

Effect of Histone Deacetylase Inhibitor to Osteosarcoma through Specificity Protein 1/ Phospholipase D1 Pathway

DAWEI. CAI¹, TANGBO. YUAN¹, ZHEN JIN AND JIAN QIN*¹

Department of Orthopaedics, The Second Affiliated Hospital of Nanjing Medical University, Nanjing, Jiangsu 210011,

¹Department of Orthopaedics, Sir Run Run Hospital, Nanjing Medical University, Nanjing, Jiangsu 211000, P.R. China

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To study the effect of histone deacetylase inhibitor to osteosarcoma by targeting specificity protein 1 and regulating Phospholipase D1. Human osteosarcoma cell line MG63 cells were cultured in Dulbecco's Modified Eagle's medium containing penicillin (100 U/ml) and 10 % fetal bovine serum at 37°. siRNA-specificity protein 1 and psiCHECK-Phospholipase-WT plasmid were constructed and transfected into MG63 cells. Dual luciferase reporter gene assay were conducted to verified the combination site of specificity protein 1 and Phospholipase. MG63 cells were treated with tumor-specific antigen (5 g/ml) or PBS (negative control) for 48 h at 37°. Western blot was used to test the phospholipase D1, nuclear specificity protein 1 and cytoplasmic specificity protein 1 in MG63 cells. Cell Counting Kit-8 and clone formation assay were conducted to analyze the difference in cell viability and proliferation of tumor-specific antigen -treated MG63 cells and tumor-specific antigen free cells. Western blot result showed that nuclear and cytoplasmic specificity protein 1 in tumor-specific antigen treated MG63 cells both decreased dramatically compared to tumor-specific antigen free cells ($p < 0.05$). Phospholipase in tumor-specific antigen treated MG63 cells also reduced ($p < 0.05$). Fluorescence analysis showed that the ratio of Firefly luciferase to Renilla luciferase in MG63 cells transfected with psiCHECK-Phospholipase-WT plasmid and si-specificity protein 1 was significantly lower than that of si-NC ($p < 0.05$), while the ratio of firefly luciferase to Renilla luciferase in MG63 cells transfected with psiCHECK-Phospholipase-WT plasmid and si-specificity protein 1 had no significant difference with si-NC control ($p < 0.05$). After tumor-specific antigen treatment for 24 h, the proliferation rate of MG63 cells was ($65.20 \% \pm 2.31 \%$), which was significantly lower than that of PBS control group ($96.63 \% \pm 1.14 \%$) ($p < 0.05$) by CCK-8 method. Cell formation assay showed that the amount of MG63 cells treated with tumor-specific antigen decreased significantly ($p < 0.05$). The gene expression of specificity protein 1 and Phospholipase in tumor-specific antigen -treated MG63 cells decreased significantly compared with tumor-specific antigen free cells ($p < 0.05$). The expression of survivin in tumor-specific antigen -treated MG63 cells were significantly lower than that in tumor-specific antigen free cells ($p < 0.05$). The gene expression levels of p21 in tumor-specific antigen -treated MG63 cells were significantly higher than that in tumor-specific antigen free cells ($p < 0.05$). The expression of cyclin A2 had no significant difference between two groups ($p < 0.05$). Tumor-specific antigen can increase the early apoptosis rate, late apoptosis rate and total apoptosis rate of MG63 cells compared to the non- tumor-specific antigen -treatment cells ($p < 0.05$). Specificity protein 1 could bind and target with Phospholipase promotor/enhancer region. Histone deacetylase inhibitor can reduce the proliferation and apoptosis of osteosarcoma cells through decreasing the expression of Phospholipase by inhibiting the protein expression and nuclear migration of specificity protein 1).

