evaluated the expression of CD44 as a cancer stem cell marker and of CXCR4 in human HNSCC tissue samples. Afterwards, we monitored the concentration of SDF-1 in peripheral blood samples of HNSCC patients and healthy donors. We showed that CD44 and CXCR4 are expressed in human HNSCC tissues. Markedly, CD44 showed a high expression in HNSCC cells bordering cancer stromal cells. CXCR4 was mainly expressed in HNSCC tumor nests, but not in the surrounding stromal cells. No significant difference was noted between the SDF-1 concentration in the peripheral blood of HNSCC patients compared to healthy donors. We showed that CD44, as a stem cell marker in HNSCC, is located mainly at the borderline of HNSCC tumor nests with the surrounding cells. In addition, we demonstrated that CXCR4 as the corresponding receptor to SDF-1 is highly expressed in HNSCC tumor nests, but not in the tumor stroma. We collected evidence that SDF-1-CXCR4 interaction may be a crucial pathway in cell trafficking in the cancer stem cell niche of HNSCC, while SDF-1 was not detected in the peripheral blood of HNSCC patients. The SDF-1-CXCR4 axis may play an important role in the cancer stem cell theory of HNSCC. As SDF-1 $\alpha$  also exhibits a multitude of functional effects on HNSCC cells, such as migration and polarization, it may be

possible that the SDF-1-CXCR4 axis is also involved in the pathophysiology of the progression, recurrence and metastasis of malignant disease. Understanding these interactions may help to gain further insight into these mechanisms and as such help to discover new strategies of therapy.

## Introduction

Although advances have been made in the surgical and non-surgical therapy of head and neck squamous carcinoma (HNSCC), the mortality rate from this disease has remained nearly constant over the last few years. This is mainly due to the development of therapy-resistant local and regional recurrences (1). Strategies of treatment apart from surgery, such as chemotherapy and radiation, eradicate a majority of proliferating cells in malignant tumors, but there is increasing evidence that there is a subpopulation of resistant tumor cells that cannot be reached by these regimens called cancer stem cells (CSCs) (2). As a result, it is imaginable that these cells are essential and responsible for initiation, but also maintenance and recurrence of malignant disease. CSCs have features of somatic stem cells such as self-renewal, differentiation and extensive proliferation ability. In recent years, the CSC hypothesis has also been assigned to HNSCC (3,4). Prince *et al* showed that CD44<sup>+</sup> cells that were isolated out of the bulk of an HNSCC tumor, but not the CD44<sup>-</sup> cancer cells, gave rise to new tumors in a mouse model. These cells typically comprise <10% of the cells in an HNSCC tumor (3). Other markers have been evaluated as cancer stem cell markers in HNSCC as well, such as ALDH1 (5) and CD133 (6). Current research aims to discover a combination of including and excluding surface markers to finally isolate the 'real CSC in HNSCC'.

CD44 is an integral cell membrane glycoprotein. It comprises different isoforms that arise from alternative splicing of a region of variable exons. Its apparent molecular mass ranges from 85 to 250 kDa, as its isoforms differ in the primary amino acid sequence and the amount of N- and O-glycosylation (7). At least 20 variants of CD44 have been reported due to the alternative splicing of 10 exons that encode the membrane's proximal portion of the extracellular domain (8-10).

In 1991, Günthert *et al* (11) and Hofmann *et al* (12) showed that the expression of CD44 gave metastatic potential to

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*Abbreviations:* HNSCC, head and neck squamous cell carcinoma; CSCs, cancer stem cells; SDF, stromal cell-derived factor

*Key words:* head and neck squamous cell carcinoma, cancer stem cells, stromal cell-derived factor, CXCR4, stem cell niche

a non-metastatic cell line in a rat carcinoma model. Several analyses have indicated that there is a correlation between the expression of CD44 and progression, metastasis and prognosis of malignant disease and this has been shown in different types of epithelial malignancies, e.g. gastric cancer (13), hepatocellular carcinoma (14) or gynecological malignancies such as breast (15) or ovarian cancer (16). The in-depth analysis of expression markers such as CD44 in tissue samples of HNSCC patients may reveal their role as potential prognostic biomarkers or therapeutic targets, e.g. for antigen-directed immunotherapy.

