Abstract. The receptor activator of nuclear factor κB ligand/receptor activator of nuclear factor κB (RANKL/RANK) pathway is crucial for the migration of RANK-expressing cancer cells. The ubiquitin-proteasome protein degradation pathway plays a significant role in tumor metastasis. However, the relationship between these two pathways in tumor cell migration is unclear. In the present study, we explored the effect of the proteasome inhibitor bortezomib (PS-341) on RANKL-induced MDA-MB-231 breast cancer cell migration. Transwell migration assay showed that RANKL-induced MDA-MB-231 cell migration was significantly blocked by the decoy receptor osteoprotegerin (OPG), and was also inhibited by the PI3-K inhibitor LY294002. Western blotting results showed that Akt was rapidly activated by soluble RANKL treatment. PS-341 significantly enhanced RANKL-induced MDA-MB-231 cell migration. Further study showed that the enhancement of migration by PS-341 involved upregulation of activated Akt and RANK. Our results for the first time support the theory that PS-341 treatment may be unsuitable for RANK-positive breast cancer patients.

Introduction

Breast cancer is the most common malignancy in women. Although early-stage breast cancers are not life threatening, development of metastatic breast cancer is responsible for the majority of cancer-related deaths and the poor quality of life of patients. Currently available treatments are unable to eradicate metastatic cancer, and the survival rate for these patients is low. Further therapeutic advances require new strategies based on a fundamental understanding of breast cancer biology.

Recent studies have shown that the receptor activator for nuclear factor κB ligand/receptor activator for nuclear factor κB (RANKL/RANK) pathway is critical for RANK-expressing cancer cells to home to bones, and RANKL increased migration of RANK-expressing cancer cells, including breast cancer cells (1-3). However, the regulatory mechanism of the RANKL/RANK pathway in cancer cell migration remains largely unknown.

The ubiquitin-proteasome protein degradation pathway plays an essential role in the orderly proteolysis of intracellular proteins. In cancer cells, this pathway affects numerous activities that are important for tumor growth and metastasis (4). The proteasome inhibitor bortezomib (PS-341) has been licensed for the treatment of refractory multiple myeloma and mantle cell lymphoma. Although PS-341 is active against breast cancer cell lines (5,6), the effect of PS-341 on RANKL-induced breast cancer cell migration has yet to be clarified.

In the present study, we explored the effect of PS-341 on the RANKL/RANK pathway and demonstrated for the first time that PS-341 enhanced RANKL-mediated MDA-MB-231 cell migration. This increased migration was associated with upregulation of RANK expression and phospho-Akt levels.

Materials and methods

Cell culture. The MDA-MB-231 breast cancer cell line was obtained from the Type Culture Collection of the Chinese Academy of Sciences (Shanghai, China). MDA-MB-231 cells were grown in Leibovitz L-15 medium (Gibco, Carlsbad, CA, USA) containing 10% FBS, according to the recommendations of the supplier.

Reagents and antibodies. Recombinant soluble RANKL (sRANKL) and recombinant human osteoprotegerin (OPG; rOPG) were purchased from Cytolab/Peprotech Asia (USA) and PS-341 was purchased from Millennium Pharmaceuticals, Inc. (Cambridge, MA, USA). The specific PI3-K inhibitor LY294002 was obtained from Sigma (St. Louis, MO, USA). Rabbit anti-actin antibodies were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Rabbit anti-Akt and rabbit anti-p-Akt (Ser473) were obtained from Cell Signaling Technology (Danvers, MA, USA).
RANK expression analysis. The MDA-MB-231 cells were seeded at 5x10^5 cells/well in 6-well plates. Surface RANK expression was determined by flow cytometry following incubation with either 25 µg/10^6 cells mouse anti-RANK antibody (R&D Systems, Minneapolis, MN, USA) or isotype control (R&D) followed by FITC-conjugated anti-mouse secondary antibody. Fluorescence was assessed using a FAScan sorter (BD Biosciences, San Jose, CA, USA). Finally, the data were analyzed with WinMDI software.

