Krüppel-like factor 4 promotes high-mobility group box 1-induced chemotherapy resistance in osteosarcoma cells

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Osteosarcoma is the most common primary malignant bone tumor, and the frequent acquisition of chemoresistance is often an obstacle to achieving favorable outcomes during chemotherapy. Recently, Krüppel-like factor 4 (KLF4) has been shown to be associated with chemoresistance in a few tumors; however, the involvement of KLF4 in chemotherapy resistance in osteosarcoma cells remains unknown. In this study, quantitative real-time PCR and western blot analysis revealed that KLF4 expression was significantly increased in response to cisplatin, methotrexate and doxorubicin treatment in osteosarcoma cells, and knockdown of KLF4 increased sensitivity to these anticancer drugs by decreasing cellular clonogenic ability and increasing apoptosis. Moreover, our data suggest that KLF4-regulated drug resistance might, at least partially, positively regulate chemoresistance in osteosarcoma cells in our previous study. In summary, this study highlights the significance of KLF4/HMBG1 interaction in regulating chemotherapy resistance, and suggests that targeting KLF4/high-mobility group box 1 may be a therapeutic strategy for osteosarcoma chemotherapy.

Materials and Methods

Cell culture and drug treatment. The osteosarcoma cell lines (MG-63, SaOS-2 and U-2 OS) were originally obtained from the American Type Culture Collection (ATCC, Rockville, MI, USA).
USA), and cultured in Eagle’s minimum essential medium supplemented with 10% FBS in a 5% CO₂ and humidified atmosphere at 37°C. For drug treatment, cisplatin (Sigma-Aldrich, Taufkirchen, Germany) was dissolved in N,N-dimethylformamide at a concentration of 15 mg/mL and added to the culture medium in the indicated concentrations. Methotrexate was dissolved in 0.01N NaOH and further diluted in PBS to a concentration of 10 mg/mL. Doxorubicin (Sigma-Aldrich) was dissolved in sterile water at a concentration of 20 mM and added to the culture medium in the indicated concentrations.

RNA preparation and quantitative real-time PCR. Total RNA was extracted by Trizol reagent, and first strand cDNA was synthesized with SuperScript II (Invitrogen, Carlsbad, CA, USA). The transcript levels were detected by quantitative real-time PCR analysis using SYBR Green Taq ReadyMix (Takara, Japan). The expression levels of HMGB1 and KLF4 were normalized to that of β-actin mRNA, which served as an endogenous control. The following sequences of PCR primers were used: β-actin: 5'-AGGGGGCGGACTGTACATTACT-3' (forward), 5'-GGGCGGACCACATGTACCTCCTC-3' (reverse); KLF4: 5'-GCCGCT GTGATGACGAGGACATCCTG GGGCGTTCTTGTGTTGCTCT-3' (reverse); HMGB1: 5'-TCAAAGGAGAACATCTGG CCTGTGCTT-3' (forward), 5'-TCAAAGGAGAACATCTGG CCTGTGCTT-3' (reverse). qRT-PCR was performed on the ABI 7500 thermocycler (Applied Biosystems, Carlsbad, CA, USA). The relative expression levels were calculated using the 2^(-ΔΔCt) method.

Western blot. Cells were lysed with 200 μL of lysis buffer and subjected to western blot analysis.16) Approximately 50 mg of total protein was separated by 10% SDS-PAGE, transferred to a PVDF membrane and incubated with the appropriate antibodies. The antibodies to HMGB1 and KLF4 were purchased from GeneChem (Shanghai, China). β-actin antibody was obtained from Kangchen Biotechnology (Shanghai, China). The protein bands were visualized using ECL detection (Applygen, Beijing, China).

Kruppel-like factor 4 expression plasmid and small interfering RNA transfection. The coding sequence of KLF4 was amplified and cloned into pcDNA3.1 vector to generate KLF4 expression plasmid (pcDNA3.1-KLF4), and the empty pcDNA3.1 vector was used as control (pcDNA3.1-Con). Primers used for KLF4 coding sequence are as follows: 5'-GGATCCATGAGGACGGAC GACCCCTGGC-3' (forward) and 5'-GAATTCCTAGATGACGAGGACATCCTG GGGCGTTCTTGTGTTGCTCT-3' (reverse). Specific siRNA for KLF4 (si-KLF4) and scrambled siRNA (negative control, si-NC) were purchased from GeneChem (Shanghai, China). si-KLF4 and scrambled siRNA were transfected to a PVDF membrane and incubated with the appropriate antibodies. The antibodies to HMGB1 and KLF4 were purchased from GeneChem (Shanghai, China). The protein bands were visualized using ECL detection (Applygen, Beijing, China).