The analysis of the cancer stem cell niche theory, where CSCs come into contact with their supportive cells surrounding them, may provide information concerning cell trafficking and underlying mechanisms, such as tumor expansion, recurrence and metastatic progression. The interaction between SDF-1 $\alpha$  and its receptor CXCR4 may play an important role in this field. SDF-1 $\alpha$  is a multifunctional cytokine that is expressed and secreted by several tissues, e.g. endothelium and stromal cells (17,18), and is one component of the bulk of an HNSCC tumor. SDF-1 has a single open reading frame of 282 nucleotides encoding a polypeptide of 93 amino acids. It arises in two forms, SDF-1 $\alpha$  (amino acids 24-88) and SDF-1 $\beta$  (amino acids 24-93) by alternative splicing (18-20). SDF-1 $\alpha$  is the only proven chemoattractant for hematopoietic progenitor cells (HPCs) to date (18,21-23). Thus, SDF-1 $\alpha$  is considered to be one of the key regulators for HPC trafficking between the peripheral circulation and the bone marrow (17,18,23,24). Faber *et al* and others have demonstrated that SDF-1 $\alpha$  induces polarization and podia formation in HPCs and leukemic cells (18,25), two properties that represent prerequisites for directed locomotion. SDF-1 $\alpha$  alone showed a moderate effect on cell proliferation in CD34<sup>+</sup> cells (18,26), and its effect on survival or apoptosis of HPCs remains controversial (18,26-28). Furthermore, the SDF-1-CXCR4 axis plays a crucial role in the regulation of homing and adhesion to the supportive cellular microenvironment in the hematopoietic stem cell niche (29). It remains unclear whether this interaction is also important in the cancer stem cell niche of malignant epithelial tumors such as HNSCC.

Signal transduction pathways initiated by the binding of SDF-1 $\alpha$  to CXCR4 are not fully understood. Mechanisms involved in CXCR4 signaling and downstream are multifaceted and include Gi-protein-mediated activation of intracellular components (30).

Herein, we monitored the expression of CD44 as a cancer stem cell marker and of CXCR4 as a potential target of interaction between CSCs and their supportive cells in the cancer stem cell niche in human HNSCC tissue samples. Accordingly, we evaluated SDF-1 serum levels in the peripheral blood of HNSCC patients compared to healthy donors to find evidence whether soluble interactions in the cancer stem cell niche of an HNSCC tumor can be detected in the periphery of human blood circulation. These findings may facilitate the development of therapeutic strategies particularly aimed at CSCs or particularly with regard to the SDF-1-CXCR4 axis aimed at the interaction of CSCs with their cellular cancer stem cell niche. Differences in the concentration of SDF-1 in the peripheral blood of HNSCC patients compared to healthy humans might lead to new strategies of tumor detection and the role of SDF-1 as a tumor marker.

Table I. HNSCC tissue collection.

Patient	Gender	Age (yrs.)	Tumor location	TNM status
1	M	71	Larynx	T4N0
2	M	68	Larynx	T3N2b
3	M	63	Larynx	T4N0
4	M	NA	Oropharynx	T3N2c
5	M	67	Oropharynx	T4N2M0
6	F	50	Oropharynx	T3N2c
7	M	76	Hypopharynx	NA
8	M	59	Hypopharynx	T4N2b
9	F	63	Hypopharynx	T2N2b

M, male; F, female; NA, information not available.

## Materials and methods

**Tissue and peripheral blood sample collection.** A total of 9 HNSCC tissue samples from tumor patients was selected from a tissue database collected from 1997 to 2010 at the Department of Otorhinolaryngology, Head and Neck Surgery at the University of Mannheim. Samples were fixated immediately after excision by freezing in liquid nitrogen. All samples were confirmed by pathology after H&E staining. The histological and clinical characteristics of all tumor samples are summarized in Table I.

In addition to the tissue collection, peripheral blood samples were obtained from HNSCC patients and healthy donors. Peripheral blood was collected before, during, but never after tumor surgery. A group of 11 healthy blood donors served as a control. Blood samples were centrifuged at 2500 rpm for 10 min. Afterwards, serum samples were harvested and fixated by freezing at -80°C. Characteristics of the peripheral blood samples are summarized in Table II.

