RESEARCH ARTICLE



Valproic acid inhibits the angiogenic potential of cervical cancer cells via HIF-1α/VEGF signals

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Abstract

Purpose Cervical cancer is one of the most prevalent malignancies in women worldwide. Therefore, the investigation about the molecular pathogenesis and related therapy targets of cervical cancer is an emergency. The objective of the present study is to investigate the effects of valproic acid (VPA), a histone deacetylase inhibitor, on the angiogenesis of cervical cancer.

Methods The effects and mechanisms of VPA on in vitro angiogenesis and vascular endothelial growth factor (VEGF) expression of human cervical cancer HeLa and SiHa cells were investigated.

Results Our present study reveals that 1 mM VPA can significantly inhibit the in vitro angiogenic potential and VEGF expression of human cervical cancer HeLa and SiHa cells. Further, the transcription and protein levels of hypoxia inducible factor- 1α (HIF- 1α), and not HIF- 1β , are significantly inhibited in VPA-treated cervical cancer cells. Over expression of HIF- 1α can obviously reverse VPA-induced VEGF down regulation. VPA-treatment decreases the activation of Akt and ERK1/2 in both HeLa and SiHa cells in a time-dependent manner. The inhibitor of Akt (LY

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294002) or ERK1/2 (PD98059) can inhibit VEGF alone and cooperatively reinforce the suppression effects of VPA on HIF-1 α and VEGF expression.

Conclusion Collectively, our data reveal that the inhibition of PI3K/Akt and ERK1/2 signals are involved in VPA-induced HIF-1 α and VEGF suppression of cervical cancer cells.

Introduction

Cervical cancer is one of the most prevalent malignancies in women worldwide, especially in developing countries. On a global scale, it is estimated that there are 530,000 new cases and 275,000 resultant deaths each year [1]. Human papilloma virus (HPV) infection is the leading cause of cervical cancer. Genetic and epigenetic alternations are also important for the tumorigenesis and development of cervical cancer besides HPV. Surgeries accompanied with radiotherapy and chemotherapy have been considered as the major treatment method for cervical cancer [2]. Generally, chemotherapy is used to treat patients with metastasis or recurrence at times. Although some causes of cervical cancer have been illustrated, the precise mechanisms for the progression, metastasis and angiogenesis are still largely unknown. There is an urgent need to investigate the molecular pathogenesis and related therapy targets of cervical cancer.

Epigenetic modifications have been suggested to act as an important role in cancer cell biology [3, 4]. Histone deacetylases (HDACs), which remove acetyl groups from the *N*-acetyllysines on histone and non-histone proteins, have garnered great interest as anti-cancer therapeutic targets [5]. Valproic acid (VPA, 2-propylpentanoic acid), which has long been used clinically for treatment of epilepsy and bipolar disorder without significant toxicity [6], is suggested to be a HDAC inhibitor (HDACI) in recent years [7]. Numerous studies show its anti-cancer effects in various cancer models including lung, renal, bladder, and cervical cancer via inhibition of cell proliferation and inducing cell apoptosis [8–11]. However, other roles and mechanisms of VPA on the progression of cervical cancer remain to be further illustrated.

Angiogenesis, which describes the formation of new blood vessels from the existing vasculature, is critical for tumor growth and metastasis [12]. Therefore the antiangiogenic agents have gathered special attention for the development of anti-cancer drugs. Within tumors, circulating cells such as bone marrow derived endothelial progenitor cells, macrophages and fibroblasts with malignant cells are able to secrete pro-angiogenic factors including vascular endothelial growth factor (VEGF) and then induce tumor blood vessel formation [13]. Recent studies indicated that HDACIs such as FK228 can inhibit the expression of angiogenic factors, including VEGF and basic fibroblast growth factor (bFGF), and therefore suppress the angiogenesis of cancer cells [12, 14, 15]. Although few studies also suggested that VPA can modulate the progression of angiogenesis in cancer cells [16-18], there is insufficient amount of data on its effects in cervical cancer cells. The present study investigates the roles and related mechanisms of VPA on the angiogenic potential of cervical cancer cells and suggests that VPA can down-regulate VEGF via HIF-1a signals.

