

Original Article

NOVA1 acts as an oncogene in osteosarcoma

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Abstract: Osteosarcoma is one of the most common bone tumors in young patients. NOVA1 (neuro-oncological ventral antigen 1) is a neuron-specific RNA binding-protein and belongs to the Nova family. Previous studies showed that NOVA1 played crucial roles in the development of several tumors. The objective of our study was to study the role of NOVA1 in the osteosarcoma. In our study, we showed that NOVA1 expression was upregulated in osteosarcoma cell lines and tissues. The expression of NOVA1 was upregulated in 22 (22/30; 73%) osteosarcoma cases compared to that in the adjacent tissues. Overexpression of NOVA1 promoted osteosarcoma cell viability, colony formation and invasion. Furthermore, knockdown of NOVA1 suppressed osteosarcoma cell viability, colony formation and invasion. These data suggested that NOVA1 acted as an oncogene in the development of osteosarcoma.

Keywords: Osteosarcoma, RNA binding-proteins, NOVA1, oncogene

Introduction

Osteosarcoma is one of the most common bone malignant tumors in young patients [1-4]. Despite the development of various treatment strategies which combine multiagent chemotherapy, surgery and radiotherapy, the five year survival rate of osteosarcoma patients is still unsatisfied [4-8]. About 80% of osteosarcoma patients will develop metastatic disease or local relapse after surgical treatment [9-12]. The detail molecular mechanisms underlying osteosarcoma progression and carcinogenesis remain uncovered [13-16]. Hence, it's urgent to find novel markers and develop new treatment strategies for osteosarcoma.

NOVA1 (neuro-oncological ventral antigen 1) is a neuron-specific RNA binding-protein and one member of Nova family [17-22]. It was first identified in a neurologic disease characterized by failure of the motor suppression [23]. NOVA1 is proved to play significant roles in the development of motor system and motoneurons survival [23]. It can regulate the alternative process of various mRNAs that is crucial for synaptic activity [24]. Recent evidences have showed that NOVA1 plays crucial roles in the development of several tumors [18, 21, 25-27]. For example, Kim et al [20]. demonstrated that NOVA1 expression was inhibited in the micro-environment of gastric cancer, and attenuated

NOVA1 expression in the gastric cancer cells was correlated with cancer progression and poor prognosis. Shen et al [21]. showed that miR-339 was upregulated in gastric cancer tissues and ectopic expression of miR-339 inhibited gastric cancer cell invasion, migration and viability through targeting NOVA1. Ectopic expression of NOVA1 in the miR-339-overexpressing gastric cancer cells impaired the inhibitory function of miR-339. However, the role of NOVA1 in osteosarcoma is still unknown.

In this study, we demonstrated that NOVA1 expression was upregulated in osteosarcoma cell lines and tissues. Overexpression of NOVA1 promoted osteosarcoma cell viability, colony formation and invasion. Furthermore, Knockdown of NOVA1 suppressed osteosarcoma cell viability, colony formation and invasion. These data suggested that NOVA1 acted as an oncogene in the development of osteosarcoma.

Materials and methods

Specimens and cell lines cultured and transfection

Surgically excised tissues from 35 cases with osteosarcoma tissues and adjacent normal tissues were collected in the Heilongjiang provincial hospital. All the samples were collected after obtaining the written informed consent

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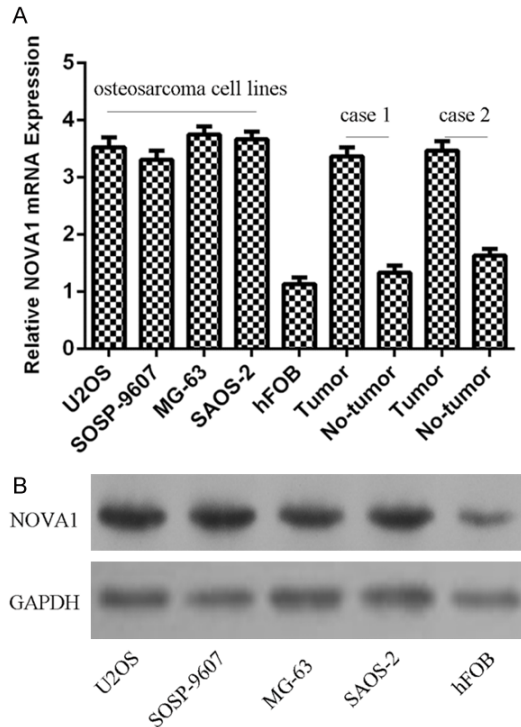


Figure 1. NOVA1 expression was upregulated in osteosarcoma cell lines. A. The mRNA expression of NOVA1 in the osteosarcoma tissues and cell lines was determined by using qRT-PCR. B. The protein expression of NOVA1 was determined in the osteosarcoma cell lines (MG-63, U2OS, SOSP-9607 and SAOS-2) and hFOB by using Western blot.

following to the protocol was approved by the Institutional Review Board of the Heilongjiang provincial hospital. The tissues were snap-frozen in the liquid nitrogen and stored. The following osteosarcoma cell lines were used in our study: MG-63, U2OS, SOSP-9607 and SAOS-2 and osteoblast cell line (hFOB). These cells were collected from the Cell bank Center of the Institute of the Chinese Academy of Medical Sciences (Beijing, China) and were cultured in the DMEM (Dulbecco's modified Eagle's medium; Gibco; Invitrogen) supplemented with FBS (fetal bovine serum; GIBCO, USA), streptomycin, and penicillin. NOVA1 vector and the control vector were purchased from the GenePharma (Shanghai, China). The cells were transfected with vector using the Lipofectamine 2000 (Invitrogen, USA) according to the instruction.

