

# Aurora-B Promotes Osteosarcoma Cell Growth and Metastasis Through Activation of the NPM1/ERK/NF- $\kappa$ B/MMPs Axis

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**Purpose:** Osteosarcoma (OS) is the most common primary malignant tumor of the bone in young adolescents and children. We explored the underlying mechanism of Aurora-B in promoting OS cell proliferation and metastasis.

**Patient and Methods:** Bioinformatics was employed to predict the substrate of Aurora-B. IHC and Western blot were used to confirm the correlation between Aurora-B and NPM1. ERK/NF- $\kappa$ B pathway-related proteins were detected by Western blot and immunofluorescence (IF). CCK8, wound healing, transwell, and Tunel assays were used to identify the cell proliferation, migration and apoptosis potential. Spontaneous metastasis xenografts were established to confirm the role of Aurora-B and NPM1.

**Results:** Aurora-B promotes NPM1 phosphorylation on Ser125. The phosphorylation of NPM1<sup>Ser125</sup> induced by Aurora-B activates the ERK/NF- $\kappa$ B signaling. Further study revealed that Aurora-B promotes proliferation, migration and inhibits apoptosis via phosphorylating NPM1 in vitro and in vivo.

**Conclusion:** Aurora-B promotes OS malignancy via phosphorylating NPM1<sup>Ser125</sup> and activating ERK/NF- $\kappa$ B signaling.

**Keywords:** Aurora-B, osteosarcoma, NPM1, ERK, NF- $\kappa$ B

## Introduction

Osteosarcoma (OS) is the most common malignant tumor of the bone and is prevalent in both young adolescents and children.<sup>1</sup> Standard treatment for OS mainly includes the surgical removal of localized lesions, combined with multiple chemotherapeutics.<sup>2</sup> These approaches increase the 5-year survival rate of patients in the absence of metastasis from less than 20% to almost 70%. However, no further breakthroughs for OS management have occurred, owing to the drug resistance and metastasis.<sup>3</sup> Novel and advanced molecular agents for targeted OS therapies are therefore urgently required.

Aurora-B forms a complex with INCEP, Survivin and Borealin and participates in a range of cell biological processes to ensure the correct interaction of kinetochores and spindle microtubules. Aurora-B deficiency will lead to the arrest in metaphase in non-malignant cells.<sup>4</sup> Numerous studies have demonstrated that Aurora-B is aberrantly expressed in various malignant tumors<sup>5-8</sup> and facilitates cell cycle progression through the regulation of cell cycle-related targets. Gonzalez-Loyola et al reported that Aurora-B activates CDK1 via suppressing the p53/p21WAF1/CIP1 axis to promote cell cycle progression and cell survival.<sup>7</sup> Guise

and colleagues showed that Aurora-B phosphorylates Class IIa HDACs to induce cell cycle progression.<sup>9</sup> In addition, Aurora-B facilitates lymphoma cell proliferation and survival through the activation of AKT/mTOR signaling.<sup>10</sup> Although the overexpression or amplification of Aurora-B contributes to tumor cell malignancy,<sup>11–13</sup> the role of Aurora-B in regulating OS cell growth and metastasis remains elusive.

In this study, we found that NPM1 is overexpressed in osteogenetic OS tissues. Aurora-B was found to promote OS cell growth and metastasis through its ability to phosphorylate NPM1<sup>Ser125</sup> and activate ERK/NF- $\kappa$ B signaling. These data suggest that targeting the Aurora-B/NPM1/ERK/NF- $\kappa$ B axis represents a novel therapeutic candidate for the prevention of OS metastasis.

## Materials and Methods

### Bioinformatics

The bioinformatics prediction tool OncoPrint was used to examine NPM1 expression in OS (<https://www.oncoPrint.com/resource/login.html>), database 200063 s at were selected; The Molecular Interaction Database was used to explore the correlation between NPM1 and Aurora-B.

### Cell Culture and Transfection

The OS cell lines U2-OS and 143B were ordered from the cell bank of Type Culture Collection of Chinese Academy of Sciences (Shanghai, China). Cells were maintained in DMEM (Gibco) plus 10% FBS. Lentivirus-Vectors were compound by GENECHM (Shanghai, China). The sequence was as follows: shAurora-B, 5'-AGAGCTGCACATTTGACGA-3'; shRNA-NC, 5'-TTCTCCGAACGTGTCACGT-3'. 143B and U2-OS cells were infected with  $5 \times 10^5$  PFU Lentivirus-Vectors (MOI=50) and treated with 0.6  $\mu$ g/mL puromycin for cell selection.

### Specimens and Immunohistochemical Analysis

A total of 87 tissue sections histologically diagnosed with OS were obtained from the First Affiliated Hospital of Nanchang University, China. No patients received therapy prior to biopsy. All experiments were approved by the ethics committee of the First Affiliated Hospital of Nanchang University (Jiangxi, China; NO.Y2019-126) and followed the Declaration of Helsinki. All the subjects were informed of the contents, latent risks, objectives and signed written informed consents.

Immunoperoxidase procedures (Maxvision procedure) were performed to detect relative protein expression. Briefly, sections were heated in citrate buffer for 20 min and probed with anti-Aurora-B (1:500, Abcam, MA, USA) and pNPM1<sup>Ser125</sup> (1:1000, Abcam) antibodies overnight. Stained sections were scored by two doctors, respectively. Tissues showing yellow and brown nuclear staining were used to indicate a positive result, and tissues lacking staining in the nucleolus were considered negative. Immunohistochemical expression of Aurora-B and pNPM1<sup>Ser125</sup> were determined as “-”, “+”, “++”, and “+++” according to the coloring intensity and the percentage of positively stained tumor cells.