Key words: Osteosarcoma, histone deacetylase inhibitor, trichostatin A, specific protein 1, phospholipase D1

Osteosarcoma is a common and highly malignant bone-related tumor. It occurs frequently in 10 to 30 y old adolescents, ranking number two in cancer-related deaths in children and adolescents^[1]. In recent years, increasing new drugs have been used to treat osteosarcoma, but its

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*Address for correspondence

E-mail: jian_qin_njmu@sina.com

prognosis is still less than satisfactory^[2], the 5 y survival rate of osteosarcoma is only 50-60 %. Poor prognosis is associate to complex cell microenvironment of osteosarcoma which involves multiple factors and multiple pathways. Common chemotherapy lack sensitive targets for tumor cell recognition, and can easily lead to drug resistance. Therefore, osteosarcoma has its special pathogenesis and special target unlike other cancer. The treatment strategy based on traditional target and single target become more and more limited. Finding new targets to improve the efficacy of drug therapy, or developing new clinical drugs to achieve multiple targets of one drug are still the hopeful research directions for clinical breakthrough.

Human nuclear transcription factor specific protein 1 (SP1), a member of SP/KLF family, is widely expressed *in vivo*. It can specifically recognize GC rich elements in deoxyribonucleic acid (DNA) and participate in the transcriptional regulation of many genes. At present, SP1 is widely accepted as a tumor promoting protein. The expression of SP1 protein is low or absent in normal tissues, but high in gastric cancer, pancreatic cancer, breast cancer and thyroid tumor cells^[3]. Phospholipase D1(PLD1) is an important enzyme that regulates the metabolism of phospholipids. It catalyzes the hydrolysis of phosphatidylcholine to produce phosphonic acid (PA). PA plays an important role in cancer-related signal pathway. Most of its downstream target proteins have been confirmed to be related to cell proliferation and metastasis^[4]. Both SP1 and PLD1 are independent tumor risk factors and interact with each other^[5]. It is still not clear how SP1 and PLD1 regulate in osteosarcoma cells. In our previous study, we compared the tumor samples of 137 patients with osteosarcoma and found that the contents of SP1 and PLD1 in osteosarcoma tissues were significantly higher than those in adjacent tissues. The follow-up results showed that the increased expression of the two proteins in osteosarcoma tissue could lead to poor prognosis of patients with osteosarcoma^[6], SP1 and PLD1 may be potential therapeutic targets for osteosarcoma.

Histone deacetylase inhibitor (HDACi) is a new kind of antitumor drugs, which can block cell cycle, induce cancer cells apoptosis and differentiation. In 2006, the Food and Drug Administration (FDA) officially approved the first HDACi, Zolinza (vorinostat) to treat cutaneous T-cell lymphoma. At present, Zolinza has developed to treat a variety of hematological and solid tumors^[7,8]. In order to further study the effect and the mechanism of HDACi for osteosarcoma, we

choose a typical HDACi trichostatin A (TSA) to treat osteosarcoma cells *in vitro*, and intend to demonstrate if TSA can antitumor by targeting SP1 and PLD, and clarify the effect of SP1 after intervention of TSA. We intend to explore the specific mechanism of combined action of SP1 and PLD1 in tumor progression process, and provide a new target and a new drug for clinical treatment of osteosarcoma, and a new idea for clinical reversal of chemotherapy resistance of osteosarcoma.

MATERIALS AND METHODS

Materials:

Human osteosarcoma cell line MG63 (ATCC, CAT. NO. CRL-1427); penicillin, fetal bovine serum (FBS) (Sigma-Aldrich, USA), Lipofectamine 2000™ (Invitrogen, CAT. NO. 11668-027), PLD1 rabbit polyclonal antibody (Absin, Shanghai, CAT. NO. ABS137279), SP1 rabbit polyclonal antibody (AtaGenix CAT. NO. ATA37845), nuclear and cytoplasmic protein extraction kit (Beyotime, CAT. NO. P0027), Cell Counting Kit (CCK-8, DOJINDO, CK04), VaripKan Flash Microplate Reader (Thermo Scientific, USA), Trizol reagent (Invitrogen, CAT. NO. 15596-026).

