

EphA2 silencing in nasopharyngeal carcinoma leads to decreased proliferation, invasion and increased sensitization to paclitaxel

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Abstract. EphA2 is frequently overexpressed and functionally altered in a variety of human cancers. However, its roles in human nasopharyngeal carcinoma (NPC) remain unclear. To investigate the roles of EphA2 in the development and progression of NPC, we initially evaluated the expression pattern of EphA2 protein in NPC tissues using western blotting and CCK-8 assay. Fluorescence-activated cell sorting analysis and invasion assay were conducted to observe the effects of EphA2 inhibition *in vivo*. Our results demonstrated that EphA2 was overexpressed in NPC specimens and the expression of EphA2 was significantly associated with T classification, advanced clinical stage and lymph node metastasis. Moreover, human NPC 5-8F cells were infected with lentiviral vector-mediated EphA2-specific shRNA, which resulted in the significant inhibition of cell growth, invasion of 5-8F cells and markedly enhanced the sensitivity of 5-8F cells to the chemotherapeutic agent paclitaxel *in vitro*. Collectively, our results demonstrate that EphA2 is involved in malignant cell behavior and is a potential therapeutic target in human NPC.

Introduction

Nasopharyngeal carcinoma (NPC), an Epstein-Barr virus (EBV)-associated cancer, is highly prevalent in southeast Asia, especially in the Cantonese region around Guangzhou in China (1). NPC is distinct from other cancers of the head and neck in terms of its epidemiology, histopathology, clinical characteristics, methods of treatment and patterns of failure (2). Although patients benefit from multi-modal treatment,

including radiotherapy, chemotherapy and biological therapy, limited improvement in 5-year survival has been achieved over the last few decades (3). Cancer metastasis and therapy failure account for the low survival rates in patients with NPC (4,5). However, the molecular mechanism of the development and progression of NPC remains poorly understood. It is of great clinical value to understand the molecular mechanism of this cancer and identify valuable predictive markers as well as novel therapeutic strategies.

The Eph protein family constitutes the largest group of trans

Table I. Correlation between the clinicopathological characteristics and protein expression of EphA2.

Characteristics	No. of samples	Relative protein expression of EphA2	t-value	P-value
Age (y)				
<46	24	0.86±0.37	0.959	0.343
≥46	23	0.76±0.30		
Gender				
Male	37	0.79±0.33	0.571	0.571
Female	10	0.86±0.36		
Smoking history				
Smoker	29	0.86±0.35	1.423	0.162
Nonsmoker	18	0.72±0.29		
T classification				
T1 + T2	20	0.68±0.28	2.272	0.028
T3 + T4	27	0.90±0.34		
Clinical stage				
I + II	19	0.64±0.21	3.069	0.004
III + IV	28	0.92±0.36		
Metastasis				
Yes	27	0.91±0.34	2.616	0.012
No	20	0.67±0.28		
EB-virus infection				
Yes	26	0.87±0.34	1.338	0.188
No	21	0.74±0.32		

pathological parameters in patients with NPC. A lentiviral RNAi system was employed to knock down the expression of EphA2 in NPC 5-8F cells. The roles of EphA2 in NPC cell invasion and chemotherapy *in vitro* conditions were investigated.

Materials and methods

Patients and tissue preparation. A total of 47 fresh undifferentiated NPC (WHO type III) tissues and 21 cases of non-carcinoma epithelial tissues (NCET) from the nasopharynx were obtained from the Department of Otolaryngology of Xiangya Hospital, Central South University, China, between October 2010 and March 2011. None of the patients had any history of previous malignancies, or history of radiotherapy or chemotherapy. Metastases were diagnosed by clinical examination and imaging evaluation. The clinical stage of all the patients was classified according to the 2008 NPC staging system of China. Specimens were snap-frozen immediately and stored in liquid nitrogen for total

manufacturer's instructions. In brief, cells at a concentration of 3.0×10^3 /well were cultured in triplicate in 96-well plates with 10% FBS at 37°C in humidified 5% CO₂ atmosphere for various periods and exposed to fresh media every other day. At 24, 48, 72 and 96 h post-transfection, cell proliferation was measured by absorbance at an optical density of 490 nm.

Cell apoptosis and cell cycle analysis by flow cytometry. Fluorescence-activated cell sorting analysis was carried out. Following incubation in serum-free medium for 24 h, the cells were washed with PBS at 4°C twice and their concentration was altered to 1×10^6 /ml. Annexin V/Cy5 and propidium iodide were added for incubation for 30 min. Fluorescence-activated cell sorting analysis was carried out using the FACS Calibur instrument (Becton Dickinson, Franklin Lakes, NJ, USA).