### qPCR

Total RNA from OS cells was extracted with Trizol (Invitrogen). Aurora-B expression was evaluated by qRT-PCR. GAPDH was used as an endogenous reference gene. The final reaction mixture of all the amplification reactions was 20  $\mu$ L. Primer sequences were listed as follows: Aurora-B sense 5'-AGAAGGAGAACTCCTACCCCT-3', Aurora-B antisense 5'-CGCGTTAAGATGTCGGGTG-3'; GAPDH sense 5'-CAGGGCTGCTTTAACTCTGGT-3', GAPDH antisense 5'-GATTTTGGAGGG ATCTCGCT-3'. The comparative Ct method was used to calculate relative gene expression.

### Wound Healing Assays

Cells ( $5 \times 10^6$ ) were seeded into six-well plates and cultured for 12 h. Confluent cells were scratched using a slender pipette tip and washed using PBS and complete DMEM. Images were taken 24 h later and compared to T=0. Image J (NCBI) was used to quantify the migrated distances.

### Transwell Invasion Assays

Cell invasion assay was assessed using matrigel Chamber assays (BD Bioscience, NJ, USA) according to the manufacturer's protocol. Briefly, cells ( $2 \times 10^5$ ) suspended in 250  $\mu$ L of serum-free DMEM were added to the upper chamber insert and complete medium (15% FBS) was used as a chemoattractant and added to the bottom chamber. After 24 h, chambers with invaded cells were fixed and stained with Crystal Violet Solution (Beyotime, China). Five high-power images were randomly taken and the number of transmembrane cells quantified using Image J (NCBI).

### CCK8 Assays

Cells (~4000) were counted and plated into 96-well plates and cultured in complete media (10% FBS) for 0 h, 24 h and

48 h, respectively. CCK-8 reagent (10  $\mu$ L, Dojindo Laboratories, Japan) was added for 2 h and cell viability was defined as the absorbance at 450nm detected on a microplate reader.

## Tunel Assays

Apoptotic activity in OS cells was assessed through TUNEL staining (Roche) according to the manufacturer's protocols. Briefly, samples were fixed in freshly prepared 4% PFA and blocked in 3% H<sub>2</sub>O<sub>2</sub> in methanol. Cells were then stained with TUNEL reagent for 1 h followed by incubation with 50  $\mu$ L of converter-POD and DAB. Slides were imaged via microscopy.

## Western Blot Analysis

Cells were lysed in RIPA lysis buffer with PMSF and Bradford assays were performed to quantitate total protein concentrations. Approximately 20  $\mu$ g of protein was loaded onto the sample wells and transferred onto Nitrocellulose membranes. Membranes were incubated with 5% fat-free milk and probed with primary antibodies (anti-Aurora-B 1:5000; anti-NPM1, anti-pNPM1<sup>Ser125</sup>, anti-ERK1/2 and anti-pERK1/2<sup>Thr202/Tyr204</sup> 1:1000; anti-MMP9, anti-MMP2, 1:5000; anti-XIAP, anti-BCL-XL, anti- $\beta$ -actin, 1:2000) for 8 h. Primary antibodies were purchased from Abcam (Cambridge, MA, USA). Species-specific secondary antibodies (dilution ratio of 1:2000) were purchased from ZSGB-BIO, China. Immune complexes were visualized using EasySee Western Blot kits (Transgen, China).

## Immunofluorescence

U2-OS and 143B cells plated onto 24-well plates were grown to 70% confluency and treated with 25  $\mu$ M 2-CIHyA for 30 min. Cells were fixed with 4% PFA and blocked in 5% (w/v) BSA for 30 mins. Cells were labeled with NF- $\kappa$ B p65 antibodies (1:100 dilution; Abcam) and washed in PBST. Slides were incubated with Rhodamine (TRITC)-Conjugate Goat anti-rabbit IgG (1:25; ZSGB-BIO, China). PBST-washed slides were treated with VECTASHIELD Anti-fade mounting medium (VECTOR Laboratories, Inc., 30 Ingold Road, Burlingame, CA 94,010 USA). Images were obtained using a confocal laser scanning microscope (Zeiss/LSM 710, Germany). For quantification of immunofluorescence images, a minimum of 50–100 NF- $\kappa$ B positive cells were counted to determine the number of NF- $\kappa$ B nucleus translocation cells.

## Spontaneous Metastasis Xenografts

Male BALB/c nude mice (4-weeks old) were provided by the Department of Laboratory Animal Science of Nanchang University (Nanchang, China). Spontaneous metastasis xenografts were established as previously described. Mice were randomly selected (n=12) and intratumorally injected with Aurora-B, shAurora-B, NC- Aurora-B and NC- shAurora-B lentivirus (10<sup>7</sup> PFU, 100  $\mu$ L) for 7 days. Mice were sacrificed after 6 weeks of orthotopic transplantation. Tumor sizes were assessed on a graduated cylinder filled with water in which changes in water height indicated tumor volume. Lung metastatic nodules were assessed by H&E staining and stereomicroscope (Nikon SMZ1500). All animal experiments were approved by the First Affiliated Hospital of Nanchang University (Jiangxi, China; NO.Y2019-126), and performed according to the National Institutes of Health (NIH) Guide for the Care and Use of Laboratory Animals (1996).