**Immunofluorescence labeling.** To detect the expression of CD44 and CXCR4 in HNSCC tissue samples, tumors underwent fixation by freezing in liquid nitrogen as mentioned above. Specimens were sectioned (5- $\mu$ m), air-dried and fixated in acetone for 10 min. Afterwards, the sections were treated with 4% paraformaldehyde (PFA) for 10 min at room temperature. After three washing steps with PBS, tumor samples were treated with 1% serum (goat) for another 10 min. Sections were then incubated with the CD44/CXCR4 antibody (mouse monoclonal, 1:100; Abcam, Cambridge, UK) for 1 h at 37°C followed by incubation with a secondary biotinylated antibody (anti-mouse, 1:100) for 30 min. After further washing steps with PBS, the sections were treated with Streptavidin-Cy3 (1:1000)/Streptavidin-Alexia 488 (1:500) for 30 min at room temperature. Subsequently, sections were stained with DAPI after washing with PBS. Finally, slices were covered in FluorSave and dried to be evaluated by fluorescence microscopy.

**Enzyme-linked immunosorbent assay.** Serum levels of SDF were measured with a human SDF ELISA kit (R&D Systems,

Wiesbaden, Germany). A monoclonal antibody against soluble SDF was adsorbed to microwells in 96-well microtiter plates. Samples, including those with standards with known SDF concentrations, were pipetted into these wells. During the first incubation, the SDF antigen was added to the wells. After washing, a biotinylated monoclonal antibody specific for SDF was incubated, and the enzyme (streptavidin-peroxidase) was added. Following incubation and washing to remove all unbound enzymes, a substrate solution was added, which catalyzed a reaction on the bound enzyme and induced a colored reaction product. The intensity of this colored product is directly proportional to the concentration of SDF present in the samples.

**Statistical analysis.** All results are plotted as mean  $\pm$  standard deviation. To estimate the probability of differences, we adopted the Student t-test (two-tailed distribution, two-sample equal variance). A probability value  $<0.05$  was considered to denote a statistically significant result.

## Results

**Tissue and peripheral blood sample collection.** A total of 9 HNSCC tissue samples and the 29 blood serum samples were selected from a database collected from 1997 to 2010 at the Department of Otorhinolaryngology, Head and Neck Surgery at the Faculty of Medicine Mannheim, University of Heidelberg. Tissue samples used in the study were derived from 2 female and 7 male patients aged 50-76 years (mean age, 65 years). Blood samples were derived from 5 female and 24 male patients aged 48-76 years (mean age, 60 years). Sites of the primary tumors were the larynx, oropharynx, hypopharynx and oral cavity. Patient characteristics are summarized in Table I for tissue sample collection and in Table II for the blood sample collection. A group of 11 healthy blood donors, aged 26-87 years (mean age, 50 years) served as a control for experiments concerning the SDF concentration in peripheral blood of HNSCC patients.

**Expression of CD44 in HNSCC tissues.** Immunofluorescence labeling was performed to measure the expression of CD44 in HNSCC tissue samples via staining with Cy3. In HNSCC tissue samples, expression of CD44 was established in all cases. Its expression was mainly found in the cells forming the invasive front of the tumor. This invasive front is in direct contact with the tumor-surrounding stromal cells, and this interaction can be postulated as the cancer stem cell niche of HNSCC (Fig. 1A.4). Tumor samples stained in our experiments were mainly obtained from the superficial portions of the HNSCC tumors, although it was noted that parts located in the center of the tumor had a weaker staining pattern for CD44 (Fig. 1A.4). In general CD44 exhibited a membranous staining pattern (Figs. 1A.2 and A.3, and 2B.1). Nuclei were stained with DAPI.

**Expression of CXCR4 in HNSCC tissues.** Immunofluorescence labeling was performed to measure the expression of CXCR4 in HNSCC tissue samples by staining with Alexia 188. CXCR4 was present in all tissue samples stained. CXCR4 exhibited a surface staining pattern as well as cytoplasmic expression (Fig. 2B.2 and B.3). Specific expression of CXCR4

Table II. HNSCC patients enrolled for blood sample collection.

Patient	Gender	Age (yrs.)	Tumor location	TNM
1	M	54	Larynx	T4N2xMx
2	F	55	Larynx	T1N0
3	M	62	Larynx	T4N0
4	M	70	Larynx	T4N0
5	M	68	Larynx	T3N2b
6	M	68	Larynx	NA
7	M	55	Larynx	T4N2b
8	M	74	Larynx	T4N0
9	M	59	Larynx	NA
10	M	57	Larynx	T4N1
11	M	55	Oral cavity	T4N1
12	M	69	Oral cavity	T1N2b
13	F	54	Oral cavity	T1N2b
14	M	48	Oral cavity	NA
15	M	49	Oral cavity	T3N1
16	M	51	Oral cavity	NA
17	M	66	Oral cavity	T2N0
18	F	50	Oropharynx	T3N2c
19	M	64	Oropharynx	NA
20	M	60	Oropharynx	T4N2c
21	M	64	Oropharynx	T3N2
22	F	55	Oropharynx	T4N3
23	M	61	Hypopharynx	T4N2b
24	M	66	Hypopharynx	T2N0
25	M	76	Hypopharynx	NA
26	M	59	Hypopharynx	T4N2b
27	F	62	Hypopharynx	T2N2b
28	M	51	Hypopharynx	NA
29	M	57	Hypopharynx	T3N2c