Western blotting. To prepare total cell lysates, cell pellets were washed twice with ice-cold PBS and solubilized in 1% Triton lysis buffer [1% Triton X-100, 50 mM Tris-HCl (pH 7.4), 150 mM NaCl, 10 mM EDTA, 100 mM NaF, 1 mM Na3VO4, 1 mM phenylmethylsulfonyl fluoride and 2 µg/ml apro tinin] on ice. Cell lysates were centrifuged at 12,000 x g for 30 min at 4°C and the supernatants were separated. Proteins were eluted by heat treatment at 100°C for 5 min with 3X sampling buffer. Total proteins (30-50 µg) were subjected to SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and electronically transferred to nitrocellulose membranes. After blocking with 5% skim milk in TBST [10 mM Tris (pH 7.4), 150 mM NaCl and 0.1% Tween-20], the blots were probed with the indicated primary antibodies at 4°C overnight, followed by the horseradish peroxidase-conjugated specific goat anti-mouse or goat anti-rabbit secondary antibody as indicated for 30 min at room temperature. Finally, proteins were detected using an enhanced chemiluminescence reagent (SuperSignal Western Pico Chemiluminescent Substrate; Pierce, USA) and visualized with Electrophoresis Gel Imaging Analysis System (DNR Bio-Imaging Systems, Jerusalem, Israel).

Transwell migration assay. Cells were washed twice in culture medium with 0.1% bovine serum albumin. A 24-well chemotaxis chamber (8-µm pore size; Corning, NY, USA) was used for this experiment. Prior to performing the migration assay, cells were pre-treated with different concentrations of inhibitors or the appropriate solvent control (dimethyl sulfoxide; DMSO) for 60 min. A volume of 200 µl (1x10^5 cells/ml) from each sample was loaded onto the upper well. The medium (0.5 ml) containing sRANKL with or without the OPG or DMSO was added to the bottom well. The plates were incubated for 16 h at 37°C. Following incubation, the porous inserts were carefully removed and the cells on the lower surface of the membrane were stained and counted in at least five different fields. The results were expressed as the percentage of migrated cells as compared to the control (untreated cells). Each experiment was performed at least three times.

Statistical analysis. Experimental data are presented as the means ± standard deviation. The significance of the difference between the groups was assessed by the Student’s two-tailed t-test, and P<0.05 was considered significant in all statistical analyses. All mean values were calculated from at least three independent experiments. Statistical analyses were carried out using SPSS 16.0 for Windows (SPSS, Inc., Chicago, IL, USA).

Results
RANKL and RANK interaction direct the migration of MDA-MB-231 breast cancer cells. Firstly, flow cytometry was used to examine RANK expression in MDA-MB-231 cells. The result showed that MDA-MB-231 cells expressed RANK on the cell surface (Fig. 1A). Subsequently, stimulation of the MDA-MB-231 cells by sRANKL significantly increased cell migration. Moreover, treatment of cells with decoy receptor of RANKL, OPG, significantly blocked RANKL-induced cell migration. Akt is involved in RANKL-induced MDA-MB-231 cell migration. Next, the downstream signaling of RANKL/RANK in MDA-MB-231 cells was examined. Since the PI3-K/Akt pathway is closely correlated with cell invasion and migration (2,7-10), the effect of Akt on the RANKL/RANK pathway was explored. The results showed that Akt was transiently activated by sRANKL in MDA-MB-231 cells (Fig. 2A). The sRANKL-induced migration was significantly inhibited by the PI3-K inhibitor LY294002 (50 µM) (Fig. 2B).
These results indicated that the PI3-K/Akt pathway was involved in RANKL-induced breast cancer cell migration.