Real-time quantitative PCR (qPCR)

Total RNA from cells or tissues was extracted by using the Trizol reagent (Invitrogen, CA, USA) according to the instruction. qRT-PCR was performed to detect the mRNA expression of NO-

VA1 and GAPDH on the ABI 7500 System (ABI) using the SYBR Green Mix (Takara). Ct ($2^{-\Delta\Delta Ct}$) method was used to measure the relative expression of each group. The following sequences: NOVA1, (forward: GGGTTCCTAGACCTGGAC; reverse: CGCTCAGTAGTACCTGGGTAA); GAPDH, (forward: TGCACCACTGCTTAGC; reverse: GGCATGGACTGTGGTCATGAG).

Western blotting

Cells or tissues were lysed with lysis buffer and the total proteins concentration was measured by using the BCA Protein Assay. Total protein was separated on the 12% SDS-PAGE and transferred to PVDF membranes (Bio-Rad). The membrane was blocked with FBS for 1 hour and then incubated the primary antibodies (NOVA1, GAPDH, Sigma, USA) overnight and band was visualized by the chemiluminescence.

Viability, invasion, colony formation assay

Cell viability was detected by the MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide). Twenty μ l MTT solution was added to the cell culture medium for about 4 hours incubation. Then, the absorbance of each well was measured at 450 nm. For cell invasion assay, Matrigel coated transwell chambers (Millipore, USA) was used. Cells were cultured in the serum-free medium on the upper chamber and medium containing FBS as the chemo-attractant was put to the bottom well. The invasive cell was stained with the crystal violet after 48 hours. For colony formation assay, cells were cultured on the 6-well plate and kept in the DMEM medium containing 10% FBS for 2 weeks. Colony was fixed and stained and the number of colony was counted.

Statistical analysis

Statistical analysis was measured by the SPSS version 17.0 (Chicago, IL, USA). Data are presented as the mean \pm SD (standard deviation). $P < 0.05$ is considered to be statistically significant. The difference between groups was determined using Student's t-test or one-way ANOVA.

Results

NOVA1 expression was upregulated in osteosarcoma cell lines

We firstly detected the NOVA1 expression level in osteosarcoma cell lines. Our result showed

