Chemokine receptor 7 enhances cell chemotaxis and migration of metastatic squamous cell carcinoma of head and neck through activation of matrix metalloproteinase-9

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Abstract. The mechanisms leading to squamous cell carcinoma of head and neck (SCCHN) metastasis are not fully understood. Although evidence shows that the chemokine receptor 7 (CCR7) and its ligand CCL19 may regulate tumor dissemination, their role is not clearly defined in SCCHN. Matrix metalloproteinases break consisting of tissue barrier to the surrounding tissue invasion and metastasis by destroying the balance of matrix degradation of the basement membrane of tumor cells and extracellular matrix (ECM). We used chemotaxis and migration assays, western blotting, gelatin zymography, actin polymerization assay, immunofluorescence staining and immunohistochemical analysis to explore whether MMP-9 can be activated by CCL19 (CCR7’s ligand) and its role in SCCHN. The experiments were performed in the metastatic SCCHN cell line PCI-37B after pre-incubation of the cells with CCL19 and SB-3CT (inhibitor of MMP-9). Our results demonstrated that CCR7 favors PCI-37B cell chemotaxis and migration, upregulation of MMP-9 protein and motivates the activity of MMP-9 protein, induces reorganization of the actin cytoskeleton and upregulation of MMP-9 protein. SB-3CT can block all these effects. Collectively, our data indicated that CCR7 regulates cell chemotaxis and migration via MMP-9 in metastatic SCCHN, and these results provide a basis for new strategies in preventing metastases of SCCHN.

Introduction

Squamous cell carcinoma of head and neck (SCCHN), a malignant tumor of epithelial origin, represents >90% of all head and neck cancers (1). In SCCHN, the 5-year survival rate is only 30-40%. The mechanisms which lead to the high level of malignancy, particularly metastasis to lymph node and low survival rate, are not completely understood, despite substantial improvements in the diagnosis and local management of, and chemotherapy for, SCCHN (2).

Chemokines are a superfamily of small chemoattractant cytokines that mediate their effects by binding to G-protein-coupled receptors. Chemokines are classified into 4 highly conserved groups: CXC, CC, C and CX3C, based on the position of the first two cysteines adjacent to the amino terminus. The CC chemokine receptor 7 (CCR7) plays a central role in regulating migration of lymphocytes to lymph nodes, such as in mature dendritic cells (DCs) (3), T and B cells (4); flow of calcium ions, changes of cytoskeletal structure, cell cycle and cell metabolism. Since both metastasis and normal migration of leukocytes involve site-directed movement across vascular barriers, non-small cell lung cancer (5,6), gastric (7) and colorectal carcinoma (8) and SCCHN (1,2,9,10) cells also use chemokine-mediated mechanisms during the metastatic process (11). However, the signaling mechanisms mediated by CCR7 and induced by CCL19 have yet to be elucidated in SCCHN cells.

The matrix metalloproteinases (MMPs) are a family of zinc-dependent proteases that are responsible for proteolytic degradation of specific extracellular matrix (ECM) components (12). MMP-9 is one of them, known as gelatinase or type IV collagenase. They are able to degrade IV collagen, which is associated with invasion and metastasis of tumor. Many researchers report high expression of MMP-9 in several types of migration and invasion cancers, such as SCCHN (13), prostate cancer (14), Hodgkin’s lymphoma (15), papillary thyroid carcinoma (16) and brain tumor (17). Thus, MMP-9 is considered the maker of metastasis and invasion of SCCHN to malignant tumor.

Recently, a new role was described for the chemokine family: signaling via chemokine receptors can modulate tumor
cell expression of MMP-9 which can then facilitate adhesion of cancer cells to and/or invasion through ECM. CCR7-induced MMP-9 expression is an important regulatory factor (18). The expression of MMP-9 was enhanced in a variety of malignant tumors, cultured tumor cells and oncogene transformed cells. In vitro migration assay confirmed that the high invasive ability of tumor is associated with the high expression of MMP-9 (19-21).

In the present study, we showed for the first time a direct relationship between CCR7 and MMP-9 in SCCHN cells. We also identified a novel mechanism for CCR7-induced invasion and migration, mediated by regulating MMP-9 in SCCHN.

Materials and methods

SCC cell lines in the head and neck. Metastatic SCCHN cell line PCI-37B, which strongly expresses CCR7, was supplied by the University of Pittsburgh Cancer Institute, USA. PCI-37B were cultured in Dulbecco's modified Eagle's medium (DMEM; Invitrogen, Carlsbad, CA, USA). DMEM was supplemented with 10% fetal bovine serum (FBS; Gibco, Carlsbad, CA, USA), 100 U/ml penicillin G and 100 U/ml streptomycin.