Materials and methods

Cell culture and transfection

Human cervical cancer cell lines HeLa and SiHa were obtained from the American Type Culture Collection (ATCC) and cultured in DMEM containing 10 % FBS and antibiotics. All cells were grown at 37 °C in a humidified 5 % CO₂ incubator. VPA was dissolved in DMSO with the final concentration of DMSO in the medium less than 0.5 %. For transfection, HeLa or SiHa cells were seeded into plates and transfected with siRNA negative control (siRNA-NC: 5'-GGC TAC GTC CAG GAG CGC A-3'), si-HIF-1 α (sequence 5'-GAG GAA ACU UCU GGA UGC UGG UGA Utt-3'), pcDNA3.1 (vector control), or pcDNA3.1/HIF-1 α by use of Lipofectamine 2000 reagent (Invitrogen) according to the manufacturer's instructions.

Cell proliferation assays

For the cell proliferation assays, HeLa or SiHa cells $(7 \times 10^3 \text{ cells/well})$ were separately seeded in 96-well plates and then treated with various concentrations of VPA for the indicated times. Then cell viability was assessed using the MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay as previously described [19].

Tube formation assays

Human umbilical vein endothelial cells (HUVECs) were obtained from Pricells (Wuhan, Hubei, China), suspended in conditioned media, and seeded on growth factor-reduced Matrigel (BD Biosciences, Franklin Lakes, NJ, USA) in a 12-well plate. After incubation with different condition medium, the number of mature, well-connected capillary tubes were manually counted under an inverted Nikon Eclipse Ti microscope (Nikon USA, Melville, NY, USA) and compared among different groups.

Quantitative real-time PCR analysis (qRT-PCR)

Total RNA samples were extracted from cells using Trizol reagent (Invitrogen, Carlsbad, CA, USA). The RNA was converted into cDNAs using the RevertAid First Strand cDNA Synthesis Kit (Fermentas, K1621). The cycle number when the fluorescence first reached a preset cycle threshold (Ct) was used to quantify the initial concentration of individual templates for expression of mRNA of genes of interest. Transcripts of the housekeeping gene GAPDH in the same incubations were used for internal normalization. Primer sequences were as follows: VEGF, 5'-TCG AGT ACA TCT TCA AGC CAT CCT G-3' and 5'-TCC TAT GTG CTG GCC TTG GTG AG-3'; GAPDH, 5'-TGA ACG GGA AGC TCA CTG G-3' and 5'-TCC ACC ACC CTG TTG CTG T-3'. After normalization to GAPDH gene, expression levels for each target gene were calculated using the comparative CT method. The Dct values were calculated according to the formula $\Delta ct = ct$ (gene of interest) - ct (GAPDH) in correlation analysis, and the $2^{-\Delta\Delta ct}$ was calculated according to the formula $\Delta\Delta ct =$ Δct (control group) – Δct (experimental group) for determination of relative.

Quantification of VEGF protein

VEGF protein was measured using the Quantikine human VEGF ELISA kit from R&D Systems (Minneapolis, MN, USA) according to previous study [20]. Briefly, cells were seeded in 12-well plates and cultured to 90–100 % confluence. Cells were switched to fresh medium in the

presence or absence of 1 mM VPA. After treatment, the supernatants were collected and cell numbers of each well were counted. VEGF in the supernatant (100 μ L) was determined and normalized to the remaining cell numbers. A serial dilution of human recombinant VEGF was included in each assay to obtain a standard curve.

Western blot analysis

Cells were grown in 6-cm dishes, exposed to VPA for the indicated times, and then lysed in 500 μ L of SDS buffer (0.125 M Tris–HCl, 10 % 2-mercaptoethanol, 4 % SDS, and 10 % sucrose) and sonicated on ice for 30 s. The protein concentrations were determined with the bicinchoninic acid method. Equal quantities of proteins were loaded and separated by electrophoresis in SDS-PAGE and then transferred onto polyvinylidene fluoride membranes. The membranes were incubated with primary antibodies at 4 °C overnight. Blots were stained with horseradish peroxidase-linked goat anti-rabbit Ig secondary antibodies and developed with a chemiluminescence detection system (Fujifilm).

Statistical analyses

Statistical analyses were performed using SPSS 17.0 (SPSS, Inc., Chicago, IL, USA). All the data presented are based on an average of at least three independent experiments. The values are presented as the mean \pm standard deviation. Student's *t* test's were used to compare the differences between groups. *P* < 0.05 was used to indicate a statistically significant difference.