M, male; F, female; NA, information not available.

at the invasive front of the tumor was not noted, while CXCR4 expression was high in the tumor nests, but was not noted in the tumor surrounding stroma (Fig. 2B.4). Thus, it was demonstrated that the corresponding receptor to SDF, which is transmitted by the supportive stromal cancer stem cell niche, can be found in the tumor nest as the counterpart of this interaction. Nuclei were stained with DAPI.

**Concentration of SDF-1 in the peripheral blood of HNSCC patients.** ELISA analysis was performed to evaluate the concentration of SDF-1 in the peripheral blood of HNSCC patients according to tumor site compared to a healthy control group. There was no significant difference in the SDF-1 concentration between tumor patients and the control group. Values are listed as means: control, 2057.91 pg/ml; larynx, 2111.50 pg/ml; oral cavity, 1891.43 pg/ml; oropharynx, 2244.2 pg/ml; hypopharynx, 2000.29 pg/ml.

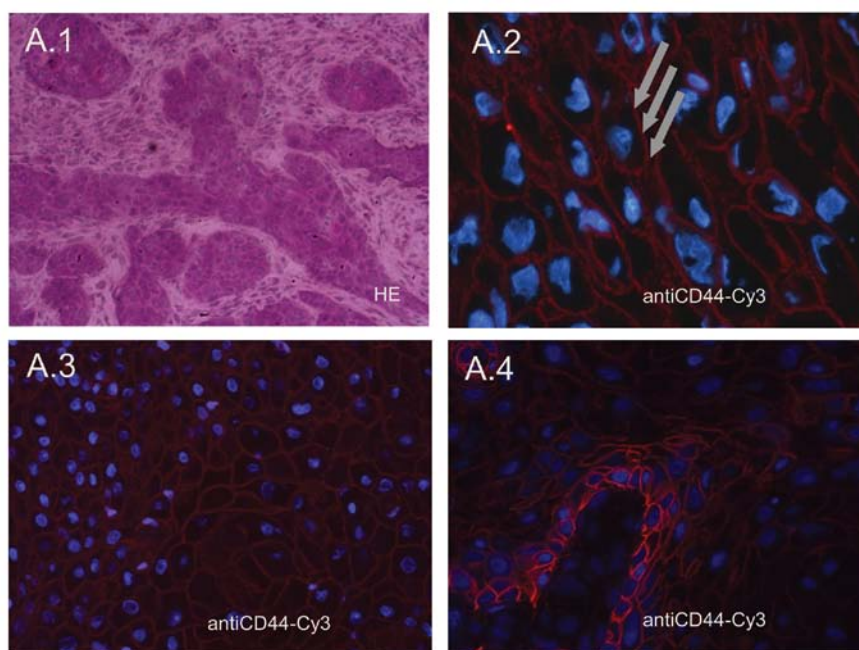


Figure 1. Expression of CD44 in HNSCC. Identification of the cancer stem cell niche. After verification of pathology by H&E staining (A.1), immunofluorescence labeling was performed to measure expression of CD44 in HNSCC tissue samples via staining with Cy3 (A.2-4). Arrows indicate the expression of CD44 marked red by Cy3. In the HNSCC tissue samples stained, expression of CD44 was established and was mainly noted in the cells forming the invasive front of the tumor (A.4). Nuclei were stained with DAPI.