Antibodies and reagents. CCL19 and CCR7-specific monoclonal antibody (mouse anti-human CCR7 antibody), MMP-2 and -9, bovine serum albumin (BSA) and fibronectin (FN), were purchased from R&D Systems (Minneapolis, MN, USA). SB-3CT (the inhibitor of MMP-9), and dimethyl sulfoxide (DMSO) were purchased from Sigma (St. Louis, MO, USA). Matrigel™ Basement Membrane Matrix was purchased from BD Biosciences Pharmingen (Rockville, MD, USA).

Cell chemotaxis assay. The chemotaxis was assayed in Transwell filter insert chambers (10-µm pore size; Corning Costar) as previously described (25). CCL19 (final density, 500 ng/ml) was placed in the lower wells. PCI-37B cells treated with or without SB-3CT (inhibition of MMP-9) and anti-CCR7 mAb in different concentrations or for different times, at 37°C in 5% CO₂, were removed from the culture flasks and added to the upper chamber. After 24 h at 37°C, cells on the upper surface of the Transwell membrane were wiped off gently with cotton swabs. The lower surface of the filters was fixed with methanol and stained with hematoxylin. Cells that migrated to the lower surface were counted under a microscope (Nikon Eclipse 80i; Nikon, Tokyo, Japan) at x200 magnification. For each experimental condition, 4-5 wells were analyzed in parallel.

Cell migration assay. The methods were the same as for the chemotaxis assay, the changes are cells were seeded onto the Matrigel-coated filter and incubated in serum-free medium with 500 ng/ml CCL19 for 36 h. Non-invading cells on the upper side of the filters were removed with a moistened cotton swab. Cells that penetrated the membrane were fixed with ice-cold methanol, stained with 0.5% crystal violet, photographed and counted under the microscope. To assess cellular migration potential, the protocol described above was used, except that Matrigel was omitted. For each experimental condition, 4-5 wells were analyzed in parallel.

Western blotting. PCI-37B cells were treated with and without the inhibitors SB-3CT and CCR7 mAb, followed by CCL19. Whole cells were harvested in lysis buffer, containing sodium diphosphate, sodium trivanadium oxygen, Tris hydrochloric acid, 1% Triton X-100, protease and phosphatase inhibitors. Then, lysates were centrifuged at 4°C, 14,000 rpm, for 30 min after sonication for 3 sec. Protein concentration of the lysate was determined by Bio-Rad protein assay dye reagent (Bio-Rad Laboratories, Richmond, CA, USA). Proteins were size-fractionated on 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to nitrocellulose filters by semi-dry blotting. The filter was blocked in phosphate-buffered saline (PBS) containing 1% skim milk, 0.1% Triton X-100, sodium chloride and Tris [Tris (hydroxymethyl) aminomethane] overnight at 4°C. The membrane was incubated with 1/1,000 diluted rabbit antibody SB-3CT (inhibition of MMP-9) for 30 min at room temperature, and incubated with horseradish peroxidase-conjugated secondary antibodies (goat anti-rabbit; Sigma). Immune complexes were visualized using enhanced chemiluminescence (ECL; Amersham Pharmacia Biotech, Piscataway, NJ, USA). Quantification of the signals was carried out by scanning densitometry using FluorChem software (version 2.0), β-actin (1:1,000) served as the internal control. For each experimental condition, 4-5 wells were analyzed in parallel.

Actin polymerization assay. PCI-37B cells pretreated with or without CCR7 mAb and MMP-9 inhibitor SB-3CT were fixed, permeabilized and stained with TRITC-labeled phalloidin. Following labeling, the samples were washed three times for 10 min each in PBS to remove the unincorporated label. F-actin distribution following CCL19 stimulation was evaluated by confocal laser scanning microscope (CLSM Leica SP2, Germany).

Immunohistochemical analysis. Seventy-eight specimens of SCCHN tumors with the adjacent metastatic (or normal) lymph nodes and 30 specimens of normal human oral mucosal tissue were obtained from the Head and Neck Tumor Center, School of Stomatology, China Medical University. All the specimens were obtained with the consent of the patients before surgery and in accordance with the Health Insurance Portability. The classification of SCCHN, including primary tumors (T), regional lymph nodes (N), distant metastasis (M) and stage grouping, was determined according to the rules of the International Union Against Cancer (UICC) for Head and Neck Cancer (tumor-node-metastasis (TNM) classification, 1997). Immunohistochemical staining used conventional horseradish peroxidase immunochemical staining methods. In brief, 5-µm sections of the specimens were deparaffinized and hydrated with 0.6% H₂O₂ in methanol to inhibit endogenous peroxidase, antigen retrieval was performed and then incubated with normal blocking serum for 10 min. Then, the sections were incubated with primary antibodies (1:100); CCR7-specific monoclonal antibody and MMP-9 overnight at 4°C. Immunodetection was performed using peroxidase labeled secondary antibody (R&D Systems) and diaminobenzidine for visualization. All sections were counterstained with hematoxylin (Sigma). Negative controls included omission of the primary antibody. The cell morphology was analyzed by microscopy (Nikon Eclipse 80i; Nikon, Tokyo,
Japan) at x100-400 magnification. According to the percentage of positive tumor cells, all cells were scored as negative (−, <10% or no staining); weak positive (+, 11-50%); positive (+++, 51-75%); or strongly positive (+++, >75%).