Matrigel invasion assay. The invasiveness of the transfected cells was evaluated in 24-well transfected chambers (Costar, Cambridge, MA, USA) according to the manufacturer's instructions. Briefly, transwell with an 8 µm diameter pore membrane was coated with 200 µl matrigel at 200 µg/ml and incubated overnight. Cells (3×10^4) in 100 µl of serum-free medium were seeded into the upper chamber of the transwell and the lower chamber was filled with 0.8ml RPMI-1640 containing 12% FBS to induce chemotaxis. Following 48 h of incubation at 37°C in a humidified 5% CO₂ atmosphere, the cells were fixed in methanol and stained with crystal violet, and the cells that invaded through the pores to the lower surface of the filter were counted under a microscope. Three invasion chambers were used per condition. The values obtained were calculated by averaging the total number of cells from three filters.

Cell viability assay. Cells were seeded in 96-well plates. Following overnight culture, they were exposed to various concentrations of paclitaxel (0.001, 0.01, 0.1, 1.5, 10, 20 and 30 nM) for 48 h in a CO₂ incubator. CCK8 assay was used to detect the chemo-sensitivity of cells. Absorbance values were expressed as percentages relative to the controls, and the concentrations resulting in 50% inhibition of cell growth (IC₅₀ values) were calculated.

Statistical analysis. Statistical analysis was performed with SPSS 17.0 software. Results of quantitative data in this study were shown as the mean ± SD. Statistical differences between groups were compared using the two-tailed Student's t-test. $P < 0.05$ was considered statistically significant.

Results

Overexpression of EphA2 protein in NPC and its correlation with clinicopathological parameters. The expression pattern of EphA2 protein was initially assayed in 47 fresh NPC specimens and 21 non-carcinoma epithelial tissue (NCET) specimens. Western blot analysis clearly revealed that the expression of EphA2 protein in NPC tissues was significantly higher than that in NCET (0.81 ± 0.33 vs. 0.43 ± 0.21 ; $t = 4.841$, $P < 0.001$) (Fig. 1). Furthermore, the association between the expression of EphA2 protein and clinicopathological characteristics of NPC was explored using the Student's t-test. EphA2 overexpression was significantly associated with NPC

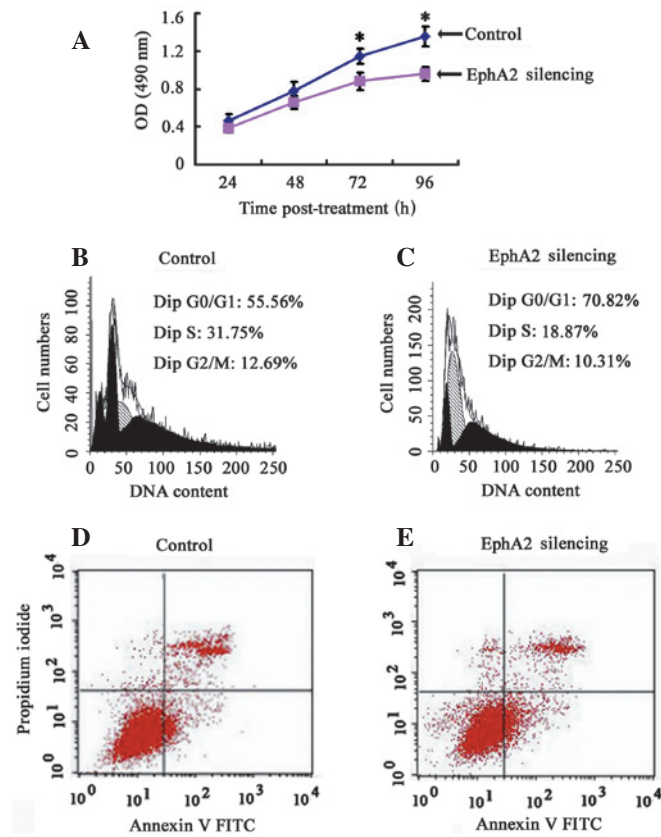


Figure 3. The effect of EphA2 on the proliferation, cell cycle distribution and apoptosis of NPC 5-8F cells. (A) Cell proliferation was measured by the CCK-8 assay every 24 h for 4 days. (B and C) Differently treated cells were stained with propidium iodide and analyzed by flow cytometry. Proportion of cells in various phases of the cell cycle. The results are the means of three independent experiments $\pm$ SD ($P < 0.05$). (D and E) Cells staining positive for Annexin V-FITC and negative for PI were considered to have undergone apoptosis. Average apoptotic rate of three independent experiments $\pm$ SD are shown. * $P < 0.05$.