Treatment of MG63 cells with TSA:

MG63 cells were resuscitated and cultured in Dulbecco's Modified Eagle's medium (DMEM) medium containing penicillin (100 U/mL) and 10 % FBS at 37° and 5 % CO₂. When MG63 cells overspread more than 80 % of the culture dish, trypsin was added for digestion and cell suspension was washed by PBS solution and collected by centrifugation. MG63 cells were diluted by DMEM medium into 2×10⁵ cells and transferred into 96 well plate at 37° and 5 % CO₂ for 24 h. Then MG63 cells were treated with TSA (5 g/ml) or PBS (negative control) for 48 h at 37° in a 5 % CO₂, six parallel wells for each treatment. After treatment with TSA, all medium was discarded and cells were harvested for following experiments.

Separation of nuclear and cytoplasmic protein:

After cell collection, we performed cytoplasmic and nuclear fractions separation by using nuclear and cytoplasmic protein extraction kit. In brief, cells treated as indication were resuspended in PBS, followed by addition to cytoplasmic protein extraction reagent A and B. Then, the samples were centrifuged, and the supernatant was collected to obtain cytoplasmic fraction. The nuclei was resuspended in nuclear protein

extraction agent. After centrifugation, the supernatant was collected as the nuclear fraction.

Western blot:

Nuclear and cytoplasmic protein was respectively treated by radioimmunoassay (RIPA) buffer and determined concentration by BCA method. The protein was heated at 95° for 5 min to lysate and was dissolved in 30 min by ice bath. Then the protein was extracted by dodecyl sulfonate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to the polyvinylidene fluoride (PVDF) membrane. The PVDF membrane was sealed by 5 % skimmed milk powder at room temperature and the added 1:1000 first antibody of SP1, PLD1 and β -actin at room temperature overnight. After washing with TBST, the PVDF membrane was added into HRP-labeled second antibody (diluted with 1:5000). After incubation for 1h at room temperature, the membrane was rinsed with TBST for three times, the positive bands were displayed with chemiluminescent reagent, and the images were processed and analyzed with gray-scale analysis software.

Down-regulation of SP1 in MG63 cells:

MG63 cells were resuscitated and cultured in DMEM medium containing penicillin (100 U/mL) and 10% FBS at 37° and 5 % CO₂. When MG63 cells overspread more than 80 % of the culture dish, trypsin was added for digestion and cell suspension was washed by PBS solution and collected by centrifugation. To down-regulation of SP1, si-SP1 was synthesized by Shanghai JIKAI GENE Chemistry Technology Co., Ltd. To synthesize siRNA-SP1, the forward primer was 5'-ATTTCGATCGGGGCGGGGCGAGC-3' and reverse primer was 5'-AUAGUUGAGUCATGT-3'. siR-SP1 was transfected into MG63 cells by Lipofectamine 2000™.

Dual luciferase reporter gene assay:

According to the sequence of PLD1 promoter/enhancer region (-154 to -109, 45bp), two reverse complementary DNA primers were designed. The primers were heated and then linked with dual luciferase reporter to construct psiCHECK-PLD1-WT plasmid. psiCHECK-PLD1-MT were constructed as mutation of psiCHECK-PLD1-WT for negative control. The construction process was completed by Shanghai Jima Company.

MG63 cells after transfection with si-SP1 or si-NC were harvested and adjusted the cell density to 1×10⁵ cells/ml. the cells were transferred on a 96 well plate

with 0.1 ml per well. It was incubated overnight in a 5 % CO₂ incubator at 37°. The psiCHECK-PLD1-WT and psiCHECK-PLD1-MT plasmid were respectively transfected into MG63 cells by Lipofectamine 2000™. After transfection, the culture medium was discarded and washed with 100 μ l PBS for two times; PLB solution was added to the cells at room temperature for 50 rpm/min for 15 min. The cell lysate was collected; 20 μ L cell lysate and 100 μ l LAR II were added to each well of the black opaque 96 well enzyme label plate, and the data were detected automatically after standing in dark for 2 s; 100 μ l STOP & GLO Regent was added into each well, and the data was detected automatically after standing in dark for 2 s.