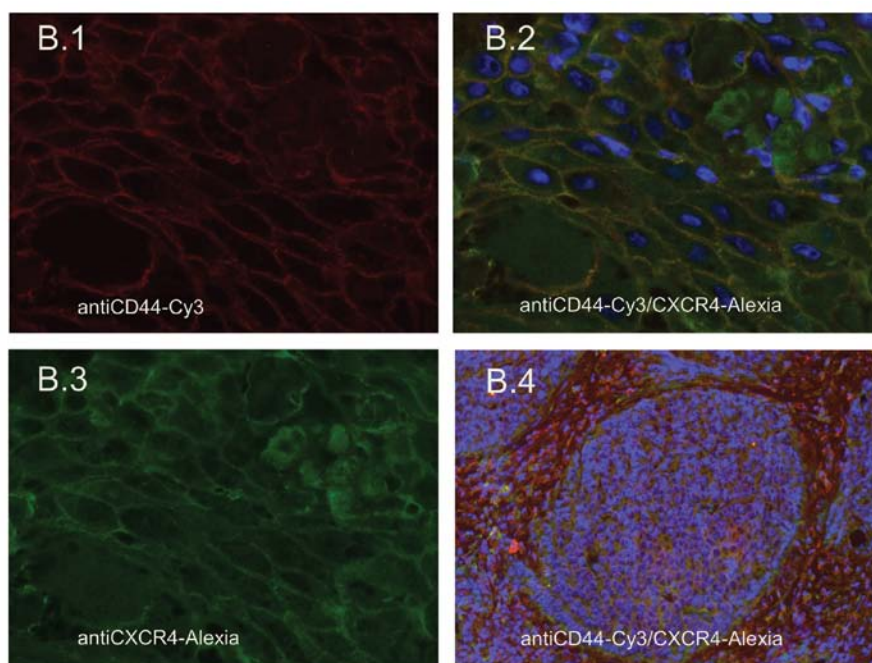


Figure 2. Expression of CD44 and CXCR4 in HNSCC tissue samples. Interactions in the cancer stem cell niche. CD44 exhibited a membranous staining pattern (red in B.1). CXCR4 was also expressed in the cytoplasm (green in B.3). CXCR4 was mainly expressed in the tumor nests, but not in the supportive cells surrounding the tumor (green in B.4). Nuclei were stained with DAPI.

## Discussion

In the past few years, the stem cell theory has become more important in the field of tumor biology, particularly in issues concerning tumor development, progression and metastasis. The cancer stem cell theory has generated new ideas and

discussion in the research of malignant disease and in therapeutic options. At present, the presence of cancer stem cells are not only known to exist in hematological malignancies but also in solid tumor entities (3,31). The cancer stem cell theory in solid tumors such as HNSCC has had dramatic consequences. To date, therapeutic interventions, e.g. surgery

or chemoradiation, are directed towards the bulk of the tumor without focusing on a small amount of specialized tumor cells which have the facilities of self-renewal, differentiation and unlimited proliferation. The question of which combination of markers should be used to isolate 'cancer stem cells' e.g. in HNSCC remains unclear. In addition, the combination of markers appears to vary in different types of tumors (3,31,32). In the hematopoietic system, CD34 has been shown to play an important role in this context (18), and in acute myeloid leukemia, CD34<sup>+</sup> cells were able to establish a new tumor in a mouse model (33). Regarding solid tumor entities, CD133 has been postulated to play an important role in malignancies of the nervous system, and results showed that CD133<sup>+</sup> cells are relatively radioresistant (34) which can also be assessed as a feature of cancer stem cells. In contrast, CD133 is a ubiquitous marker in the hematopoietic and endothelial progenitor cell system (35), and it has also been shown to be present in the neurologic system, e.g. in glioblastoma (36).

In HNSCC, the presence of cancer stem cells has also been postulated. Prince *et al* showed that CD44<sup>+</sup> cells, compared to CD44<sup>-</sup> cells, were able to engraft an entire new HNSCC tumor in a mouse model (3). According to their results, the cell selection for CD44<sup>+</sup> was sufficient to isolate cells with cancer stem cell properties out of the bulk of an HNSCC tumor (3). Unfortunately, CD44 is also expressed in ordinary cells (35). The prevalent challenge of research is to identify a combination of surface markers that can identify the subgroup of potent CSCs from the bulk of a tumor more precisely. CD44 cannot be sufficiently used as a defining CSC marker alone, since as a cell surface glycoprotein it takes part in numerous cell-cell interactions (35,37) and in humans it is present in a multitude of splice variants (35) and can be found ubiquitously. Approaches to the investigations of the 'real cancer stem cell' in HNSCC requires research with human tumor material with isolation of a small subpopulation of cells out of the bulk of the tumor using complicated cell separation steps including recurrent washing steps, lysis by DNases and FACS-sorting. To concur with the current standard of knowledge, a combination of lineage markers must be negative to separate CSCs out of the entire HNSCC tumor (CD3, CD3, CD10, CD18, CD31, CD64, CD140) (3,35,38). Other CSC markers (CD44, ALDH1) have been proposed, but only small amounts of cells carrying these markers are said to be able to initiate an HNSCC tumor in a mouse model (3,5). The cell amount remaining after this separation as described above is often minute and the remaining cells are often not useful for further experiments. Often it is necessary to revert to cell lines to perform experiments. ALDH1 is another marker postulated as a cancer stem cell marker in HNSCC (5). It was also postulated as a cancer stem cell marker in breast (39) and lung cancer (40). It was shown that high expression of ALDH1 in breast cancer is associated with poor prognosis and early metastasis (41). It is an enzyme that catalyzes the oxidation of aromatic aldehydes into carboxylic acids and it has been shown to be responsible for the resistance of progenitor cells to chemotherapeutic agents by breaking down cytotoxic drugs (35,42).