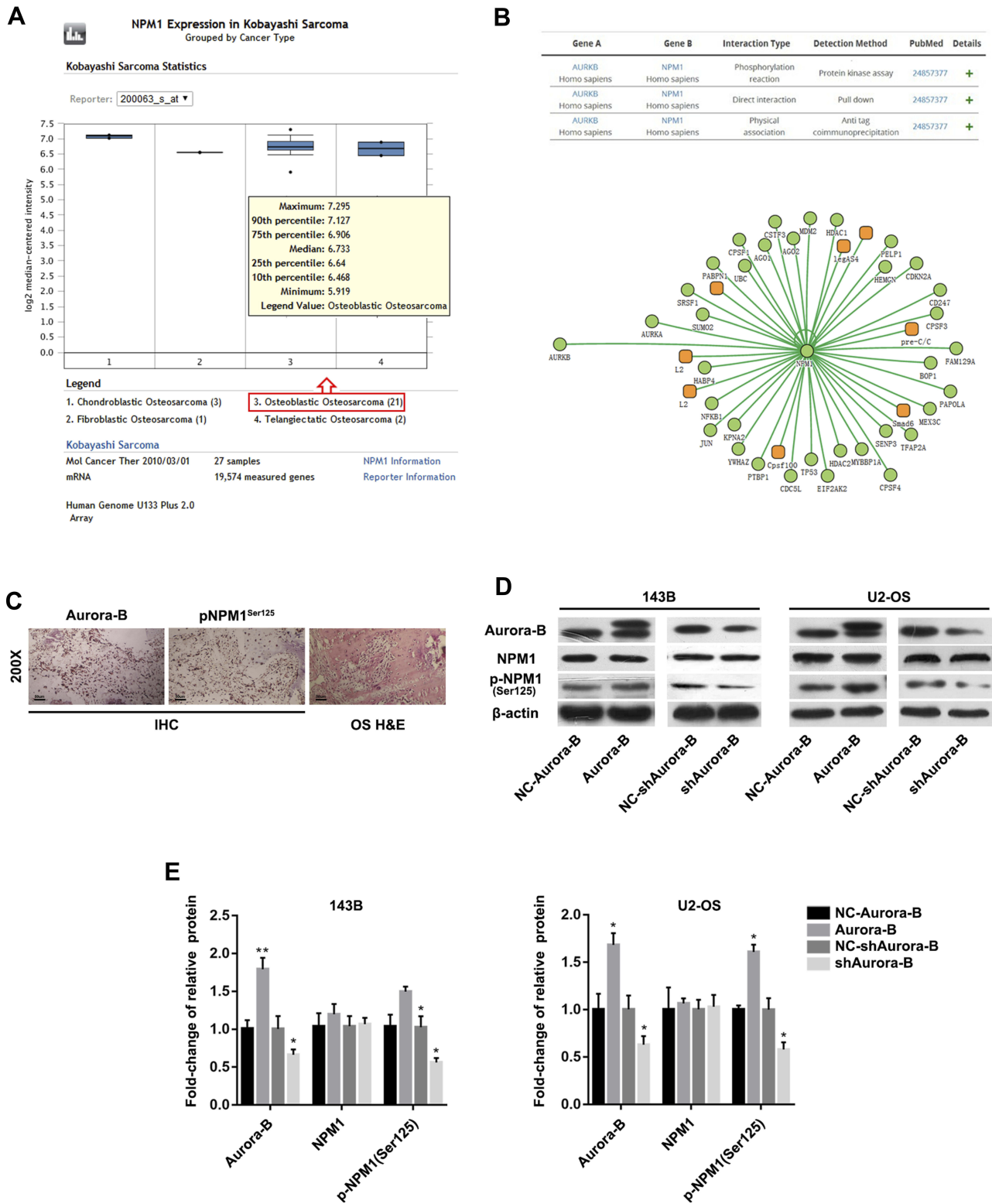
## Statistical Analysis

Non-parametric Wilcoxon rank-sum tests were used to analyze count data. The correlation between Aurora-B and pNPM1<sup>Ser125</sup> protein levels were determined by Spearman's rho tests. Measurement data are shown as the mean  $\pm$  standard deviation. A Student's *t*-test was used for two-sample analysis. A one-way ANOVA was used for multiple-sample analysis. Data were analyzed using SPSS statistical software (Version 13.0).

## Results

### Aurora-B Promotes the Phosphorylation of NPM1

To explore NPM1 expression in OS samples and the potential interaction between Aurora-B and NPM1, bioinformatics was performed using the Molecular INTERaction Database; Oncomine Database). The results indicated that NPM1 was up-regulated in osteogenetic OS samples and that Aurora-B interacts with NPM1 (Figure 1A-B). In addition, we detected the Aurora-B and p-NPM1 expression in 87 samples of osteogenetic OS by immunohistochemistry (Figure 1C). The correlation between Aurora-B and phosphorylated NPM1 expression was analyzed using bivariate Spearman assays (Table 1). The results indicated that Aurora-B was positively related to phosphorylated NPM1. We further investigated the effects of Aurora-B alterations on NPM1 phosphorylation. Cells were infected with Aurora-B, NC-Aurora-B, shAurora-B, and NC-shAurora-B lentivirus, respectively. Aurora-B, NPM1 and



**Figure 1** Aurora-B promotes the phosphorylation of NPM1. **(A)** NPM1 expression in osteosarcoma. Data were obtained from the Oncomine database. **(B)** Bioinformatics (The Molecular INTERaction Database) revealed NPM1 as a substrate of Aurora-B. **(C)** Representative images of IHC staining of Aurora-B and pNPM1<sup>Ser125</sup> and H&E stained images in osteosarcoma patients. pNPM1<sup>Ser125</sup> is shown as brownish-yellow particle deposition and was expressed in the nucleus. Experimental tissues were osteosarcoma as confirmed by H&E staining. **(D)** Western blot analysis of Aurora-B, NPM1 and pNPM1<sup>Ser125</sup> in 143B and U2-OS cells stably transfected with NC-Aurora-B overexpression, Aurora-B overexpression, NC-shAurora-B, and shAurora-B lentivirus. **(E)** Quantification of the Western blot assay. Values represent mean±SD (n = 3). \*p<0.05, \*\*p<0.01.