Results

CCL19 induces MMP-9 high expression and enhances its activity. MMP-2 and -9 are well-documented ECM-degrading enzymes whose activities are associated with SCCHN tumor invasiveness (8). To investigate whether MMP-2 and -9 play a role in the CCL19-stimulated cell invasion, MMP-2 and -9 protein and enzymatic activities were measured by western blotting and gelatin zymography. As shown in Fig. 1A and B, the expression and activity of MMP-2 were not significantly altered by CCL19. In contrast to MMP-9, both expression and activity of MMP-9 were found to be markedly elevated after CCL19 treatment (Fig. 1A and B). Thus, we continued researching...
MMP-9. We found that the expression and activity of MMP-9 were not only elevated after CCL19 treatment but were also diminished after CCR7 mAb treatment, which suggested that this was induced by CCR7 activation. At the same time, the role of CCR7 in MMP-9 activation was also blocked by SB-3CT indicating that CCR7 can activate MMP-9 (Fig. 1C and D). The expression and the activation of MMP-9 were determined by western blot analysis and gelatin zymography, quantified by computer-aided densitometry.

**CCL19 induces the migration of PCI-37B cells and SB-3CT blocks it.** We treated the cells in the absence or presence of SB-3CT, then conditioned medium of CCL19 was placed on the lower part of a Transwell unit and PCI-37B cells were added to the upper part in the absence or presence of SB-3CT. As shown in Fig. 2, CCL19 significantly enhanced the migration ability of PCI-37B cells that were specifically blocked by the SB-3CT. The inhibitive ability became stronger with the SB-3CT increasing density.

**CCL19 activates the chemotaxis ability and SB-3CT abolishes it in PCI-37B.** To investigate the correlation between the chemotaxis ability and the activities of CCR7 and MMP-9 in the metastatic SCCHN cell line, we separated the cell line into many teams and analyzed their chemotaxis ability in vitro in response to the respective chemokine ligand CCL19, SB-3CT and CCR7 mAb. These experiments showed that CCL19 enhanced chemotaxis of SCCHN significantly as compared with background control levels established with media alone. The SB-3CT and anti-CCR7 mAb significantly blocked CCL19-induced cell chemotaxis, as shown in Fig. 3.

**CCR7 induces F-actin rearrangement.** Cell motility involves regulation of the actin cytoskeleton and the actin-severing protein coflin regulates actin organization. We found that CCR7 activation leads F-actin polymerization and pseudopodia formation. In untreated cells, we observed a scattered distribution of F-actin (Fig. 5). In the cells treated with CCL19, F-actin arrays and pseudopodia were formatted, while these...
effects were blocked by SB-3CT. We therefore consider that the actin cytoskeletal rearrangement induced by CCL19 requires MMP-9.

**CCR7 and MMP-9 expressed by immunohistochemical staining have significant positive correlation in tumor tissues and metastatic lymph nodes.** Using immunohistochemistry, we investigated the location of CCR7 and MMP-9 in SCCHN tumor tissues, metastatic lymph nodes, normal lymph nodes and oral mucosal tissues. CCR7 and MMP-9 were found in the cell membrane and cytoplasm, mainly expressed in the surrounding of stroma in tumor cells and metastatic lymph node cells. Analyzing the sections of CCR7 and MMP-9 staining in normal lymph nodes and oral mucosal tissues, we observed that the number of stained cells was low and that they were not expressed (Fig. 5 and Table 1). The expression levels of CCR7 and MMP-9 were both significantly correlated with cervical lymph node metastasis and SCCHN clinical stage (P<0.05). In addition, the T3/T4 tumor size also appeared to express high levels of MMP-9 (P<0.05). However, there were no significant differences between CCR7 or MMP-9 expression and age or gender (P>0.05). A moderate correlation was observed between CCR7 and MMP-9 expression in SCCHN tumor tissues (P<0.05) and metastatic lymph nodes (P<0.05), but there was no significant correlation between normal lymph nodes (P>0.05) and normal oral mucosal tissues (P>0.05).