In a present study concerning HNSCC, ALDH1 was declared to be a more specific cancer stem cell marker in comparison to CD44 (5). In the present study, only CD44 was used as a cancer stem cell marker, and we found a high

expression of CD44 in all tissue samples tested especially at the invasive front of the tumor, where HNSCC cells come into contact with their surrounding cells, e.g. stromal cells. Our findings corroborate Sterz *et al* (43) who showed colocalization of CD44, ALDH1 and CK14 in the basal layer of HNSCC and at the border of the tumor at the tumor-stroma border. The localization of cancer stem cell candidates at the border of tumors is feasible as this is where invasion and the main tumor growth occurs. Highly differentiated cell layers were negative for these markers, and this agrees with the supposition that stem cells are localized close to the basal membrane. Invasiveness and metastasis of tumors also depend on their capacity to penetrate and rebuild the extracellular matrix (44). Malignant cells infiltrate healthy tissue by degradation of the extracellular matrix components, by breaking down vessel borders and by promoting metastasis in distant organs. It has been shown for different tumor entities, that the presence of matrix metalloproteinases plays an important role in this process and this has also been shown for HNSCC (43,45). In HNSCC, MMP-9 and -2 have been shown by Sterz *et al* to be located at the invasive front of HNSCC (43).

The SDF-CXCR4 axis is involved in several aspects of tumor progression, such as angiogenesis, metastasis and survival (46). The microenvironment of the bone marrow has been reported to support survival, differentiation and proliferation of hematopoietic progenitor cells (47), but also malignant progenitor cells of the hematopoietic system, e.g. B-cell acute lymphoblastic leukemia (B-ALL) (48). The pathway which includes the SDF-1-CXCR4 axis has been postulated to be responsible for retention of lymphoid and myeloid leukemia cells in the bone marrow (48,49). It becomes obvious that the importance of the SDF-1-CXCR4 axis in the hematopoietic system is well-discussed, but this is the first time that the SDF-1-CXCR4 axis has been reported to play a crucial role in the development and the progression of invasion and metastasis of HNSCC.

In this study, we showed that CXCR4 can be found in tumor nests of HNSCC, but not in the surrounding stromal region of the cancer stem cell niche. It has been shown by Clatot *et al* that the intratumoral level of SDF-1 is correlated with survival in HNSCC (50). In contrast, we showed in this study that the concentration of SDF-1 in the peripheral blood of HNSCC patients does not differ in comparison to healthy donors. This suggests that the SDF-1-CXCR4 axis plays a role in the cancer stem cell niche within the tumor, but not in the periphery of the blood system. In previous studies, we showed that polarization and formation of filopodia and a prominent uropod were increased in the CD44<sup>+</sup>CXCR4<sup>+</sup> HNSCC cell line UM-SCC 11A in a dose-dependent manner by SDF-1 $\alpha$ . This effect can probably be attributed to cytoskeleton rearrangements of actin-containing protrusions (18,51), and this also might be influenced by extracellular factors such as matrix metalloproteinases (51,52). Podia formation was found to occur together with cell adhesion especially to the microenvironment (18). If evidence can be found that podia formation and adhesion to the cellular cancer stem cell niche are also associated with HNSCC CD44<sup>+</sup> cells, understanding of these interactions would offer insight into new strategies of cancer-directed therapy in HNSCC. For example small-molecule agonists or antagonists of SDF-1 may be

used to interfere with the cancer stem cell niche resulting in inhibition or ideally blockage of tumor invasion and metastasis (18). Further experiments using human material are warranted to expand and specify our insight regarding the cell-cell interactions in the cancer stem cell niche of solid tumors. Based on the findings it may be possible to develop particular therapeutic strategies aimed at CSCs or the SDF-1-CXCR4 axis.

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