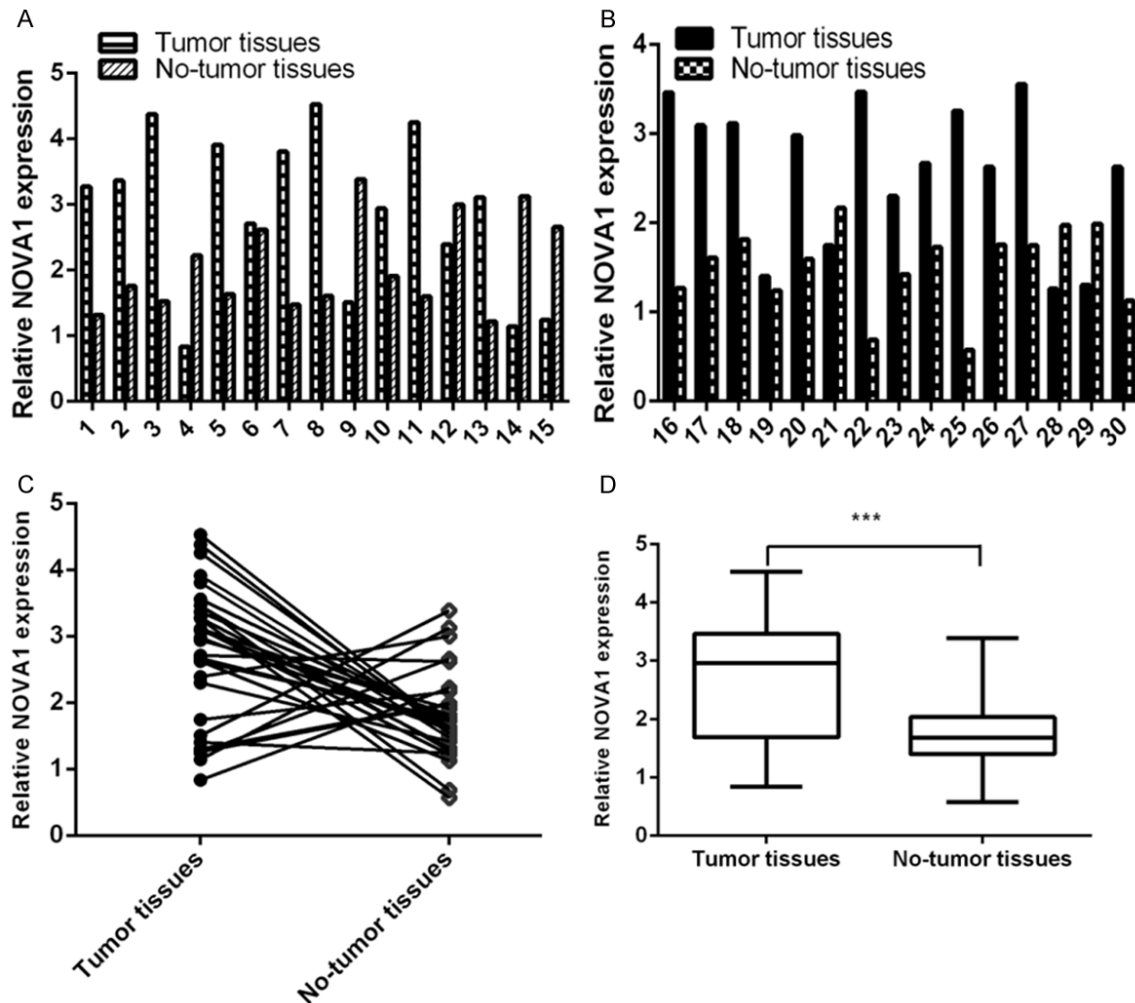


Figure 2. NOVA1 expression was upregulated in osteosarcoma tissues. A. The expression of NOVA1 in the osteosarcoma tissues was detected by using qRT-PCR. B. The expression of NOVA1 in the osteosarcoma tissues was detected by using qRT-PCR. C. The expression of NOVA1 was upregulated in 22 cases (22/30; 73%) of osteosarcoma cases compared to that in adjacent tissues. D. The expression level of NOVA1 in osteosarcoma tissues was higher than that in the adjacent tissues. ***P<0.001.

that the expression level of NOVA1 was upregulated in osteosarcoma cell lines (MG-63, U2OS, SOSP-9607 and SAOS-2) and tissue compared to that in the osteoblast cell line (hFOB) (**Figure 1A**). Moreover, the NOVA1 protein expression was also upregulated in the osteosarcoma cell lines (MG-63, U2OS, SOSP-9607 and SAOS-2) compared to that in the hFOB (**Figure 1B**).

NOVA1 expression was upregulated in osteosarcoma tissues

We next measured NOVA1 expression in 30 osteosarcoma cases. The expression of NOVA1 was shown in the **Figure 2A** and **2B**. The expression of NOVA1 was upregulated in 22 cases

(22/30; 73%) of osteosarcoma cases compared to that in adjacent tissues (**Figure 2C**). The expression level of NOVA1 in osteosarcoma tissues was higher than that in the adjacent tissues (**Figure 2D**).

Overexpression of NOVA1 promoted osteosarcoma cell viability

To study the biological function of NOVA1 in osteosarcoma cell, MG-63 cell was transfected with NOVA1 vector and control vector. The mRNA and protein expression of NOVA1 was upregulated in the MG-63 cell after treated with NOVA1 vector (**Figure 3A** and **3B**). Ectopic expression of NOVA1 promoted the MG-63 cell

