Overexpression of BPIFB1 promotes apoptosis and inhibits proliferation via the MEK/ERK signal pathway in nasopharyngeal carcinoma

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Abstract: Nasopharyngeal carcinoma (NPC) is a respiratory malignant epithelial carcinoma. Research has indicated that bactericidal/permeability-increasing fold-containing protein B1 (BPIFB1), mostly secreted by nasopharyngeal epithelia, is dysregulated in patients with NPC. This study aimed to explore the effects of BPIFB1 inviability, proliferation, apoptosis and its molecular mechanism. To confirm the effects of BPIFB1 on NPC cells, BPIFB1 was overexpressed or silenced in NPC-KT cells after being transfected with BPIFB1 or siBPIFB1 plasmids. The results showed that BPIFB1 overexpression could induce apoptosis and DNA damage in NPC-KT cells, and silenced BPIFB1 had the opposite effects. BPIFB1 overexpression can inhibit the cell cycle by being arrested at the G0/G1 phase and by regulating the MEK/ERK signaling pathway. MEK inhibitor U0126 was used to confirm the effects of BPIFB1 on the MEK/ERK pathway, and U0126 can inverse the effects of siBPIFB1. Additionally, BPIFB1 can enhance the anti-proliferative effect of chemotherapy drugs on NPC-KT cells. All the results indicated that BPIFB1 could be a potential target for the treatment of NPC.

Keywords: Nasopharyngeal carcinoma, BPIFB1, MEK/ERK pathway, cell cycle, apoptosis, proliferation

Introduction

Nasopharyngeal carcinoma (NPC), a malignant epithelial carcinoma, occurs in the head and neck region with a remarkable racial and geographical incidence [1]. The disease is rare globally but occurs more commonly in Southeast Asia and Southern China and with a gender imbalance (the male-to-female ratio is about 3:1), which indicates that genetic, ethnic, and environmental factors play an important role in the disease [2, 3]. The primary treatment of NPC is radiotherapy (RT), as it is relatively sensitive to ionizing radiation for patients in the disease’s early stages. For late stage patients, chemoradiotherapy (CCRT) is superior to RT alone and leads to a better prognosis [4, 5]. Although radiotherapy or chemoradiotherapy can control local NPC, a large number of patients in stages III and IV still experience local recurrence or distant metastasis [6]. Therefore, more studies on the mechanisms of NPC progression are essential to develop new targets and novel therapies for treating NPC.

Bactericidal/permeability-increasing (BPI)-fold-containing family B member 1 (BPIFB1), as the so called long-palate lung and nasal epithelium clone 1 (LPLUNC1), is secreted by nasopharyngeal epithelia or minor mucosal glands of the respiratory and upper digestive tracts [7, 8]. Many studies have confirmed that BPIFB1 is dysregulated in patients with respiratory tract disease. Researchers showed that BPIFB1 is downregulated in tumor tissues, negatively correlates with the severity of NPC [9], and, more importantly, relates with prognosis in NPC patients [10]. The previous studies indicated that BPIFB1 plays an important role in NPC progression [11] and could be a potential target for NPC treatment, but the details of its mechanisms remain unclear [12]. This research aimed to investigate the effects of BPIFB1 on NPC cells and its related mechanisms.
Materials and methods

Cell culture

NPC-KT is an EBV-positive epithelial cell line derived from EBV-negative nasopharyngeal epithelial Ad-AH cells and EBV-positive NPC tissue, and our supply was kindly provided by Wuhan University. The NPC-KT cells were cultured in Dulbecco’s Modified Eagles Medium (DMEM) containing 10% fetal bovine serum (FBS), supplemented with penicillin and streptomycin, at 37°C with 5% CO₂ [13].

Plasmid transfection

The BPIFB1 overexpressed mimics, BPIFB1 silencing siRNA (si-BPIFB1) and the separated negative control were purchased from Shanghai GenePharma Co., Ltd. (Shanghai, China) and transfected into NPC-KT cells using the Lipo2000 transfection reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s instructions. The sequences were cited as described [14].

Assay of cell proliferation

Cell proliferation was determined by MTT assay. The cells were seeded in a 96-well culture plate. Then 10 μL of MTT solution was added to a 100 μL medium and incubated at 37°C for 4 h. Then we added 200 μL DMSO to each well. Next, we incubated the plate at 37°C for 10 min and read the absorbance at 490 nm using an ELISA plate reader (Beckman, Miami, FL).