Cell proliferation analysis:

Cell Counting Kit was used to measure the effects of TSA on cell viability^[9]. Briefly, After indicated TSA exposure, 10 μ l of CCK8 reagent was added followed by incubation for another 2 h at 37° and 5 % CO₂. The optical density (OD) value at 450 nm was recorded by the Microplate Reader.

Clone formation assay:

1.2 % and 0.6 % low melting point agarose solutions were sterilized and kept in semisolid state by 40° water bath. 1.2 % low melting point agarose solution and 2×DMEM medium (containing 20% FPS) were mixed and transferred into 6 well plate as the bottom agar. 0.3 % soft agar was prepared by mixing 0.6% low melting point agarose solution and 2×DMEM medium (containing 20 % FPS).

The MG63 cells treated with TSA or PBS were harvested by centrifuging at 1500 rpm for 10 min. The supernatant was removed and washed twice with PBS solution. The cell density was adjusted with DMEM medium to 500 cells/ml. 1ml of cells solution and 1 ml 0.3 % soft agar were inoculated into six well plate. Culture in incubator at 37° and 5 % CO₂ -95 % air mixture for 10 d. The cells were washed with PBS twice, and fixed with methanol for 10 min. The cells were dyed with 0.1 % crystal violet for 20 min and washed with PBS. Under 400×microscope, the cells were observed in 5 fields randomly and counted the number of colonies.

Quantitative real-time polymerase chain reaction (qRT-PCR):

Total RNA was isolated with Trizol following the manufacturer's protocol. Total RNAs were reverse transcribed into cDNA using the Reverse Transcription

kit. qRT-PCR analysis was performed in SYBR GREEN PCR Master Mix in 20 μ l reactions. The related mRNA expression levels were normalized to the β -actin by $2^{-\Delta\Delta CT}$. The primers are synthesized by Shanghai Sangon Company, the primer sequence were listed in Table 1. PCR program was 95°: 30s, (95°: 5s, 60°: 34s) \times 40 cycles.

Flow cytometry:

The MG63 cells treated with TSA or PBS were harvested by centrifuging at 1500 rpm for 10 min. The supernatant was removed and washed twice with PBS solution. The cell density was adjusted to 2×10^6 cells/group. Then the MG63 cells were centrifugated to remove the supernatant and replace with pre-cooled 70% ethanol to incubate at -20° overnight. The MG63 cells were centrifugated at 1500 rpm for 5 min for removal the supernatant, then washed with PBS twice. Propidium iodide (PI) solution was added into the cells and dyed at room temperature and avoided light for 30 min. Flow cytometry was performed to measure the cell cycle within 1 h.

Statistical analysis:

SPSS 22.0 software was used for statistical analysis. Data were present as ($\bar{x} \pm s$). Student's T-test was used for comparison between two groups, $p < 0.05$ means there was statistical significance between different groups.

RESULTS AND DISCUSSION

Nuclear and cytoplasmic SP1 in TSA treated MG63 cells both decreased dramatically compared to TSA free cells ($p < 0.05$). PLD1 in TSA treated MG63 cells also reduced ($p < 0.05$), as shown in fig. 1.

As shown in fig. 2, fluorescence analysis showed that the ratio of Firefly luciferase to Renilla luciferase in MG63 cells transfected with psiCHECK-PLD1-WT plasmid and si-SP1 was significantly lower than that of si-NC ($p < 0.05$), while the ratio of firefly luciferase to Renilla luciferase in MG63 cells transfected with psiCHECK-PLD1-WT plasmid and si-SP1 had no significant difference with si-NC control ($p > 0.05$), indicating that the decrease of Firefly luciferase to Renilla luciferase ratio was due to the expression of reporter gene after SP1 binding to PLD1 promotor/enhancer. It was proved that SP1 could bind and target with PLD1 promotor/enhancer region (fig. 2).