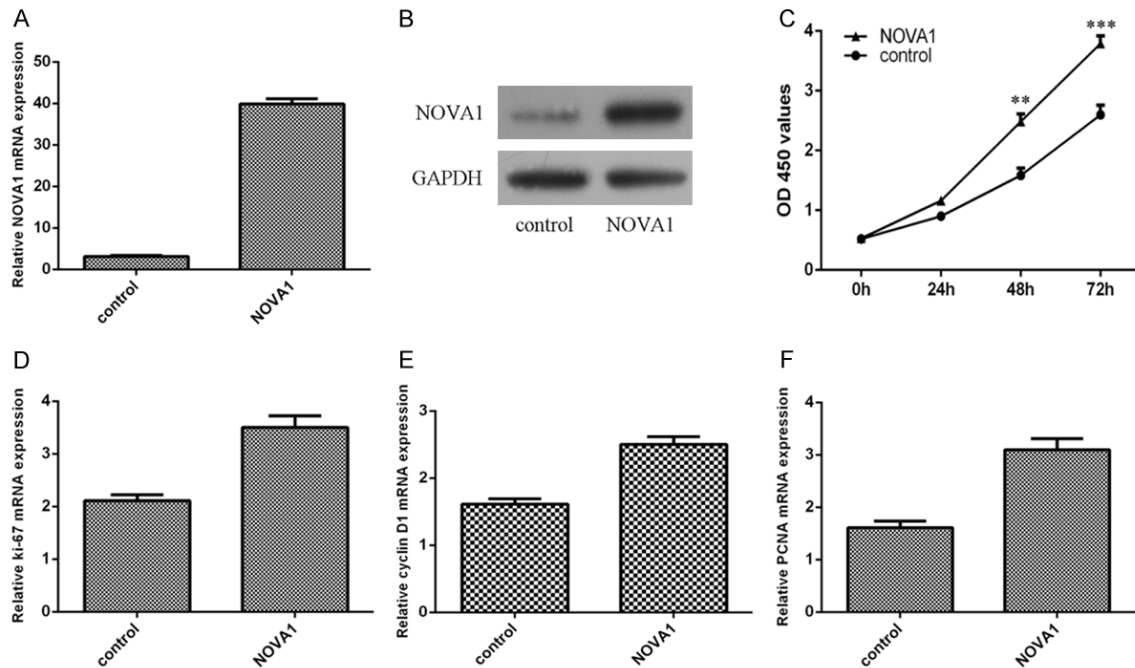


Figure 3. Overexpression of NOVA1 promoted osteosarcoma cell viability. A. The mRNA expression of NOVA1 in the MG-63 cell after treated with NOVA1 vector was measured by qRT-PCR. B. The protein expression of NOVA1 in the MG-63 cell after treated with NOVA1 vector was measured by Western blot. C. Ectopic expression of NOVA1 promoted the MG-63 cell viability. D. The expression of ki-67 was determined by qRT-PCR. E. The expression of cyclin D1 was determined by qRT-PCR. F. The expression of PCNA was determined by qRT-PCR. ** $P < 0.01$ and *** $P < 0.001$.

viability (**Figure 3C**). Moreover, overexpression of NOVA1 increased the expression of ki-67, cyclin D1 and PCNA (**Figure 3D-F**).

Overexpression of NOVA1 promoted osteosarcoma cell colony formation and invasion

Ectopic expression of NOVA1 increased osteosarcoma cell colony formation (**Figure 4A**). In addition, we showed that overexpression of NOVA1 promoted the MG-63 cell invasion by using invasion assay (**Figure 4B**).

Knockdown of NOVA1 suppressed osteosarcoma cell viability

To further detect the biological function of NOVA1 in osteosarcoma cell, MG-63 cell was transfected with si-NOVA1 vector and control vector. The mRNA and protein expression of NOVA1 was downregulated in the MG-63 cell after treated with si-NOVA1 vector (**Figure 5A** and **5B**). Inhibition expression of NOVA1 suppressed MG-63 cell viability (**Figure 5C**). Moreover, knockdown of NOVA1 decreased the expression of ki-67, cyclin D1 and PCNA (**Figure 5D-F**).

Knockdown of NOVA1 suppressed osteosarcoma cell colony formation and invasion

Knockdown of NOVA1 decreased osteosarcoma cell colony formation (**Figure 6A**). In addition, we showed that inhibition expression of NOVA1 suppressed the MG-63 cell invasion through using invasion assay (**Figure 6B**).

Discussion

In this study, we demonstrated that NOVA1 expression was upregulated in osteosarcoma cell lines and tissues. The expression of NOVA1 was upregulated in 22 cases (22/30; 73%) of osteosarcoma cases compared to that in adjacent tissues. The expression level of NOVA1 in osteosarcoma tissues was higher than that in the adjacent tissues. Overexpression of NOVA1 promoted osteosarcoma cell viability, colony formation and invasion. Furthermore, Knockdown of NOVA1 suppressed the osteosarcoma cell viability, colony formation and invasion. These data suggested that NOVA1 acted as an oncogene in the development of osteosarcoma.

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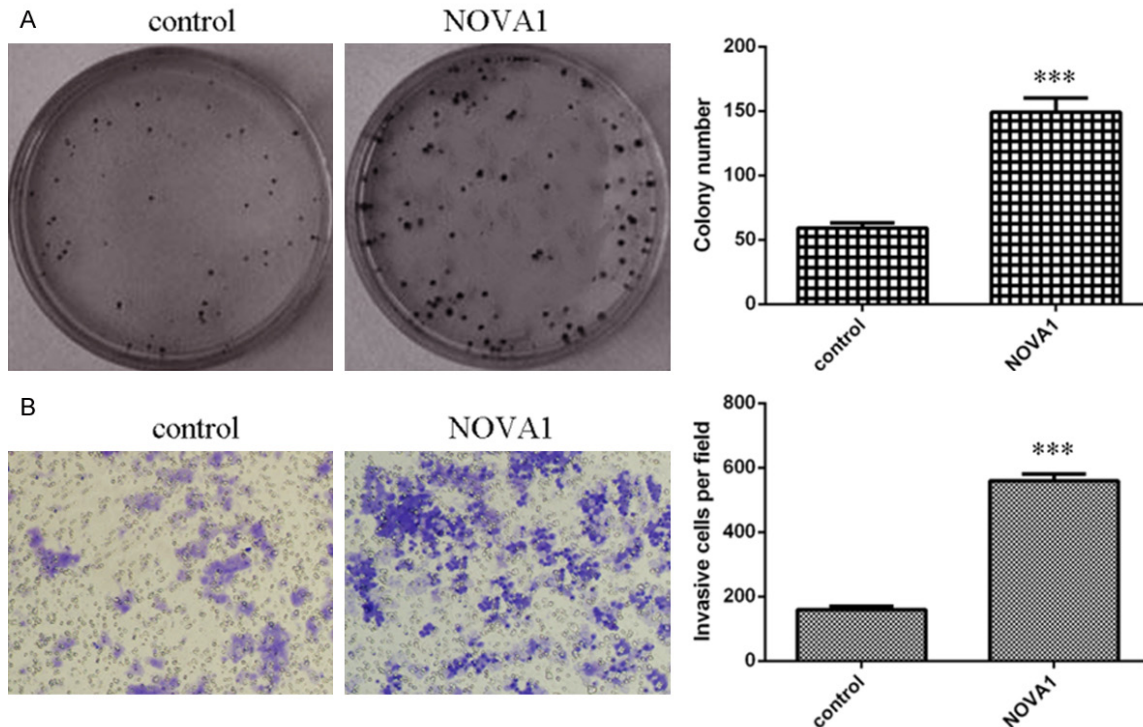