Results

The effects of VPA on the in vitro proliferation of cervical cancer cells

To explore whether VPA might be a potential therapeutic agent against cervical cancer, we investigated its effects of growth inhibition in human cervical cancer HeLa and SiHa cells. Our results indicated that VPA can inhibit the proliferation of HeLa (Fig. 1a) and SiHa (Fig. 1b) cells via both time and concentration dependent manners. The IC₅₀ of VPA in HeLa and SiHa cells at 24 h was 5.6 and 6.3 mM, respectively, and at 48 h was 2.8 and 3.4 mM, respectively. Therefore, 1 mM VPA was chosen for further studies due to having no significant effect on the proliferation of cervical cancer cells and on the basis of other previous studies [21, 22].

VPA inhibits the angiogenic potential of cervical cancer cells

Recent studies indicated that HDACIs including VPA can modulate the progression of angiogenesis in cancer cells [16–18]. Then we tested the effects of VPA on the angiogenic potential of cervical cancer cells by use of in vitro tube formation assay. Our results showed that tube formation of HUVEC cells was significantly (p < 0.01) inhibited by medium prepared from 1 mM VPA-treated HeLa and SiHa cells (Fig. 2a, b). These results suggested that VPA can inhibit the angiogenic potential of cervical cancer cells.



Fig. 1 The effects of VPA on the in vitro proliferation of cervical cancer cells. HeLa (a) or SiHa (b) were treated with various concentrations of VPA for 24 or 48 h, and then cell viability was assessed by MTT assay. Data were presented as mean \pm SD of six independent experiments



Fig. 2 VPA inhibits the angiogenic potential of cervical cancer cells. a HeLa or SiHa cells were treated with 1 mM VPA for 24 h, then the medium of HeLa or SiHa cells was used for conditioned media to test

the tube formation for HUVEC cells. **b** Quantitative analysis of tube formation assays. Data represented the average of three independent experiments. *p < 0.05 compared with control





Fig. 3 VPA inhibits the expression of VEGF in cervical cancer cells. **a** HeLa or SiHa cells were treated with 1 mM VPA for the indicated times, and then the mRNA levels of VEGF were measured by qRT-PCR; **b** HeLa or SiHa cells were treated with 1 mM VPA for the

VPA inhibits VEGF expression of cervical cancer cells

Then we determined the effects of VPA on VEGF mRNA levels in cervical cancer cells. As shown in Fig. 3a, VPA can inhibit the expression of VEGF mRNA in a time-dependent manner in both HeLa and SiHa cells. Then the protein expression of VEGF was measured in VPA treated HeLa or SiHa cells by use of ELISA. The results confirmed that VPA can significantly decrease the production of

indicated times, the protein expression of VEGF was measured by ELISA. Data represent the average of three independent experiments. *p < 0.05 compared with control

VEGF protein in both cell lines (Fig. 3b). Collectively, our results suggested that VPA can inhibit the expression of VEGF in cervical cancer cells.

HIF-1α mediates the suppression effects of VPA on VEGF expression of cervical cancer cells

Hypoxia inducible factor 1 (HIF-1) can activate the expression of the VEGF gene by binding to the hypoxia response element (HRE) in the VEGF promoter [23]. HIF-



Fig. 4 HIF-1 α mediates the suppression effects of VPA on VEGF expression of cervical cancer cells. **a** HeLa or SiHa cells were treated with 1 mM VPA for the indicated times, and then protein levels of HIF-1 α or HIF-1 β were measured by use of Western blot analysis; **b** HeLa or SiHa cells were treated with 1 mM VPA for 12 h, the mRNA of HIF-1 α or HIF-1 β was measured by qRT-PCR; **c** HeLa or SiHa cells were transfected with pcDNA3.1 (vector) or pcDNA/HIF-

1 is composed of HIF-1 α and HIF-1 β subunits [24]. To determine whether HIF-1 is involved in VPA-induced VEGF suppression, we examined the effects of VPA on HIF-1 α and HIF-1 β protein levels. The results showed that VPA can obviously decrease the expression of HIF-1 α in both HeLa and SiHa cells, however, VPA had no significant effect on HIF-1 β protein levels (Fig. 4a). This was further confirmed by the results that VPA can significantly inhibit the mRNA levels of HIF-1 α , but not HIF-1 β , in both HeLa and SiHa cells (Fig. 4b).