p-NPM1<sup>Ser125</sup> expression were confirmed by Western blot. The results indicated that p-NPM1<sup>Ser125</sup> expression was higher in cells infected with Aurora-B compared to control groups. Lower p-NPM1<sup>Ser125</sup> levels were observed in cells infected with shAurora-B lentivirus compared to control groups. However, the total levels of NPM1 did not differ between the groups (Figure 1D and E). The reason for the different bands of Aurora-B may be related to different splice variants/isoforms. Collectively, these results suggest that Aurora-B promotes NPM1 phosphorylation on Ser125.

## Aurora-B Activates ERK/NF- $\kappa$ B Signaling Through the Phosphorylation of NPM1<sup>Ser125</sup>

To explore the downstream effects of Aurora-B-mediated NPM1 phosphorylation in OS cells, ERK1/2, pERK1/2, MMP2, MMP9, and apoptotic Bcl-xL and XIAP levels were assessed in U2-OS and 143B cells transfected with Aurora-B and scrambled controls. Western blot analysis revealed that pERK1/2, MMP2, MMP9, Bcl-xL and XIAP levels were significantly higher in Aurora-B vs control groups. In contrast, pERK1/2, MMP2, MMP9, Bcl-xL and XIAP level were significantly lower in cells infected with LV/shAurora-B compared to the control group. This loss of protein expression could be recovered by NPM1 overexpression (Figure 2A and B). Furthermore, the nuclear translocation of NF- $\kappa$ B was examined by confocal microscopy. The results revealed that NF- $\kappa$ B expression and nucleus translocation were enhanced in OS cells infected with LV/Aurora-B compared to the control group. An opposing phenotype was observed in cells transfected with NV/shAurora-B (Figure 2C and D). These results suggest that Aurora-B activates ERK/NF- $\kappa$ B signaling via phosphorylating NPM1<sup>Ser125</sup>.

**Table 1** Correlation Between Aurora-B and pNPM1<sup>Ser125</sup> Expression in Osteosarcoma

Aurora-B Expression	pNPM1 <sup>Ser125</sup> Expression			
	-	+	++	+++
-	4	1	0	0
+	2	6	2	1
++	1	3	17	2
+++	0	4	7	37

**Note:** Aurora-B and p-NPM1<sup>Ser125</sup> expression in OS tissues were detected by IHC and analyzed using Spearman's rho tests (R=0.723).

## Aurora-B-Induced NPM1<sup>Ser125</sup> Phosphorylation Promotes OS Malignant Phenotypes in vitro

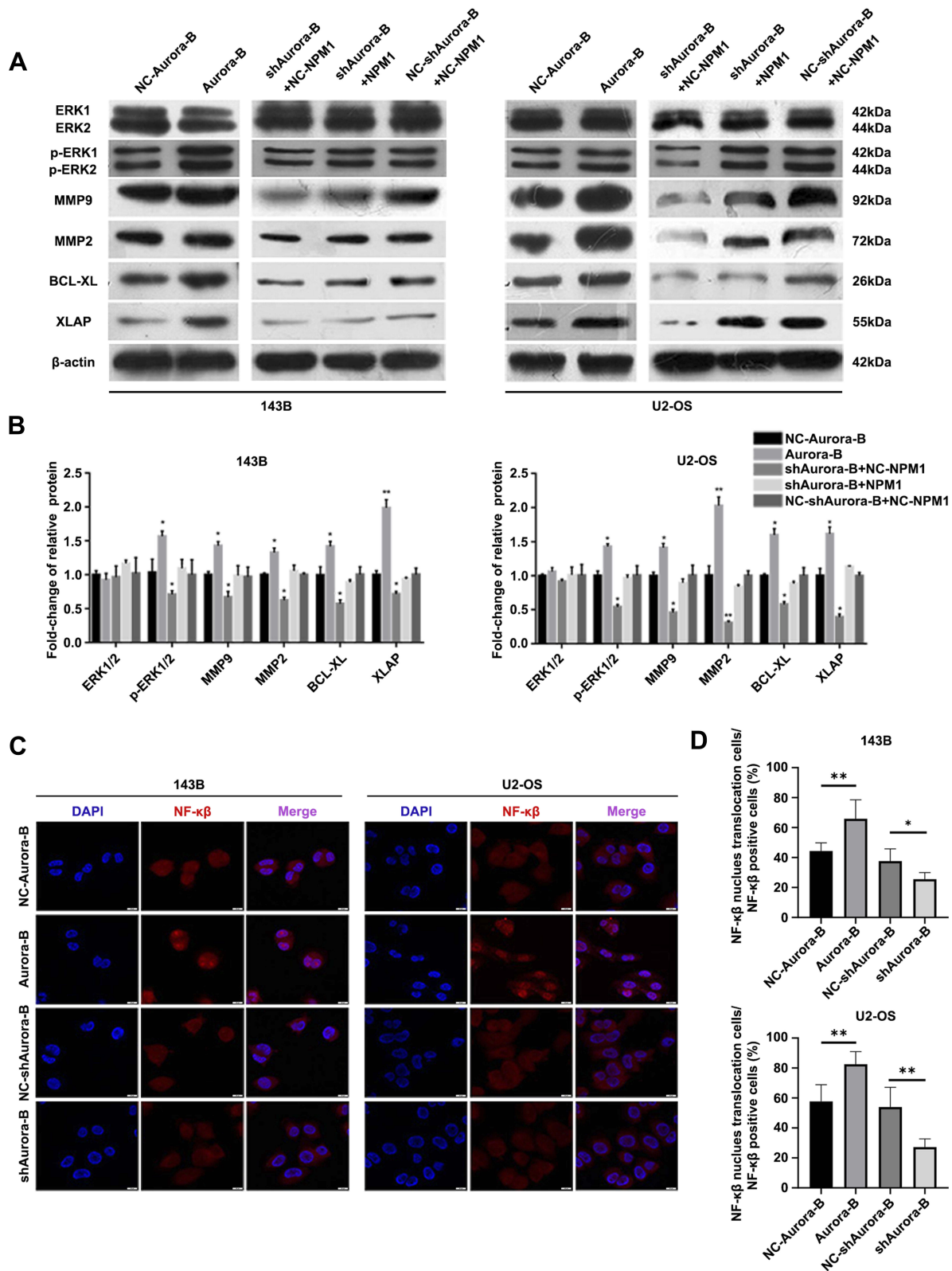
To investigate the role of Aurora-B-mediated NPM1 phosphorylation in OS progression. Wound healing, transwell invasion, CCK8 and TUNEL assays were performed to evaluate the malignant phenotypes induced by Aurora-B and NPM1 alterations. The results indicated that the metastasis and proliferation ability of OS cells significantly increased in the Aurora-B group, whilst the opposite phenotype was observed in Aurora-B silenced groups. In addition, the rates of apoptosis were significantly reduced in cells transfected with LV/Aurora-B. These effects could be reversed by the restoration of NPM1 (Figure 3A-D). These results indicate that Aurora-B promotes OS malignancy in-part through the phosphorylation of NPM1.