PS-341 enhances RANKL-induced MDA-MB-231 cell migration by upregulation of RANK and phospho-Akt. Transwell migration assay was used to explore the effect of PS-341 on RANKL-induced MDA-MB-231 cell migration. The results showed that pre-treatment with PS-341 for 4 h markedly increased RANKL-induced MDA-MB-231 cell migration (Fig. 3A). Western blotting results demonstrated that pre-treatment with PS-341 for 4 h, followed by exposure to sRANKL,
significantly increased the p-Akt levels (Fig. 3B). FACS results showed that after treatment with PS-341 for 24 h, RANK expression on MDA-MB-231 cells was upregulated (Fig. 3C and D).

These results indicated that PS-341 enhanced RANKL-induced MDA-MB-231 cell migration by upregulation of RANK and phospho-Akt levels.

Discussion

The RANKL/RANK pathway is considered the key pathway in the regulation of osteoclast and lymphocyte proliferation and survival. RANKL interacts with RANK and in turn recruits tumor necrosis factor receptor-associated factors (11) leading to activation of NF-κB, c-Jun N-terminal kinase, p38, ERK and Akt (12-14). As recently demonstrated, RANK has been found to be expressed in several solid tumors, such as breast cancer, prostate cancer, malignant bone tumors, multiple myeloma and squamous cell carcinoma (1,15), and RANKL directs RANK-expressing cancer cells to preferentially migrate into bone. However, the exact signaling pathways in tumor cell migration need to be elucidated. Tang et al (16) reported that Src and MAPK were involved in RANKL-induced MDA-MB-231 cell migration. In T47D breast cancer cells, ERK and Akt were involved in RANKL-induced cell migration (1), while in C42b prostate cancer cells, ERK, but not Akt, was involved in the RANKL/RANK pathway (3). These data showed that RANKL/RANK signaling pathways vary in different tumor cell types. In the present study, we firstly showed that Akt was involved in RANKL-induced MDA-MB-231 cell migration.

Proteasome is involved in the targeted elimination of damaged and regulatory proteins, such as transcription factors, signaling molecules and cell cycle inhibitors (17). The inhibition of the proteasome results in the abnormal accumulation of many intracellular proteins, thereby disrupting cellular homeostasis. In osteoclast differentiation and survival, the RANKL/RANK pathway was found to be regulated by the proteasome pathway (18,19). Tumor cells have been reported to be more sensitive to proteasome inhibition than normal cells, thus making proteasome a potential novel target for cancer treatment (20).

PS-341 is a selective and reversible proteasome inhibitor that results in stabilization of cell cycle regulatory proteins, inhibition of NF-κB activation, induction of apoptosis and override of Bcl-2 resistance and anti-angiogenesis (21). Moreover, PS-341 exhibits a wide range of antitumor activity and increases the activity of multiple chemotherapeutic agents (5,6,22,23). Although PS-341 was found to be active against breast cancer cell lines, in vivo models and phase I clinical trials have found it ineffective for breast cancer when used as a single agent (24), and limited efficacy of PS-341 has been noted in combination with several chemotherapeutic agents (25,26).

In the present study, the role of low-dose PS-341 (10 nM) in RANKL-induced MDA-MB-231 cell migration was explored. The results showed that PS-341 significantly enhanced RANKL-induced MDA-MB-231 cell migration. Furthermore, PS-341 treatment resulted in the upregulation of RANK expression and p-Akt levels, which suggests that the activation of the RANKL/RANK/Akt pathway may contribute to increased RANKL-induced breast cancer cell migration. Since a recent study showed that RANK expression may predict accelerated skeletal metastasis-free survival and poor prognosis in breast cancer patients (27), our results suggest that PS-341 treatment may be unsuitable for RANK-positive breast cancer patients.

Taken together, our results provide evidence that PS-341 increases the RANKL-induced MDA-MB-231 cell migration. Further study is required to confirm the effect of PS-341 in RANKL-expressing cancer cells.

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References