**Discussion**

CCR7 has been shown to interact with chemokines (CCL19, CCL21) and to modulate tumor cell migration, invasion and proliferation of metastatic squamous cell carcinoma of the head and neck (SCCHN) (9,10,24). However, the mechanisms of chemotaxis and migration and the signaling pathway involved remain poorly understood. We demonstrated that CCR7 regulates cell chemotaxis and migration via MMP-9 in metastatic SCCHN.

Remodeling of the extracellular matrix (ECM), which occurs during many physiological and pathological processes, is one of the requisite events of cellular invasion. MMPs can degrade almost all ECM proteins in the destruction of tumor cell invasion (12). MMP-9 is a well-documented ECM-degrading enzyme whose activities are associated with SCCHN tumor invasion (26). Regulating the activity of MMP-9 modulates...
the degradation of the ECM components which in turn alter cellular invasion, expression and activation.

It has been reported that CCR7 can regulate MMP-9 in lung cancer cells, thus affecting the expression of tumor (18). Chemokine CXCL12, through its specific receptor CXCR4, induced colon cancer metastasis of HT-29 cells by secretion of MMP-9 (22). The interaction of CCL21/CCR7 enhances the expression and secretion of MMP-9 in colon cancer, degradation of ECM and basement membrane, thus promoting invasion and metastasis of colon cancer (23). We speculated that CCR7-induced MMP-9 expression is an important regulatory factor. In the present study, our results (Fig. 1) showed that stimulation of CCL19 could also result in increased expression of MMP-9 in western blot analysis, and the activity in gelatin zymography. Furthermore, we used Transwell chemotaxis and migration assays to examine CCL19-induced activation of MMP-9 that significantly enhanced the chemotaxis (Fig. 3) and migration (Fig. 2) index in SCCHN, and was blocked by MMP-9 inhibitor SB-3CT, thus supporting the hypothesis that CCR7-induced MMP-9 expression is an important regulatory factor.

In the present study, our results (Fig. 1) showed that stimulation of CCL19 could also result in increased expression of MMP-9 in western blot analysis, and the activity in gelatin zymography. Furthermore, we used Transwell chemotaxis and migration assays to examine CCL19-induced activation of MMP-9 that significantly enhanced the chemotaxis (Fig. 3) and migration (Fig. 2) index in SCCHN, and was blocked by MMP-9 inhibitor SB-3CT, thus supporting the hypothesis that CCL19 concentrations in the lymph nodes probably induce SCCHN cell migration into these organs through a CCR7-mediated mechanism. High levels of actin polymerization are required for the formation of pseudopodia, which are needed for chemokine mediated cell migration and invasion into surrounding tissues and efficient metastasis formation (27). In the present study (Fig. 4), we examined TRITC-labeled phalloidin staining by inverted microscope, and observed reorganization of the actin cytoskeleton of PCI-37B was enhanced in response to CCL19, and this function was inhibited by SB-3CT. Immunohistochemical studies confirmed the presence of CCR7 and MMP-9 in the cytoplasm and cell membrane of SCCHN tumor tissues, metastatic lymph nodes, and all significantly correlated with cervical lymph node metastasis and clinical stage, but in normal lymph nodes and oral mucosal tissues they were low or absent.

CCR7 has been reported to be a novel prediction biomarker of metastasis in cancer. Our results showed that stimulation of CCL19 could also result in increased chemotaxis and migration of SCCHN cells. CCR7 induced the activation of MMP-9, and MMP-9 interacted with its counterpart molecules. As has previously been reported, we found that when MMP-9 was inhibited, the CCL19-induced chemotaxis and migration of SCCHN cells were also inhibited.

Future studies including the immunohistochemical analysis of both CCR7 and MMP-9 may be useful for predicting lymph node metastasis (23). Therefore, we suggest that CCR7 activation by CCL19 via MMP-9 may promote SCCHN cell chemotaxis, migration.

In summary, our findings emphasize the potential role of overexpression of the CCR7 in promoting cellular migration and matrix-degrading activities through MMP-9 in SCCHN cells. This information provides a mechanistic rationale for the observed MMP-9 overexpression in advanced-stage SCCHN. To our knowledge, the chemotaxis and migration of SCCHN are very complex systems. It is therefore impossible to cure the tumor by blocking only this pathway. However, it provides us with a novel idea for enhancing the invasiveness of SCCHN and may contribute to the development of improved and more specific therapeutics for the treatment of SCCHN.

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Table I. Correlations between MMP-9 expression and clinicopathological factors of SCCHN.

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*P<0.05, the internal difference of CCR7 or MMP-9 expression within clinicopathological characteristics.
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