## Aurora-B Promotes OS Tumor Formation Through Phosphorylating NPM1

To confirm the role of Aurora-B-mediated NPM1 phosphorylation in OS initiation and metastasis, lentivirus-vectors were injected into tumor tissues. The results showed that tumor size and pulmonary nodules in nude mice intratumorally injected with LV/Aurora-B increased compared to the control group. The opposite phenotype was observed in nude mice injected with LV/shAurora-B (Figure 4A-D and Table 2). In addition, Western blot analysis indicated that the expression of Aurora-B and pNPM1<sup>Ser125</sup> were significantly higher in nude mice transfected with LV/Aurora-B compared to the control group. The expression of NPM1 remained almost unchanged in each group (Figure 4E and F). These results suggest that Aurora-B accelerates tumor initiation and metastases through the phosphorylation of NPM1 in vivo.

## Discussion

Aurora-B has been proved to play a tumor-promoting role in osteosarcoma.<sup>14</sup> Our previous studies indicated that Aurora-B expression was higher in OS and promoted OS malignant phenotypes via the activation of PI3K/Akt signaling.<sup>15,16</sup> However, the inhibition of PI3K/Akt alleviated the effects of Aurora-B on OS cell malignancy only partially, suggesting the existence of other mechanisms in Aurora-B-mediated OS. Herein, we explored the downstream effectors of Aurora-B and its potential molecular