ChIP assay. ChIP assays were performed as previously described.17) Briefly, 5 × 10⁷ cells were crosslinked with 1% formaldehyde for 10 min at 37°C, and then sonicated under the condition of a Bioruptor™x2000 and fragmented chromatin were immunoprecipitated using an anti-KLF4 antibody or control goat IgG antibody and a Chromatin Immunoprecipitation Kit (Millipore, Boston, MA, USA). Immunoprecipitated DNA were purified and quantitatively analyzed by real-time PCR using a HMGB1 gene promoter-specific primer set (5'-TGGTGGCTCCATTTTGAAA-3' and 5'-ATGGTTTACGATAGTTTGA-3') to determine the enrichment. Values were presented as relative to DNA input. The assays were carried out in three replicates and relative fold change was calculated using the −ΔΔCt method.

Electrophoretic mobility shift assay. Electrophoretic mobility shift assay (EMSA) was performed using nuclear extracts according to the protocol of the Light Shift Chemiluminescent EMSA Kit (Pierce, Carlsbad, CA, USA). Double-stranded DNA probes (5'-GATGAAATGTTGGGCAAGAAGAGGCGGAGACCTGTTGGTGCTCT-3' and 5'-AGGAGACCCACAGGTTCCTCCCCCTCTTTGCACATTAC-3') were generated. Double-stranded mutant DNA probes: 5'-ATATAG AGCAAAGGGGGGGGGAGACCTAAGATG-3', 5'-CAT CATCTAGGGTCCTCCCCCTCTTGCTATAT-3'. Nucleotides were labeled using biotin 3'-end DNA labeling (Pierce). Detection of supershift mobility was assessed by reaction of KLF4 antibody (Santa Cruz Biotechnology) with nuclear extracts for 30 min before addition of the labeled nucleotides for the EMSA.

Reporter gene assays. Five different fragments of the promoter region of HMGB1 were amplified and ligated to pGL3 GFP reporter. MG-63 cells were cultured to approximately 80% of the plates, and co-transfected with HMGB1 promoter−reporter constructs and KLF4 expression plasmid using Lipofectamine 2000 (Invitrogen), according to the manufacturer’s protocol. The cells were cultured for another 4 h, and the GFP expression levels were quantified by flow cytometry.

Cell proliferation, colony formation and apoptosis assays. Cell proliferation was evaluated by MTT assay. Cells were seeded in a 96-well plate at 5000 cells per well and incubated for
24 h, then exposed to different doses of anticancer drugs for 48 h. Then, the cells were added to 10 mg/mL MTT (Sigma Chemicals, St. Louis, MO, USA). After incubation for 2 h, the reaction was terminated by removal of the supernatant followed by adding 200 μL of DMSO. The optical density at 570 nm was measured with a microplate reader (Bio-Rad, Hercules, CA, USA). For colony formation assays, approximately 500 cells were placed in complete growth media each 35-mm dish and allowed to grow for 6 h. Then, with different doses of anticancer drugs, were added to each dish. After 24 h of treatment, the anticancer drugs were removed by adding fresh complete growth media, and cells were allowed to grow until visible colonies formed, and were stained and counted. For cell apoptosis assays, cells were collected and the cell apoptosis ratio was analyzed using the Annexin V-FITC Apoptosis Detection Kit (BD Biosciences, San Diego, CA, USA), according to the manufacturer’s instructions.

**Statistical analysis.** Data are expressed as the mean ± SD. Differences between groups were analyzed by Student’s t-test or ANOVA, and P < 0.05 was considered to be statistically significant.

**Results**

Kruppel-like factor 4 is upregulated in response to chemotherapy in osteosarcoma cells. We first assayed the effects of the anticancer reagents cisplatin (Cis), methotrexate (Mtx) and doxorubicin (Dox) on the expression of KLF4 in osteosarcoma cells, respectively. As shown in Figure 1a, quantitative real-time PCR (qRT-PCR) revealed that these drugs significantly
promoted increased mRNA expression levels of KLF4 in MG-63, SaOS-2 and U-2OS cell lines after treatment with these anticancer agents for 48 h. In addition, western blot analysis showed similar results (Fig. 1b). These findings suggest that KLF4 is upregulated in response to chemotherapy in osteosarcoma cells, and, thus, may play an important role in regulating the chemoresistance in osteosarcoma cells.