The cell viability of MG63 cells treated with TSA or not was measured by CCK-8 analysis. As shown in fig. 3, after TSA treatment for 24 h, the proliferation rate of MG63 cells was ($65.20 \% \pm 2.31 \%$), which was significantly lower than that of PBS control group ($96.63 \% \pm 1.14 \%$) ($p < 0.05$).

Cell formation assay showed that the number of MG63 cells treated with TSA decreased significantly ($p < 0.05$), indicating that TSA could inhibit the proliferation of osteosarcoma cells. The photos and statistical results are shown in fig. 4.

TABLE 1: THE PRIME SEQUENCES FOR QRT-PCR

Gene	Forward primer sequence	Reverse primer sequence
SP1	5'-CACCAGAATAAGAAGGGAGG-3'	5'-GGTGGTAATAAGGGCTGAA-3'
PLD1	5'-AATCGTTGGAGTTGGACTG-3'	5'-AGACGGTGGATGACACATGA-3'
β -actin	5'-AGCGAGCATCCCCAAAGTT-3'	5'-GGGCACGAAGGCTCATCATT-3'
p21	5'-TGCTTTGTACCCTTGTGCCTCG-3'	5'-AATCTGTCATGCTGGTCTGCC-3'
Survivin	5'-TCCCTGGCTCCTCTACTG TT-3'	5'-TGTCTCCTCATCCACCTGAA-3'
Cyclin A2	5'-TCCTCGTGACTGGTTAGTTGA-3'	5'-GCACTGACATGGAAGACAGGAA-3'

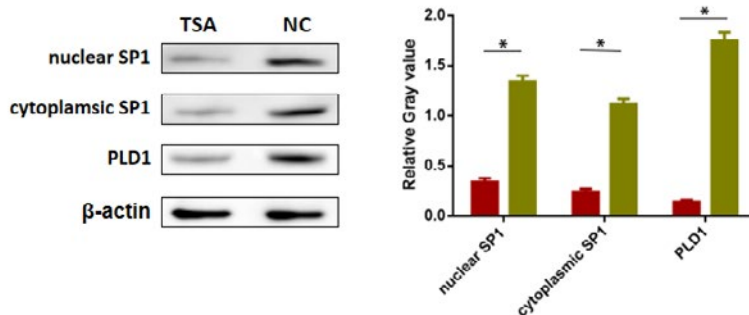


Fig. 1: Western blot of PLD1, nuclear and cytoplasmic SP1 (■): TSA (■) NC ($*p < 0.05$ vs NC, $n=6$)

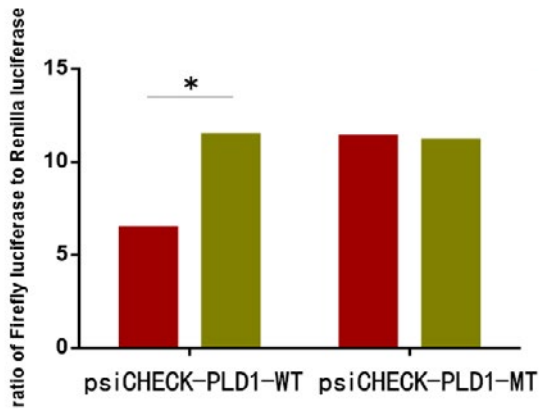


Fig. 2: Dual luciferase reporter gene assay result (■) si-SP1 (■) si-NC (*p<0.05 vs si-NC)