Figure 4. Overexpression of NOVA1 promoted osteosarcoma cell colony formation and invasion. A. Ectopic expression of NOVA1 increased osteosarcoma cell colony formation. The relative colony number was shown in the right. B. Overexpression of NOVA1 promoted the MG-63 cell invasion by using invasion assay. The relative invasive cell was shown in the right. *** $P < 0.001$.

NOVA1 is one member of the Nova family, which was firstly identified as the target in autoimmune neurologic diseases characterized through the failure of motor suppression [23]. NOVA1 was essential for development of the mammalian motor system and survival of motoneurons through regulating the alternative process of various mRNAs that is crucial for synaptic activity [28-30]. Recent studies have demonstrated that NOVA1 plays important roles in the development of various tumors [17, 19, 26, 31, 32]. For example, Kim et al [20]. showed that NOVA1 inhibition was found in the microenvironment of gastric cancer and attenuated NOVA1 expression was correlated with gastric cancer progression and poor prognosis. Yoon et al [18]. showed that NOVA1 was enriched in stromal spindle cells and T lymphocytes, while NOVA1 expression was frequently downregulated in those types of cells and gastric cancer tissues. Shen et al [21]. demonstrated that miR-339 overexpression suppressed gastric cancer cells migration, viability, tumorigenicity and invasion through inhibiting NOVA1. Zhang et al [26]. demonstrated that higher NO-

VA1 expression was correlated with increased recurrence rate and poor survival of hepatocellular carcinoma. Overexpression of NOVA1 promoted hepatocellular carcinoma cell invasion, migration and viability. However, the role and biology function of NOVA1 are still unknown. In this study, we firstly detected the expression of NOVA1 in osteosarcoma cell lines (MG-63, U2OS, SOSP-9607 and SAOS-2) and tissue compared to that in the osteoblast cell line (hFOB). Furthermore, we measured the NOVA1 expression in the 30 osteosarcoma cases. The expression of NOVA1 was upregulated in 22 cases (22/30; 73%) of osteosarcoma cases compared to that in adjacent tissues. The expression level of NOVA1 in osteosarcoma tissues was higher than that in the adjacent tissues.

We next investigated the function role of NOVA1 in osteosarcoma cell. Ectopic expression of NOVA1 promoted osteosarcoma cell line MG-63 cell viability. In line with this, we show-

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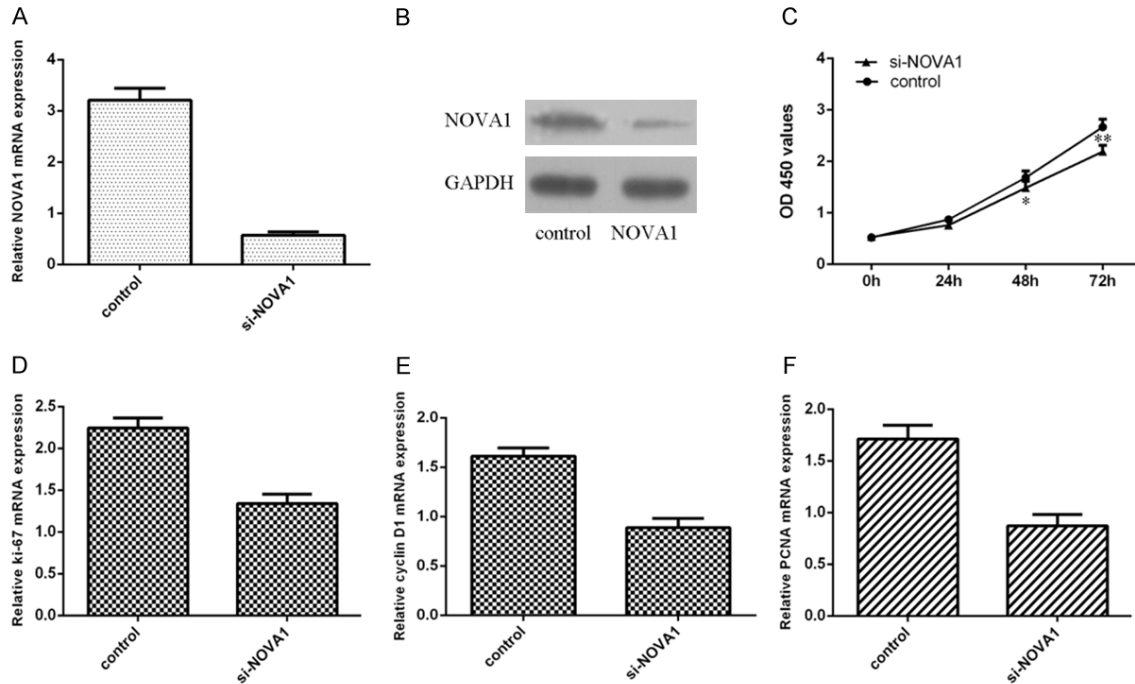