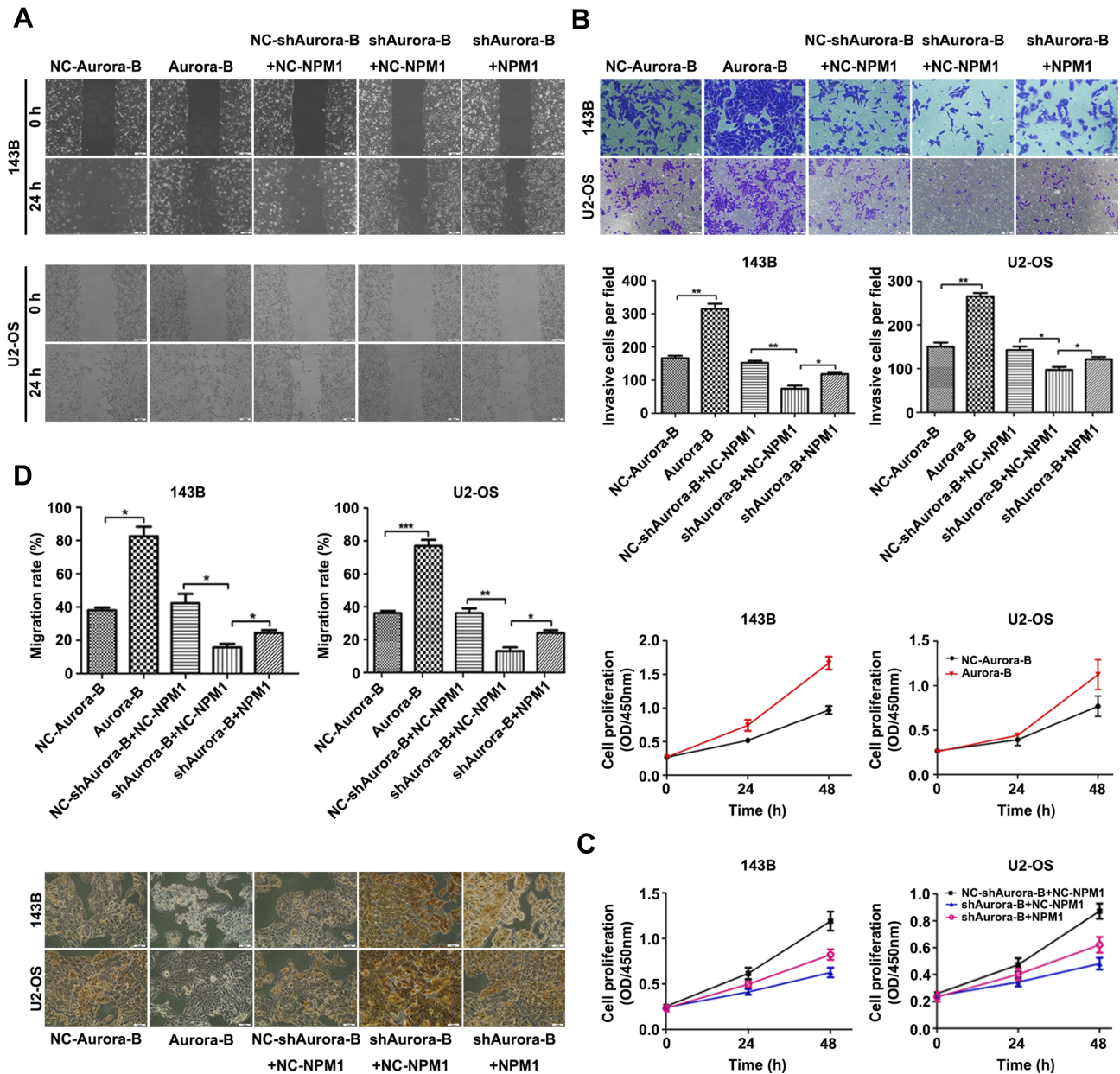


**Figure 2** Aurora-B activates ERK/NF-κβ signaling through the phosphorylation of NPM1<sup>Ser125</sup>. **(A)** Relative expression of ERK1/2, pERK1/2, MMP9, MMP2, BCL-XL and XIAP were examined by Western blotting in 143B and U2-OS cells. **(B)** Quantification of the Western blot assay. Values represent mean±SD (n = 3). \*p<0.05, \*\*p<0.01. **(C)** For immunofluorescence, NF-κβ nuclear translocation in U2-OS and 143B cells was assessed by confocal microscopy (x 400). **(D)** Quantification of cells with NF-κβ in the nucleus over total NF-κβ positive cells. \*p<0.05, \*\*p<0.01.

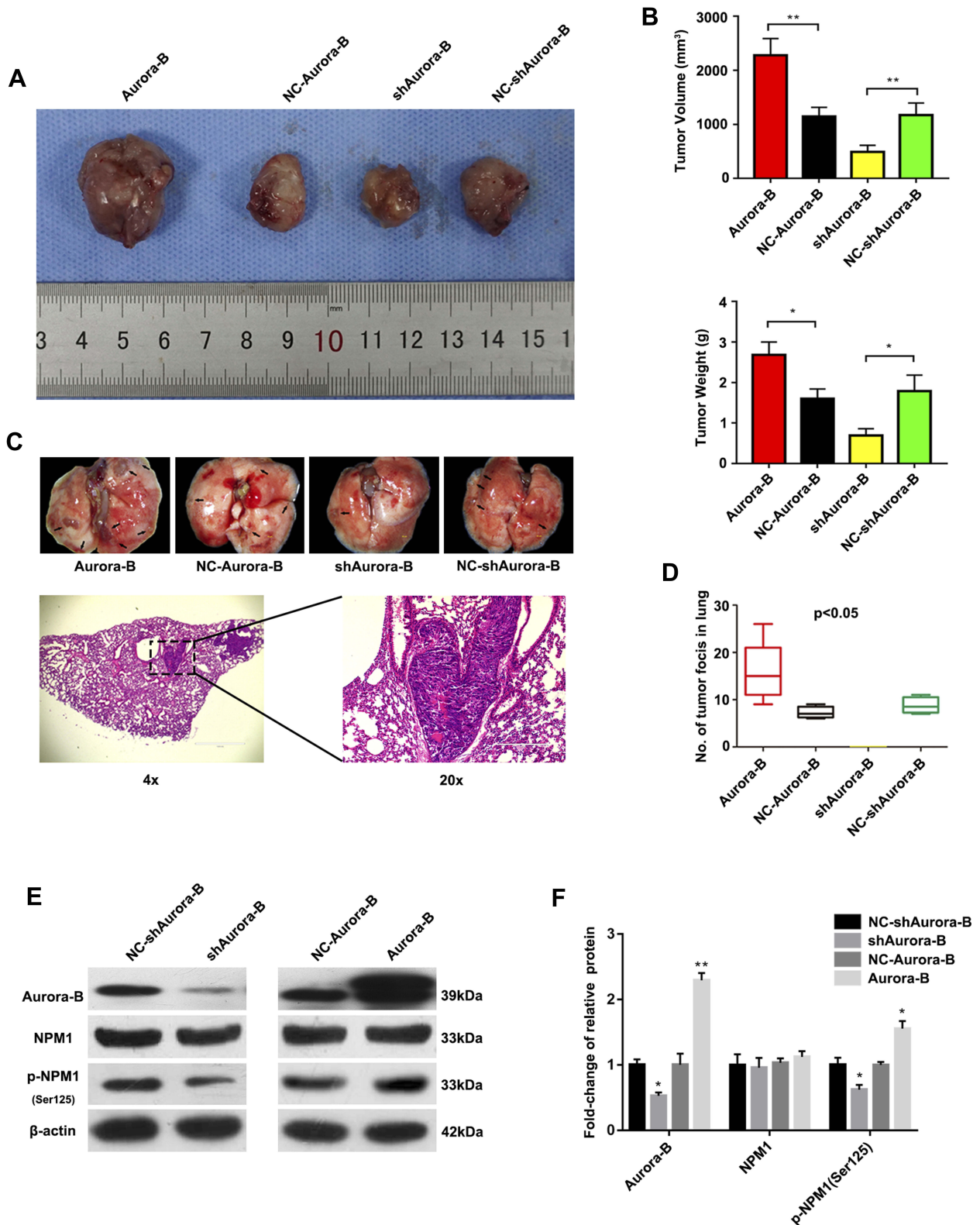
mechanisms in vitro and in vivo. Based on bioinformatics tools, we identified NPM1 as a substrate of Aurora-B, and hypothesized that Aurora-B interacts with NPM1, thus affecting the OS phenotype.