**Suppression of Krüppel-like factor 4 increases sensitivity to chemotherapy in vitro.** To examine the potential role of KLF4 in the regulation of anticancer drug sensitivity of osteosarcoma cells, we established KLF4 knocked down osteosarcoma cells by KLF4-specific siRNA (MG-63/si-KLF4, SaOS-2/si-KLF4) and the corresponding control cells by scrambled siRNA (MG-63/si-Con and SaOS-2/si-Con). qRT-PCR and western blot analysis validated that MG-63/si-KLF4 and SaOS2/si-KLF4 cells expressed significantly lower mRNA and protein levels of KLF4 than in the corresponding control cells. Using the MTT assays, we found that knockdown of KLF4 in these cells rendered them significantly more sensitive to Dox-induced, Cis-induced and Mtx-induced cell injury, respectively (Fig. 2a), and this was also associated with drastically decreased clonogenic ability (Fig. 2b) and high levels of apoptotic cell death (Fig. 2c). In contrast, when KLF4 was overexpressed in MG-63 and SaOS-2 cells, we observed an opposite effect on Dox-induced, Cis-induced and Mtx-induced cell injury (Fig. 3a), clonogenic ability (Fig. 3b) and apoptotic cell death (Fig. 3c). Taken together, these data suggest that targeted suppression of KLF4 increases sensitivity to chemotherapy in osteosarcoma cells in vitro.

**Krüppel-like factor 4 positively regulates high-mobility group box 1 expression in osteosarcoma cells.** Our previous report revealed that chemotherapy agents, including Cis, Mtx and...
Dox, induced HMGB1 expression in osteosarcoma cells.\(^{(16)}\) To determine whether KLF4 is responsible for HMGB1 transcriptional regulation, we examined the effect of KLF4 knockdown or overexpression on the expression of HMGB1 in MG-63 and SaOS-2 cells. qRT-PCR and western blot analysis both revealed that knockdown of KLF4 inhibited mRNA and protein expression levels of HMGB1 in MG-63 and SaOS-2 cells, while overexpression of KLF4 resulted in increased HMGB1 expression (Fig. 4). This suggests that KLF4 may positively regulate HMGB1 expression in osteosarcoma cells.

**Kruppel-like factor 4 binds directly to the high-mobility group box 1 promoter and regulates its transcription.** To determine whether KLF4 directly regulates HMGB1 expression, we firstly performed promoter reporter assays to identify the KLF4-responsive region on the HMGB1 promoter. MG-63 cells were co-transfected with pcDNA3.1-KLF4 expression plasmid and pGL3 GFP reporter with five different fragments of the promoter region of HMGB1 (Fig. 5a). As shown in Figure 5b, progressive deletion of the promoter sequence from −1899 to −456 did not affect the HMGB1 promoter activity, but deletion between −456 and −240 significantly attenuated the promoter activity. The data indicate that the region between −456 and −240 is critical for responding to KLF4. In addition, EMSA also revealed that KLF4 could directly bind to the region between −456 and −240 on the HMGB1 promoter. As shown in Figure 6a, preincubation of nuclear extracts from Cis-treated MG-63 cells with MHGB1 probes resulted in DNA/protein complex formation, which was competed on with the addition of unlabeled (cold) probes (Fig. 6a, lanes 2–3), whereas preincubation with mutant unlabeled probes had no effect on DNA/protein complex formation (Fig. 6a, lane 4). Moreover, addition of KLF4 antibody resulted in the formation of a shift band (Fig. 6a, lane 5). These data suggest that KLF4 is able to bind to the KLF4-responsive core region on the HMGB1 promoter. Finally, ChIP assays combined with quantitative PCR (ChIP-qPCR) were performed in MG-63 and SaOS-2 cells treated with 50 uM Cis for 48 h and it was found that KLF4 could directly bind to the KLF4-responsive core region on the HMGB1 promoter in MG-63 and SaOS-2 cells in vivo. As shown in Figure 6b, immunoprecipitation of the chromatin/protein complex with KLF4 resulted in the enrichment of KLF4 protein at the HMGB1 promoter region near the region between −456 and −240. The promoter sequence of HMGB1 was also analyzed using Matinspector Professional (www.genomatix.de) and TESS (www.cbil.upenn.edu). The KLF4-binding site was existed at region −379 to −335 bp upstream the TSS of HMGB1. Based on our recent study and previous work,\(^{(16)}\) Fig. 4. **Kruppel-like factor 4 (KLF4) positively regulates high-mobility group box 1 (HMGB1) expression in osteosarcoma cells.** (a,b) KLF4 was overexpressed by transfecting with KLF4 expression plasmids (pcDNA3.1-KLF4) or knocked down by KLF4-specific siRNA (si-KLF4) in MG-63 and SaOS-2 cells, and qRT-PCR was performed to validate the KLF4 mRNA levels. (c) The expression levels of KLF4 as well as HMGB1 protein were examined by western blot, which revealed that KLF4 positively regulates HMGB1 expression in osteosarcoma cells. *P < 0.05 versus the control.
here we provide a model of KLF4-induced chemotherapy in osteosarcoma cells, in which increased expression of KLF4 during chemotherapy activates HMGB1, thus further promoting autophagosome maturation and autophagy (Fig. 7). Therefore, KLF4/HMGB1 interaction is a potential therapeutic target for use in osteosarcoma.