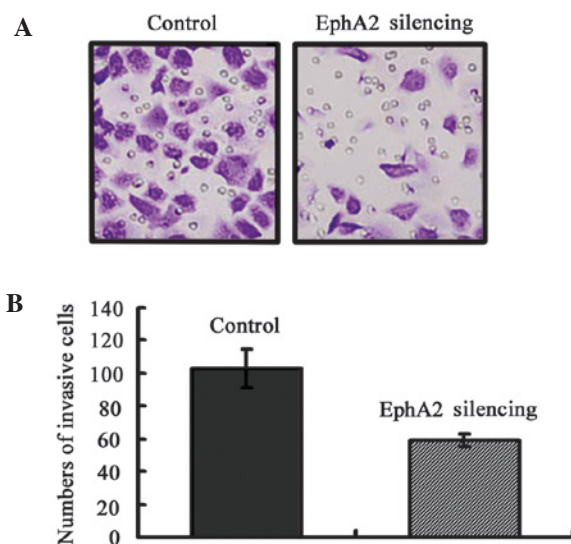


Figure 4. Knockdown of EphA2 inhibited the invasion of NPC 5-8F cells *in vitro*. (A) Transwell invasion assay showed that the knockdown of EphA2 inhibited cell invasion of 5-8F. Images were captured at a magnification of $\times 200$. Representative of three independent experiments. (B) Graphical representation of the number of invasive 5-8F cells per microscopic field. Data were shown as the mean $\pm$ SD from three independent experiments.

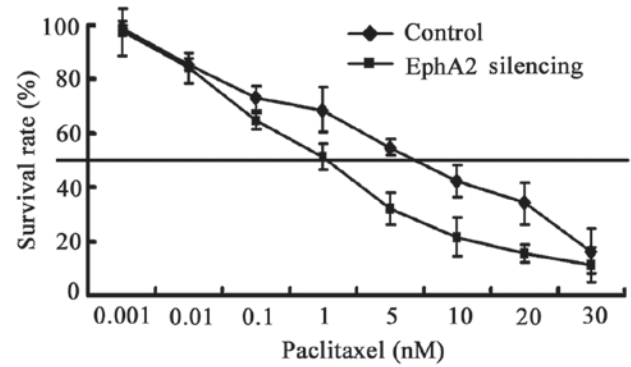


Figure 5. Knockdown the expression of EphA2 sensitized 5-8F cells to paclitaxel. The

used chemotherapeutic agent in NPC. 5-8F cells were exposed to paclitaxel following transfection with EphA2-shRNA for 48 h. Cell viability was evaluated using CCK-8 assays. The IC₅₀ value of cells transfected with EphA2-shRNA and control-shRNA were 1.6 and 5.6 nM, respectively (Fig. 5). The results show that the sensitivity of 5-8F to paclitaxel is enhanced by inhibiting the expression of EphA2 3.5-fold.

Discussion

Our previous investigations have clearly demonstrated EphA2 overexpression in human laryngeal and hypopharyngeal squamous cell carcinoma, and EphA2 regulation of the growth and cervical lymph node metastasis of human laryngeal and hypopharyngeal squamous cell carcinoma *in vitro* and *in vivo* (20,22). Although NPC is also classified as a subtype of head and neck squamous cell carcinoma, it possesses unique and distinct epidemiology, clinical characteristics, etiology and histopathology (2). Therefore, the association between EphA2 expression and clinicopathological parameters in patients with NPC, and its role in growth, invasion and paclitaxel chemotherapy of NPC were investigated for the first time in the present study.

Results of the present study initially revealed that the expression level of EphA2 protein was significantly higher in NPC samples than that in non-carcinoma tissues obtained from the nasopharynx. Moreover, the results of 47 NPC specimens assayed by western blotting further demonstrated that a high protein expression of EphA2 was significantly associated with T classification, clinical stage and cervical lymph nodes metastasis, which was consistent with recent publications on NPC (23) and other reports in urinary bladder cancer (24), ovarian cancer (13) and lung cancer (25,26), indicating the importance of EphA2 in the malignant progression of patients with NPC.

Metastasis at the early stages is one of the most frequent clinical features in patients with NPC. Approximately 70-80% of new patients with NPC present with cervical lymph node metastasis, and approximately 4.2% of those patients have distant metastasis to the bone, lung, liver, and central nervous system (27). Metastasis severely reduces the possibility of successful treatment and overall survival time (28). Therefore, identification of molecular markers for metastasis would be beneficial in designing optimized and individualized therapeutic regimens for patients with NPC. The present study revealed the association of EphA2 with lymph node metastasis in NPC, suggesting that EphA2 could be considered as an indicator for the propensity to metastasize. Furthermore, EphA2 inhibition was able to reduce the invasive ability of NPC 5-8F cells *in vitro*, supporting the theory that EphA2 is involved in the process of invasion and metastasis in patients with NPC.

Since acquired resistance to chemotherapy negatively affects the outcome of patients with NPC (29) and the finding that EphA2 knockdown led to inhibition of NPC proliferation *in vitro*, the potential therapeutic function of EphA2 in combination with a chemotherapeutic drug was further explored in the current study. Paclitaxel is a tubulin-stabilizing anti-cancer agent and now widely used in the therapy of patients with NPC. We found that knockdown of the EphA2 protein significantly enhanced the cytotoxicity of paclitaxel in human NPC

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