Cell cycle analysis

The cell cycle analysis was measured by flow cytometry. Briefly, the cells were fixed in cold, 70% alcohol at 4°C overnight, treated with 0.25% Triton X-100 for 5 min, stained in a propidium iodide solution at a final concentration of 50 μg/mL (Sigma-Aldrich, St. Louis, Mo. USA) for 1 h at 37°C. The cell cycle analysis was performed by a FACS can flow cytometer (BD Biosciences, Bedford, MA).

Hoechst stains

The blue fluorescent Hoechst dye (Sigma-Aldrich Chemical Co., St Louis, USA) creates cell permeable nucleic acid stains and is sensitive to DNA conformation and the chromatin state and thus can be used to detect nuclear damage. NPC-KT cells were washed or resuspended in a PBS solution (pH 7.4) and then stained in the Hoechst staining solution (1 μg/mL) for 20 min. Then the cells were examined under an inverted fluorescence microscope.

RNA isolation and qRT-PCR

Total RNA was isolated from NPC-KT cells using the TRIzol reagent (Invitrogen, Breda, Netherlands) and transcribed into cDNA using a First Strand cDNA Synthesis kit (Promega, San Luis Obispo, CA, USA) according to the manufacturer’s instructions. Real-time quantitative PCR (RT-qPCR) was performed using specific primers and a SYBR green PCR kit (GenePharma, Shanghai, China) and GAPDH was used as an internal control. The sequences of the primers are described elsewhere [15].

Protein extraction and Western blotting

The pretreated NPC-KT cells were lysed in a cold lysis buffer (RIPA solution with phosphatase and protease inhibitor cocktail (Sigma-Aldrich, St. Louis, Mo. USA) and centrifuged at 12,000 g for 10 min. Protein concentrations were measured using the Bradford method (Pierce® BCA Protein Assay Kit, Thermo Scientific, Rockford, USA) and degenerated at 95°C. Equal amounts of proteins were separated on Invitrogen™NuPAGE™Bis-Tris gels (10%) and transferred onto PVDF membranes (Millipore, MA). After blocking, primary and secondary antibody incubation (primary antibody: BIPFB1, caspase-3,Bcl-2, Bax, CyclinD1, Cyclin B1, MEK, p-MEK, ERK, p-ERK, GADPH (Cell Signaling Technology, Inc. Beverly, MA) at 1:1,000 dilution; secondary antibody at 1:10000 dilution), the proteins were visualized using an ECL system according to the manufacturer’s instructions.

Statistical analysis

Data were analyzed using one-way ANOVA with Tukey’s test using Prism 6 (GraphPad Software, Inc). Data are presented as the mean ± standard deviation (SD) and P<0.05 was considered statistically significant.

Results

BPIFB1 promoted the apoptosis of NPC-KT cells

To investigate the effects of BPIFB1 on the apoptosis of NPC-KT cells, BPIFB1 was overex-
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Figure 1. The effects of BPIFB1 on the cell apoptosis of NPC-KT cells. NPC-KT cells were transfected with BPIFB1 mimics or siRNA and the negative control (A, B) BPIFB1 expression of mimics or siRNA transfected NPC-KT cells was
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measured by PCR (A) and Western blotting (B). (C) DNA damage was measured by Hoechst staining. (D) The cell apoptosis was analyzed by flow cytometry. (E) The apoptosis related protein expression was assessed by Western blotting. Data were presented as the mean ± SD. n=3, *P<0.05, **P<0.01 vs. Control.