Figure 5. Knockdown of NOVA1 suppressed osteosarcoma cell viability. A. The mRNA expression of NOVA1 in the MG-63 cell after treated with si-NOVA1 was measured by qRT-PCR. B. The protein expression of NOVA1 in the MG-63 cell after treated with si-NOVA1 vector was measured by Western blot. C. Inhibition expression of NOVA1 promoted the MG-63 cell viability. D. The expression of ki-67 was determined by qRT-PCR. E. The expression of cyclin D1 was determined by qRT-PCR. F. The expression of PCNA was determined by qRT-PCR. * $P < 0.05$ and ** $P < 0.01$.

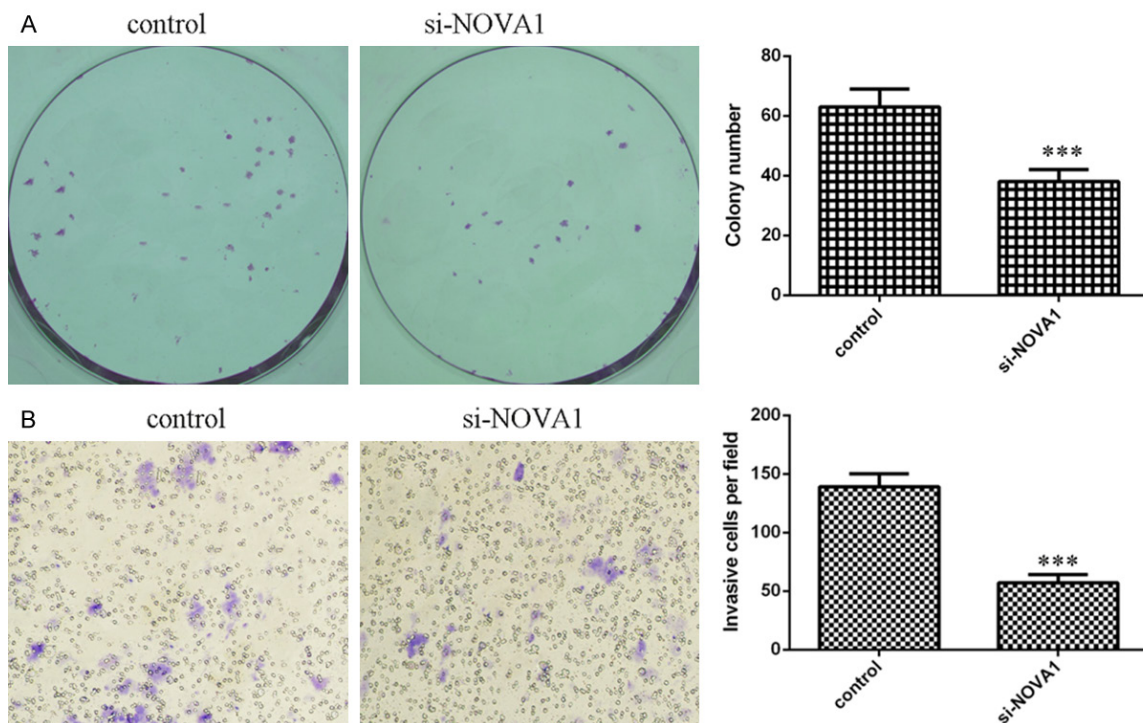


Figure 6. Knockdown of NOVA1 suppressed osteosarcoma cell colony formation and invasion. A. Knockdown of NOVA1 decreased osteosarcoma cell colony formation. The relative colony number was shown in the right. B. Inhibition expression of NOVA1 suppressed the MG-63 cell invasion through using invasion assay. The relative invasive cell was shown in the right. *** $P < 0.001$.

ed that overexpression of NOVA1 increased the expression of ki-67, cyclin D1 and PCNA. Furthermore, we demonstrated that overexpression of NOVA1 promoted the MG-63 cell colony formation and invasion. In addition, we showed that knockdown of NOVA1 suppressed osteosarcoma cell line MG-63 cell viability. We also showed that inhibition of NOVA1 decreased the expression of ki-67, cyclin D1 and PCNA. In addition, we demonstrated that suppression of NOVA1 inhibited MG-63 cell colony formation and invasion. These results suggested that NOVA1 played as an oncogene in the development of osteosarcoma.