NPM1 is a ubiquitously expressed nucleocytoplasmic shuttling protein that plays an active role in ribosomal assembly, chromatin remodeling, DNA repair, replication, and transcription.<sup>17-19</sup> Accumulating evidence suggest that NPM1 is directly implicated in cancer progression. High

levels of NPM1 expression have been found in various malignant tumors and are associated with oncogenic progression and chemoresistance, including prostate, hematological tumors and high-grade serous ovarian adenocarcinoma.<sup>20-22</sup> Sawazaki et al reported that nucleophosmin is strongly associated with the recurrence and prognosis of urothelial carcinoma patients with radical nephroureterectomy.<sup>23</sup> Chen et al revealed that NPM1 inhibits apoptosis through its effects on Akt activity in breast cancer.<sup>24</sup> However, the upstream



**Figure 3** Aurora-B-induced NPM1<sup>Ser125</sup> phosphorylation promotes OS malignant phenotypes in vitro. (A) Cell migration distances at 0 h and 24 h (200×). \*p<0.05, \*\*p<0.01, \*\*\*p<0.001. (B) Representative images of transwell invasion assays. Chambers were crystal violet stained for 30 min. Cells in five random fields were imaged and counted \*p<0.05, \*\*p<0.01. Scale bar: 50 μm. (C) OD 450 was detected at 0 h, 24 h, and 48 h. Mean ± S.D. (D) Representative images of TUNEL assays. Degree of apoptosis was assessed by brown staining. Darker colors indicate higher levels of apoptosis.



**Figure 4** Aurora-B promotes OS tumor formation through phosphorylating NPM1. **(A)** Spontaneous metastasis xenograft models in nude mice. Mice were intratumorally injected with the indicated lentivirus vectors. Six weeks later, tumors were dissected and representative images were obtained. **(B)** Quantitative tumor volumes and weight,  $V = (\text{Length} \times \text{Width}^2) / 2$ . (mean  $\pm$  S.D.;  $n = 12$ . \* $p < 0.05$ , \*\* $p < 0.01$ ). **(C)** Representative images of collected lungs in Aurora-B, shAurora-B, NC-Aurora-B and NC-shAurora-B groups (upper panels). Black arrows indicate metastases ( $\times 7.5$ ). Representative H&E stained images in Aurora-B groups (lower panels), (Left:  $\times 100$ . Right:  $\times 400$ ). **(D)** Pulmonary metastatic foci in related groups were quantified (mean  $\pm$  S.D.  $p < 0.05$ ). **(E)** Western blot analysis of protein expression. **(F)** Quantification of the Western blot assay. Values represent mean  $\pm$  SD ( $n = 3$ ). \* $p < 0.05$ , \*\* $p < 0.01$ .



**Table 2** Number of Nude Mice with/Without Lung Metastasis

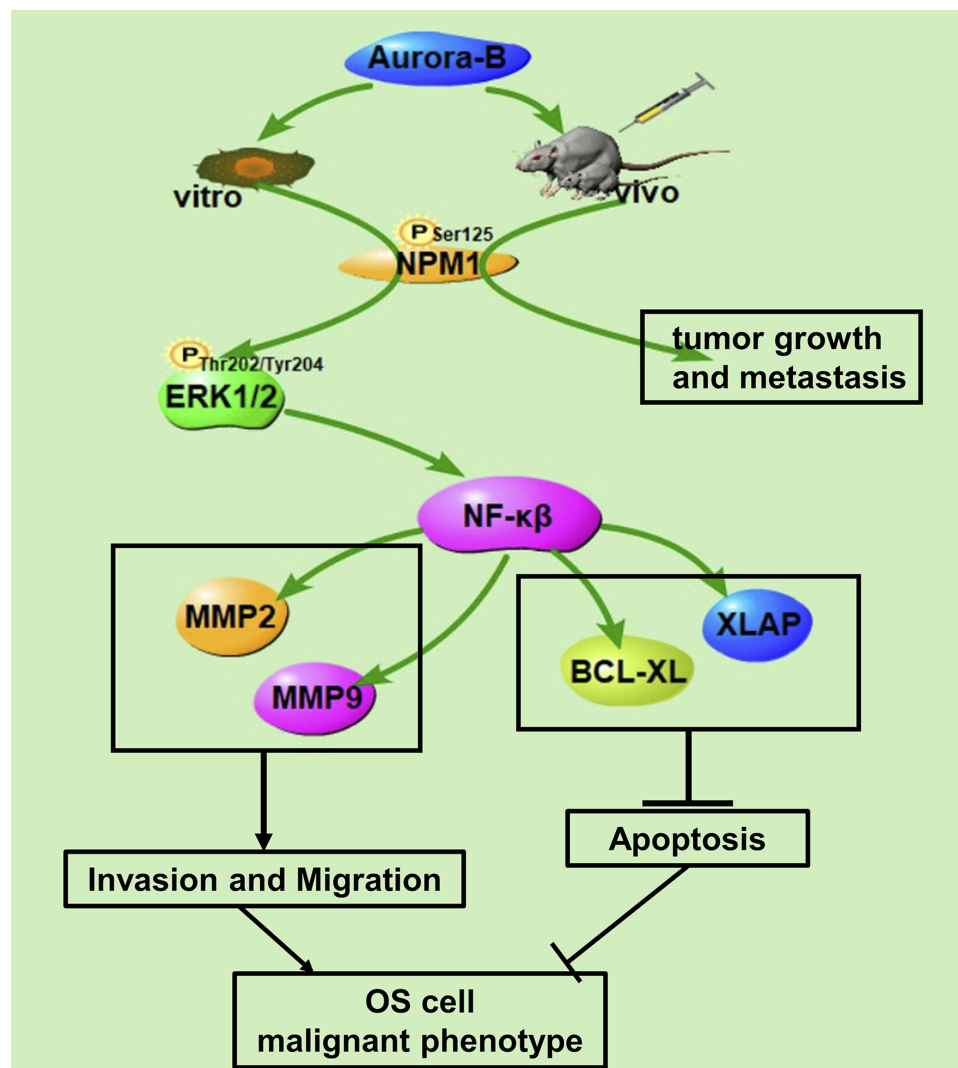
Groups (n=12 Mice/Group)	With Lung Metastasis	Without Lung Metastasis
Aurora-B	11	1
NC-Aurora-B	3	9
shAurora-B	1	11
NC-shAurora-B	4	8