Figure 2. The effect of BPIFB1 on the cell cycle of NPC-KT cells. NPC-KT cells were transfected with BPIFB1 mimics or siRNA and the negative control. A. The proliferation was assessed using an MTT assay. B. The cell cycle distribution was analyzed by flow cytometry. C. The protein levels of the cyclins were analyzed by Western blotting. Data are the means ± SD, n=3, *P<0.05, **P<0.01 vs. Control.

pressed or silenced by the BPIFB1 plasmid or siRNA. As shown in Figure 1, nuclear damage, cell apoptosis, and the apoptosis protein level (Bax, Bcl, Caspase 3) were examined. The transfection efficiency was confirmed by Western blotting and RT-PCR analysis. The BPIFB1 mRNA (Figure 1A) and the protein level (Figure 1B) in the BPIFB1 group were remarkably increased and in the siBPIFB1 group they were decreased compared with the negative
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control. Hoechst staining results (Figure 1C) indicated that the cell proportion with DNA damage (arrows in Figure 1C) was increased when BPIFB1 was overexpressed and decreased when BPIFB1 was silenced. Furthermore, the apoptosis results (Figure 1D) measured by flow cytometry were in accordance with the Hoechst staining results. The amount of apoptotic cells was increased when BPIFB1 was overexpressed and decreased when BPIFB1 was silenced. Apoptotic protein expressions were measured by Western blotting. As shown in Figure 1E, BPIFB1 overexpression increased the expression of the pro-apoptosis protein Bax and increased caspase 3, anti-apoptosis Bcl-2, and in siBPIFB1 the opposite was the case. All the results demonstrated that BPIFB1 promoted the apoptosis of NPC-KT cells.

**BPIFB1 inhibited the proliferation of NPC-KT cells**

To evaluate the effects of BPIFB1 on the proliferation of NPC-KT cells, the cell viability, cell cycle, and cyclin were investigated. As shown in Figure 2A, the MTT results demonstrated that BPIFB1 overexpression (BPIFB1 group) could inhibit the proliferation of NPC-KT cells, and BPIFB1 silencing (siBPIFB1) promoted proliferation. To explore the mechanism by which BPIFB1 inhibited cell proliferation, we evaluated the cell cycle by flow cytometry. The amount of the cells in the G0/G1, S, and G2/M phases is illustrated in Figure 2B. The fraction in the G0/G1 phase was higher in the BPIFB1 overexpression group, and the fraction in the S phase was lower. The result of the BPIFB1 group was in the opposite. Furthermore, cell cycle proteins (cyclin D1 and cyclin B1) were measured by Western blotting. As shown in Figure 2C, the level of cyclins was decreased when BPIFB1 was overexpressed and increased when BPIFB1 was silenced. All the results demonstrated that BPIFB1 inhibits the cell cycle of NPC-KT cells.

**BPIFB1 inhibited the activation of the MEK/ERK signaling pathway**

The MEK/ERK signaling pathway plays an important role in NPC cell viability, proliferation, and chemoresistance. To investigate the effects of BPIFB1 on the MEK/ERK pathway, the expression of phosphorylated MEK (p-MEK), ERK (p-ERK), and total MEK, ERK were analyzed by Western blotting. To demonstrate the activation of ERK and MEK, p-ERK/ERK and p-MEK/MEK were used. As shown in Figure 3, p-MEK/MEK and p-ERK/ERK were decreased in the BPIFB1 overexpressed group, and increased in BPIFB1 silenced group, which indicated that BPIFB1 could inhibit the activation of the MEK/ERK signaling pathway.

**BPIFB1 reduced NPC-KT cell viability via the MEK/ERK signaling pathway**

To confirm that BPIFB1 inhibited cell viability through the MEK/ERK signaling pathway, siBPIFB1 was transfected into NPC-KT cells, and the MEK inhibitor U0126 was added to the medium in another group. As in the previous results, in the siBPIFB1 cells, NPC-KT cell viability increased (Figure 4A), DNA damage decreased (Figure 4B), the proportion of apop-
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Figure 4. BPIFB1 regulated cell viability of NPC-KT cells via the MEK/ERK signaling pathway. NPC-KT cells were transfected with siRNA and/or treated with the MEK inhibitor U0126. A. The proliferation was assessed by MTT assay. B. DNA damage was measured by Hoechst staining. C. The cell apoptosis was analyzed by flow cytometry. D. The cell cycle distribution was analyzed by flow cytometry. E. The protein levels of MEK and p-ERK were analyzed by Western blotting. Data are the means ± SD, n=3, *P<0.05, **P<0.01 vs. Control, *P<0.05, **P<0.01 vs. Model.

totic cells decreased (Figure 4C), mitosis was enhanced (Figure 4D), and pERK/ERK and pMEK/MEK increased (Figure 4E). All the effects were inversed by the MEK inhibitor U0126. These results suggest that BPIFB1 reduced NPC-KT cell viability and proliferation via the MEK/ERK signaling pathway.