In conclusion, NOVA1 acted as an oncogene in the initiation and development of osteosarcoma. Given that overexpression of NOVA1 promoted osteosarcoma cell viability and invasion in the osteosarcoma, NOVA1 may be a potential therapeutic target for osteosarcoma.

Disclosure of conflict of interest

None.

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References

- [1] Wang Y, Jia LS, Yuan W, Wu Z, Wang HB, Xu T, Sun JC, Cheng KF and Shi JG. Low miR-34a and miR-192 are associated with unfavorable prognosis in patients suffering from osteosarcoma. *Am J Transl Res* 2015; 7: 111-119.
- [2] Delebinski CI, Georgi S, Kleinsimon S, Twardziok M, Kopp B, Melzig MF and Seifert G. Analysis of proliferation and apoptotic induction by 20 steroid glycosides in 143B osteosarcoma cells in vitro. *Cell Prolif* 2015; 48: 600-610.
- [3] Tang J, Shen L, Yang Q and Zhang C. Overexpression of metadherin mediates metastasis of osteosarcoma by regulating epithelial-mesenchymal transition. *Cell Prolif* 2014; 47: 427-434.
- [4] Novello C, Pazzaglia L, Conti A, Quattrini I, Polino S, Perego P, Picci P and Benassi MS. p53-dependent activation of microRNA-34a in response to etoposide-induced DNA damage in osteosarcoma cell lines not impaired by dominant negative p53 expression. *PLoS One* 2014; 9: e114757.
- [5] Liu Y, Zhu ST, Wang X, Deng J, Li WH, Zhang P and Liu BS. MiR-100 inhibits osteosarcoma cell proliferation, migration, and invasion and enhances chemosensitivity by targeting IGFIR. *Technol Cancer Res Treat* 2016; 15: NP40-48.
- [6] Zhu Z, Tang J, Wang J, Duan G, Zhou L and Zhou X. MiR-138 acts as a tumor suppressor by targeting EZH2 and enhances cisplatin-induced apoptosis in osteosarcoma cells. *PLoS One* 2016; 11: e0150026.
- [7] Dai N, Qing Y, Cun Y, Zhong Z, Li C, Zhang S, Shan J, Yang X, Dai X, Cheng Y, Xiao H, Xu C, Li M and Wang D. miR-513a-5p regulates radiosensitivity of osteosarcoma by targeting human apurinic/apyrimidinic endonuclease. *Oncotarget* 2016; [Epub ahead of print].
- [8] Mori F, Sacconi A, Canu V, Ganci F, Novello M, Anelli V, Covello R, Ferraresi V, Muti P, Biagini R, Blandino G and Strano S. miR-181c associates with tumor relapse of high grade osteosarcoma. *Oncotarget* 2015; 6: 13946-13961.
- [9] Ge L, Zheng B, Li M, Niu L and Li Z. MicroRNA-497 suppresses osteosarcoma tumor growth in vitro and in vivo. *Oncol Lett* 2016; 11: 2207-2212.
- [10] Geng S, Gu L, Ju F, Zhang H, Wang Y, Tang H, Bi Z and Yang C. MicroRNA-224 promotes the sensitivity of osteosarcoma cells to cisplatin by targeting Rac1. *J Cell Mol Med* 2016; 20: 1611-1619.
- [11] Shen L, Chen XD and Zhang YH. MicroRNA-128 promotes proliferation in osteosarcoma cells by downregulating PTEN. *Tumour Biol* 2014; 35: 2069-2074.
- [12] Sun XH, Geng XL, Zhang J and Zhang C. miRNA-646 suppresses osteosarcoma cell metastasis by downregulating fibroblast growth factor 2 (FGF2). *Tumour Biol* 2015; 36: 2127-2134.
- [13] Wang L, Shao J, Zhang X, Xu M and Zhao J. microRNA-377 suppresses the proliferation of human osteosarcoma MG-63 cells by targeting CDK6. *Tumour Biol* 2015; 36: 3911-7.
- [14] Salah Z, Arafeh R, Maximov V, Galasso M, Khawaled S, Abou-Sharieha S, Volinia S, Jones KB, Croce CM and Aqeilan RI. miR-27a and miR-27a* contribute to metastatic properties of osteosarcoma cells. *Oncotarget* 2015; 6: 4920-4935.
- [15] Han K, Chen X, Bian N, Ma B, Yang T, Cai C, Fan Q, Zhou Y and Zhao TB. MicroRNA profiling identifies MiR-195 suppresses osteosarcoma cell metastasis by targeting CCND1. *Oncotarget* 2015; 6: 8875-8889.
- [16] Sarver AL, Thayanithy V, Scott MC, Cleton-Janssen AM, Hogendoorn PC, Modiano JF and Subramanian S. MicroRNAs at the human 14q32 locus have prognostic significance in osteosarcoma. *Orphanet J Rare Dis* 2013; 8: 7.
- [17] Villate O, Turatsinze JV, Mascali LG, Grieco FA, Nogueira TC, Cunha DA, Nardelli TR, Sammeth