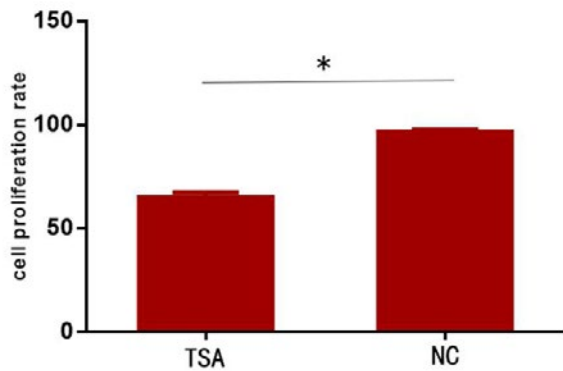


Fig. 3: CCK-8 assay result (*p<0.05 vs NC, n=6)

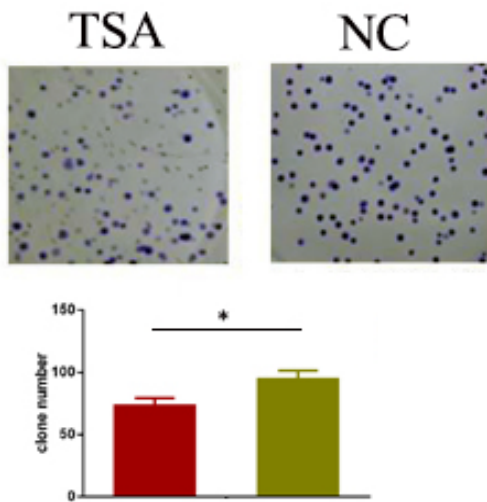


Fig. 4: Photos and result of clone formation assay (■): TSA (■) NC (*p<0.05 vs NC, n=5)

As shown in fig. 5, the gene expression of SP1 and PLD1 in TSA-treated MG63 cells decreased significantly compared with TSA free cells (p<0.05). The gene expression levels of survivin which inhibits cells

apoptosis in TSA-treated MG63 cells were significantly lower than that in TSA free cells (p<0.05). The gene expression levels of p21 which induces cells apoptosis in TSA-treated MG63 cells were significantly higher than that in TSA free cells (p<0.05). The expression of cyclin A2 gene had no significant difference between two groups (p<0.05).

As shown in fig. 6, TSA can increase the early apoptosis rate, late apoptosis rate and total apoptosis rate of MG63 cells compared to the non-TSA-treatment cells (p<0.05) (fig. 5).

HDACi is a new class of antitumor drugs by blocking cell cycle, inducing apoptosis and differentiation. Histone deacetylation is harmful to DNA and histone for preventing dissociation of protein octamer, then inhibits the specific binding of various transcription factors and co transcription factors with DNA binding sites and

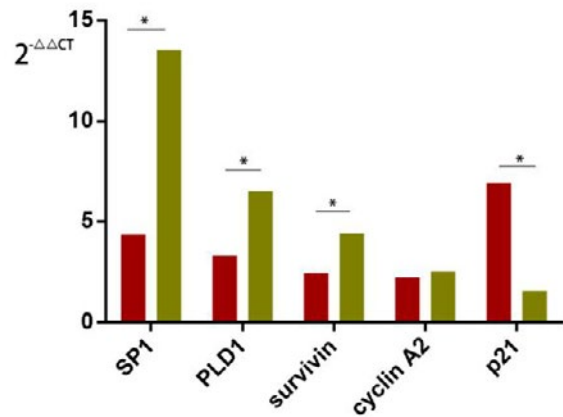


Fig. 5: PCR assay result (■): TSA (■) NC (*p<0.05 vs NC)