BPIFB1 enhanced the inhibition of the chemotherapeutic agent on the viability of NPC-KT cells

To investigate the role of BPIFB1 in chemoresistance, NPC-KT cells were transfected with a BPIFB1 plasmid and incubated with cisplatin.
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Figure 5. BPIFB1 enhanced the chemo-sensitivity of NPC-KT cells. NPC-KT cells were transfected with BPIFB1 plasmids and treated with cisplatin (Cis) (A) or Paclitaxel (Pct) (B), and the proliferation was measured using an MTT assay. Data are the means ± SD, n=3, *P<0.05, vs. Control, #P<0.05 vs. Pct or Cis group.

Discussion

BPIFB1 or LPLUNC1, secreted by nasopharyngeal epithelia or the minor mucosal glands of the respiratory and upper digestive tracts [8], has been proved to regulate host immune responses in respiratory diseases [16] and is upregulated in COPD and relevant to its severity [17], but it is downregulated in NPC tissues [18]. The effects and mechanism of BPIFB1 in NPC are reported as limited, and whether BPIFB1 downregulation is associated with the progression and prognosis of NPC has not been clarified. In this research, the effects and the molecular mechanisms of BPIFB1 on the proliferation, apoptosis, cell cycle, and chemoresistance of NPC cells were investigated. Our results indicated that the overexpression of BPIFB1 inhibited cell growth, promoted apoptosis through cell cycle arrest at the G0/G1 phase, and regulated the MEK/ERK signaling pathway. Additionally, BPIFB1 could enhance the inhibitory effects of chemotherapeutic agents on cancer cells, which suggests that BPIFB1 could be a potential target for the treatment of NPC.

Although the studies on BPIFB1 are limited, there are still studies on the effects and mechanisms of BPIFB1 on NPC. Wei et al. first demonstrated that BPIFB1 markedly inhibited NPC cell migration, invasion, and metastatic abilities via the inhibition of VTN (a BPIFB1-interacting protein), leading to the inactivation of the FAK/Src/ERK signaling pathway [19]. BPIFB1 also inhibited NPC cell growth by downregulating MAPK and the cyclin D1/E2F pathways [12], and the inflammation related signaling pathway (signal transducer and activator of transcription 3 (STAT3)) activity [20]. The previous studies were in accordance with our findings that BPIFB1 could regulate the cell cycle and the MAPK/ERK signaling pathway. The MAPK pathway is shared by four downstream cascades, including the Jun amino-terminal kinases (JNK1/2/3), the extracellular signal-related kinases (ERK1/2), p38 and ERK5, among which ERK is reported to be related to cell proliferation, migration, and apoptosis and plays an important role in tumor progression and cancer treatment [21]. In our research, BPIFB1 could inhibit NPC cell proliferation and promote cell apoptosis through the inactivation of the MEK/ERK signaling pathway.

Chemoresistance is a crucial challenge in the treatment of cancer and is related to poor prognosis in cancer patients [22]. NPC is relatively sensitive to radiotherapy, but for late stage patients, chemoradio resistance was still a challenge and affects the prognosis of NPC patients [4]. We et al. first demonstrated that BPIFB1 enhanced the sensitivity of NPC cells to ionizing radiation by inhibiting VTN (a BPIFB1-binding protein) expression [23]. Our results confirmed the finding that BPIFB1 overexpression could increase the chemosensitivity of NPC-KT cells. More interestingly, researchers found that BPIFB1 plays an important role in inducing innate immune responses. BPIFB1, along with PA-MSHA (Pseudomonas aeruginosa mannose sensitive hemagglutination) can activate CD14/TLR4/MyD88 and induce the upregulation of pro-inflammatory cytokines [24]. This research indicated that BPIFB1 may also regulate the inflammatory response in the tumor microenvironment and induce the apoptosis of cancer cells.
Conclusion

Our research demonstrated that BPIFB1 can affect cell proliferation and the cell cycle, promote the apoptosis of NPC cells by regulating the MEK/ERK signaling pathway, and enhance the inhibition effects of chemotherapeutic agents on cancer cells. The results indicated that BPIFB1 could be a potential target for the treatment of NPC.

Disclosure of conflict of interest

None.

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References

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