- M, Salunkhe VA, Esguerra JL, Eliasson L, Marselli L, Marchetti P and Eizirik DL. Nova1 is a master regulator of alternative splicing in pancreatic beta cells. *Nucleic Acids Res* 2014; 42: 11818-11830.
- [18] Yoon SO, Kim EK, Lee M, Jung WY, Lee H, Kang Y, Jang YJ, Hong SW, Choi SH and Yang WI. NOVA1 inhibition by miR-146b-5p in the remnant tissue microenvironment defines occult residual disease after gastric cancer removal. *Oncotarget* 2016; 7: 2475-2495.
- [19] Zhi F, Wang Q, Deng D, Shao N, Wang R, Xue L, Wang S, Xia X and Yang Y. MiR-181b-5p down-regulates NOVA1 to suppress proliferation, migration and invasion and promote apoptosis in astrocytoma. *PLoS One* 2014; 9: e109124.
- [20] Kim EK, Yoon SO, Jung WY, Lee H, Kang Y, Jang YJ, Hong SW, Choi SH and Yang WI. Implications of NOVA1 suppression within the microenvironment of gastric cancer: association with immune cell dysregulation. *Gastric Cancer* 2017; 20: 438-447.
- [21] Shen B, Zhang Y, Yu S, Yuan Y, Zhong Y, Lu J and Feng J. MicroRNA-339, an epigenetic modulating target is involved in human gastric carcinogenesis through targeting NOVA1. *FEBS Lett* 2015; 589: 3205-3211.
- [22] Storchel PH, Thummler J, Siegel G, Aksoy-Aksel A, Zampa F, Sumer S and Schratt G. A large-scale functional screen identifies Nova1 and Ncoa3 as regulators of neuronal miRNA function. *EMBO J* 2015; 34: 2237-2254.
- [23] Ueki K, Ramaswamy S, Billings SJ, Mohrenweiser HW and Louis DN. ANOVA, a putative astrocytic RNA-binding protein gene that maps to chromosome 19q13.3. *Neurogenetics* 1997; 1: 31-36.
- [24] Ratti A, Fallini C, Colombrita C, Pascale A, Laforenza U, Quattrone A and Silani V. Post-transcriptional regulation of neuro-oncological ventral antigen 1 by the neuronal RNA-binding proteins ELAV. *J Biol Chem* 2008; 283: 7531-7541.
- [25] Kim EK, Yoon SO, Kim SH, Yang WI, Cho YA and Kim SJ. Upregulated neuro-oncological ventral antigen 1 (NOVA1) expression is specific to mature and immature T- and NK-cell lymphomas. *J Pathol Transl Med* 2016; 50: 104-112.
- [26] Zhang YA, Zhu JM, Yin J, Tang WQ, Guo YM, Shen XZ and Liu TT. High expression of neuro-oncological ventral antigen 1 correlates with poor prognosis in hepatocellular carcinoma. *PLoS One* 2014; 9: e90955.
- [27] Pitkanen E, Cajuso T, Katainen R, Kaasinen E, Valimaki N, Palin K, Taipale J, Aaltonen LA and Kilpivaara O. Frequent L1 retrotranspositions originating from TTC28 in colorectal cancer. *Oncotarget* 2014; 5: 853-859.
- [28] Jelen N, Ule J, Zivin M and Darnell RB. Evolution of Nova-dependent splicing regulation in the brain. *PLoS Genet* 2007; 3: 1838-1847.
- [29] Alkelai A, Greenbaum L, Rigbi A, Kanyas K and Lerer B. Genome-wide association study of antipsychotic-induced parkinsonism severity among schizophrenia patients. *Psychopharmacology (Berl)* 2009; 206: 491-499.
- [30] Iourov IY, Vorsanova SG, Liehr T, Kolotii AD and Yurov YB. Increased chromosome instability dramatically disrupts neural genome integrity and mediates cerebellar degeneration in the ataxia-telangiectasia brain. *Hum Mol Genet* 2009; 18: 2656-2669.
- [31] Amer M, Elhefnawi M, El-Ahwany E, Awad AF, Gawad NA, Zada S and Tawab FM. Hsa-miR-195 targets PCMT1 in hepatocellular carcinoma that increases tumor life span. *Tumour Biol* 2014; 35: 11301-11309.
- [32] Gimenez M, Marie SK, Oba-Shinjo S, Uno M, Izumi C, Oliveira JB and Rosa JC. Quantitative proteomic analysis shows differentially expressed HSPB1 in glioblastoma as a discriminating short from long survival factor and NOVA1 as a differentiation factor between low-grade astrocytoma and oligodendroglioma. *BMC Cancer* 2015; 15: 481.