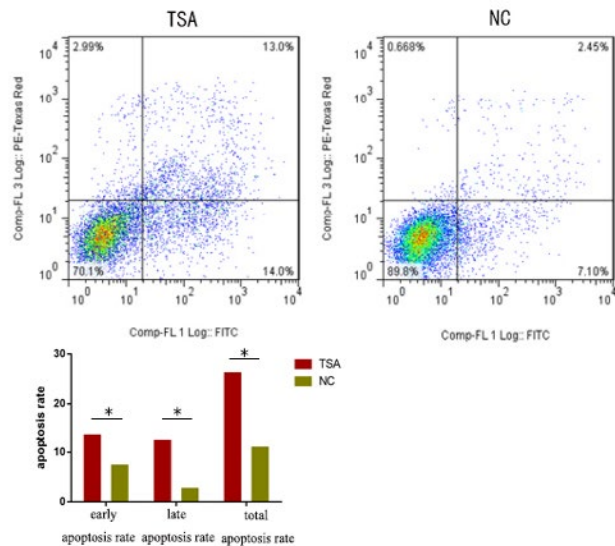


Fig. 6: Flow cytometry result for apoptosis rate (■): TSA (■) NC (*p<0.05 vs NC)

gene transcription. For the treatment of osteosarcoma, HDAC inhibitors such as AR-42 and TSA can activate Akt, p53, VEGF / DR3 related pathways and promote osteosarcoma cell apoptosis. It has been reported that the expression of SP1 in bladder cancer cells was decreased by TSA. In recent years, several studies have suggested that transcription factor SP1 is a tumor promoter for colon cancer^[10] and prostate cancer^[11]. SP1 plays an important role in regulating tumor related gene expression by self-modifying protein complex structure and HDAC inhibitors^[12]. Our previous studies have shown that SP1 inhibitor mithramycin A and transfection of SP1 siRNA to human osteosarcoma cell line MG63 could decrease the protein expression and promoter activity of PLD1, indicating that SP1 may participate in the pathogenesis and progress of osteosarcoma by regulating PLD1 expression. PLD1 present widely in different tissues and cells and are activated by a wide variety of receptor tyrosine kinases (RTKs) and G protein-coupled receptors (GPCRs)^[13]. Biological function of PLD1 includes regulated exocytosis, endocytosis, proliferation, cell migration, Golgi-ER trafficking, autophagy, and apoptosis, which encompasses almost all signaling-driven processes^[14]. Interest has become increasingly focused on PLD1 in the context of cancer. Therefore, based on the literatures and our previous research results, we speculate that HDACi may target and regulate SP1 and effect the expression of PLD1, thus inhibit the proliferation and promote cells apoptosis of osteosarcoma cells.

In this article, we used TSA, a typical and extensive HDACi, to treat osteosarcoma cell line MG63 *in vitro*. qPCR and Western blot results showed that after TSA treatment, SP1 in nucleus and cytoplasm of MG63 both decreased significantly, and PLD1 also decreased significantly. These results indicated that HDACi can inhibit the expression of SP1 protein and nuclear migration, thus inhibiting the expression of PLD1. In addition, this study demonstrated that SP1 could bind to the promoter / enhancer fragment of PLD1 gene by dual luciferase reporter gene assay. So SP1 is the regulatory transcription factor of PLD1 in osteosarcoma cells. TSA inhibits the expression of PLD1 by inhibiting SP1. Then, CCK-8 test and clone formulation analysis showed that TSA could reduce the proliferation level and cell activity of MG63 cells. The results of qPCR showed that TSA could inhibit survivin and increase p21^[15]. The results of flow cytometry showed that TSA could promote the early and late apoptosis rates of MG63 cells.

In conclusion, SP1 could bind and target with PLD1 promoter/enhancer region. HDACi can reduce the proliferation and apoptosis of osteosarcoma cells through decreasing the expression of PLD1 by inhibiting the protein expression and nuclear migration of SP1.

Authors' contributions:

Jian qin conceived and designed the experiments; Tangbo Yuan and Zhen Jin performed the experiments; Dawei Cai analyzed the data and wrote the paper.

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Conflicts of interest:

The authors report no conflicts of interest.

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