**Note:** Metastasis foci in the lungs were stained by H&E and counted by two pathologists using stereomicroscopy.

molecular mechanisms through which NPM1 promotes cancer progression remain unclear. In this study, we investigated whether Aurora-B modulates NPM1 to promote OS metastasis. Bioinformatics predictions (the Molecular INTERaction Database; the Oncomine Database) revealed that NPM1 was overexpressed in osteogenic OS tissue and may interact

with Aurora-B. We found that Aurora-B expression was positively associated with phosphorylated NPM1 in OS tissue and cell lines. Western blot analysis also confirmed the effects of Aurora-B on the modulation of NPM1 phosphorylation. Although more assays should be performed to identify the interaction between Aurora-B and NPM1, our results were consistent with Shandilya et al who found that NPM1 can be phosphorylated by Aurora-B and participate in mitotic progression.<sup>25</sup> Further studies demonstrated that the inhibitory effects induced by Aurora-B silencing could be partially reversed by NPM1 overexpression. These results suggest that Aurora-B promotes OS cells malignancy in-part, by phosphorylating NPM1<sup>Ser125</sup>.

ERK/NF- $\kappa$ B signaling plays a critical role in malignant tumor proliferation, invasion and metastasis.<sup>26</sup> ERK1/2 are highly conserved, universal extracellular-regulated kinases



**Figure 5** Proposed scheme of molecular basis for effect of Aurora-B/NPM1/ERK/NF- $\kappa$ B axis in osteosarcoma according to our results.

that participate in a range of pathologies. The phosphorylation of ERK1/2 induces NF- $\kappa$ B activation in tumor cells.<sup>27</sup> Nuclear factor- $\kappa$ B (NF- $\kappa$ B) is a well-known transcription factor consisting of five components, NF- $\kappa$ B1 (p50), NF- $\kappa$ B2 (p52), RelA (p65), RelB and Rel.<sup>28</sup> NF- $\kappa$ B participates in cell cycle regulation, apoptosis, and cytokine secretion via initiating protein transcription. Aberrant NF- $\kappa$ B expression was observed in various malignant tumors, including gastric cancer,<sup>29</sup> breast cancer,<sup>30</sup> and prostate cancer.<sup>31</sup> NF- $\kappa$ B activation initiates the transcription of MMP-2 and MMP-9, the increased MMPs proteins enhance the invasiveness of tumor cells through the degradation of extracellular matrix (ECM).<sup>32,33</sup> Besides, NF- $\kappa$ B silencing promotes apoptosis in hematopoietic tumors via inhibiting Bcl-xL and XIAP.<sup>34</sup> Recent studies revealed that NPM1 silencing inhibits prostate cancer cell growth, migration and invasion via reducing the phosphorylation of ERK1/2.<sup>35</sup> We therefore hypothesized that Aurora-B-mediated NPM1 phosphorylation affects the OS phenotype via activating ERK/NF- $\kappa$ B signaling. Herein, we report that Aurora-B overexpression enhances pERK1/2, MMP9, MMP2, BCL-XL, XLAP expression and the translocation of NF- $\kappa$ B/p65. In addition, the inhibitory effects of Aurora-B on pERK1/2, MMP9, MMP2, BCL-XL and XLAP can be recovered by NPM1 overexpression, confirming our hypothesis.

## Conclusion

Our results show that Aurora-B promotes OS cell growth and metastasis via activating NPM1/ERK/NF- $\kappa$ B. Targeting the Aurora-B and NPM1/ERK/NF- $\kappa$ B axis may therefore represent a promising strategy for OS management (Figure 5).

## Abbreviations

OS, osteosarcoma; Aurora-B, aurora kinase B-like; NPM1, nucleophosmin 1; IHC, immunohistochemistry; IF, immunofluorescence; qRT-PCR, quantitative reverse transcriptase polymerase chain reaction; DMEM, Dulbecco's Modified Eagle Media; shRNA, short-hairpin RNA; TUNEL, TdT-mediated dUTP Nick-End Labeling.

## Author Contributions

JM L and ZL L contribute to the conception and design of the study; HH S, AF P and X W researched data; JM L and WZ C contribute to the analysis and interpretation of data. HH S and Y Z edited the manuscript; ZL L contributes to the final approval of the version to be submitted. All authors contributed to data analysis, drafting or revising

the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

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## Disclosure

The authors report no conflicts of interest in this work.

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