Discussion

Osteosarcoma is a high-grade malignant bone tumor. Although the introduction of chemotherapy has reduced its mortality, the clinical effectiveness is limited by the emergence of drug resistance, which ultimately leads to poor therapeutic outcomes. Consequently, it is necessary to discover novel molecules regulating drug resistance to develop more effective targeted therapies. In the present study, we found that KLF4 expression was significantly increased in response to treatment with commonly used anticancer drugs in osteosarcoma cells. In addition, we demonstrated for the first time that knockdown of KLF4 increased sensitivity to these anticancer drugs in osteosarcoma cells in vitro, by decreasing clonogenic ability and increasing apoptosis. Finally, our data suggested that KLF4-mediated drug resistance might be through directly binding to the HMGB1 promoter and positively regulating its transcription, which highlighted the significance of KLF4/HMGB1 interaction in regulating chemotherapy resistance in osteosarcoma cells.

Krüppel-like factor 4 contains C2H2-type zinc fingers at the C-terminal and regulates the expression of target genes by binding to GC-rich or CACCC promoters. Recent studies have shown that the expression of KLF4 is altered in certain types of cancer. For example, KLF4 was downregulated in gastrointestinal (GI) tumors and negatively-regulated GI cancer EMT,13 while the expression of KLF4 was upregulated in other types of cancers, such as breast cancer,19 breast cancer,11 and head and neck squamous cell cancer,11 and functioned as an oncogene which enhanced tumor development and progression. Increasing evidence has revealed that KLF4 is associated with cancer cell chemotherapy resistance. To address the involvement of KLF4 in chemotherapy resistance in osteosarcoma cells, we first examined the expression alteration of KLF4 in osteosarcoma cells in response to anticancer reagents (Cis, Mtx and Dox), and found that these drugs significantly promoted increased KLF4 mRNA and protein expression levels.
Further *in vitro* study revealed that suppression of KLF4 by siRNA increases sensitivity to chemotherapy. In contrast, when KLF4 was overexpressed in MG-63 and SaOS-2 cells, we observed an opposite effect on sensitivity to chemotherapy. Our data are consistent with previous reports that increased KLF4 contributed to drug resistance. For example, with a higher level of KLF4, hepatocarcinoma cell line T3A-A3 was found to be more resistant to Cis than HepG2 cells, which have lower levels of KLF4 expression, and KLF4 knockdown reduced Cis resistance in T3A-A3 cells. Tai *et al.* report that enforced KLF4 expression in head and neck squamous cell carcinoma (HNSCC) SAS cells significantly increased multidrug resistance. Taken together, KLF4 may have a potential role in the regulation of chemotherapy sensitivity of reported cancers and other cancers.

Our previous study revealed that HMGB1 promotes drug resistance in osteosarcoma, however, the regulation on HMGB1 expression remains unclear. As HMGB1 contains putative KLF4-binding elements in its promoter region, we hypothesized that KLF4 might affect the expression of HMGB1. In this study, through KLF4 knockdown or overexpression, we found that KLF4 could positively regulate HMGB1 expression in osteosarcoma cells. HMGB1 promoter reporter assays, electrophoretic mobility shift assay and chromatin immunoprecipitation assays validated that KLF4 could directly bind to KLF4-binding elements (three “CACCC” elements) on the region between −456 and −240 of the HMGB1 promoter in osteosarcoma cells. Our data are consistent with a previous report that KLF4 could bind to HMGB1 in mouse RAW264.7 macrophages when encountered LPS treatment.

Autophagy is a catabolic process critical to maintaining cellular homeostasis and responding to various nutrient starvation or metabolic stress. Many anticancer reagents result in enhanced autophagy, which facilitates the cancer cells’ resistance to chemotherapy treatment, and the abrogation of autophagy potentiates the re-sensitization of therapeutic-resistant cancer cells to the anticancer treatment. Recent reports have demonstrated that HMGB1 is a critical regulator of autophagy in fibroblasts, leukemia, colon, pancreatic cancer and osteosarcoma cells. Our previous study also demonstrated that knockdown of HMGB1 or inhibition of autophagy increases apoptosis, and reverses drug resistance in osteosarcoma cells. In this study, we validated that KLF4, upregulated during chemotherapy treatment in osteosarcoma cells, is a positive regulator of HMGB1.

In summary, our results demonstrate that KLF4 functions as an inducer of chemotherapy resistance in osteosarcoma, at least partially, by binding to and activating the HMGB1 promoter, which suggests that targeting KLF4/HMGB1 may be a therapeutic strategy for osteosarcoma chemotherapy.

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Disclosure Statement

The authors have no conflict of interest to declare.

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