To verify the roles of HIF-1 α in VPA-induced VEGF suppression, HeLa or SiHa cells were transfected with pcDNA3.1 (vector) or pcDNA/HIF-1 α for 24 h, then further treated with VPA for 24 h. The over expression of HIF-1 α was confirmed by the results of Western blot analysis (Fig. 4c). Our results showed that over expression of HIF-1 α significantly attenuated VPA-induced down-

1α for 24 h, the expression of HIF-1αwas measured by use of Western blot analysis; **d** HeLa or SiHa cells were transfected with pcDNA3.1 (vector) or pcDNA/HIF-1α for 24 h and then treated with 1 mM VPA for another 24 h, the protein expression of VEGF was measured by ELISA. Data represented the average of three independent experiments. **p < 0.01 compared with control

regulation of VEGF in both HeLa and SiHa cells (Fig. 4d). Collectively, our results revealed that HIF-1 α mediates the suppression effects of VPA on VEGF expression of cervical cancer cells.

VPA suppresses the activities of PI3K/Akt and ERK1/2 in cervical cancer cells

Several upstream signals including PI3K/Akt and ERK1/2 can regulate the expression of HIF-1 α and VEGF [25, 26]. To identify which signal is responsible for the inhibition effects of VPA on expression of HIF-1 α , we determined the effects of VPA on activation of Akt and ERK1/2 by Western blot analysis. The results showed that 1 mM VPA can obviously inhibit Akt phosphorylation in both HeLa (Fig. 5a) and SiHa (Fig. 5b) cells. Further, incubation of cells with VPA also resulted in significant decrease of



Fig. 5 The effects of VPA on activation of PI3K/Akt and ERK1/2 of cervical cancer cells. HeLa (\mathbf{a}) or SiHa (\mathbf{b}) cells were treated with 1 mM VPA at the indicated times, and then total and phosphorylation levels of Akt and ERK1/2 were measured by use of Western blot analysis

phospho-ERK1/2 levels. These results suggested that both PI3K/Akt and ERK1/2 might be involved in the effects of VPA on HIF-1 α and VEGF down-regulation.

Inhibition of PI3K/Akt and ERK1/2 cooperatively reinforces VPA-induced HIF-1α and VEGF suppression

To test whether PI3K/Akt and ERK1/2 signals are involved in VPA-induced HIF-1 α and VEGF suppression, HeLa and SiHa cells were treated with PI3K/Akt inhibitor (LY 294002, LY) and ERK1/2 inhibitor (PD98059, PD) alone or combination with VPA. Our results showed that both LY 294002 and PD98059 can obviously inhibit the expression of HIF-1 α , further, the inhibition effects of VPA on HIF-1 α expression were significantly reinforced with the cooperation of LY 294002 or PD98059 in both HeLa and SiHa cells (Fig. 6a). Similar results were also observed for the effects of VPA on VEGF protein (Fig. 6b, c) and mRNA (Fig. 6d, e) expression. These data suggested that the inhibition of PI3K/Akt and ERK1/2 signals are involved in VPA-induced HIF-1 α and VEGF suppression of cervical cancer cells.

Discussion

Tumors cannot grow larger than 1-2 mm in diameter without vessel formation, therefore anti-angiogenesis is considered as a new attractive strategy to overcome cancer. In the present study, VPA, with a concentration which has no significant effect on cell proliferation, can inhibit the VEGF expression and in vitro angiogenic potential of cervical cancer cells. VPA-treatment significantly suppresses the expression of HIF-1 α , while not HIF-1 β . Over expression of HIF-1 α can reverse the suppression effects of VPA on VEGF expression. Further, the inhibition of PI3K/ Akt and ERK1/2 also participates in VPA-induced HIF-1 α / VEGF down-regulation. Collectively, our present study shows that VPA can inhibit the angiogenic potential of cervical cancer cells via HIF-1 α /VEGF signals. The identified anti-angiogenic activities of VPA in the present study suggested that it can become a valuable new addition in the attempt to develop alternative therapeutic approaches for the treatment of cervical cancer.

Inhibition of VEGF production is a promising therapeutic approach for cancer. HDACI is a new class of anticancer therapeutics which show great promise at inhibiting VEGF and angiogenesis [12]. Due to the low cost and favorable safety profile, VPA has been considered as a candidate drug for various cancer cell types [27-29]. Several recent studies also indicated that VPA can inhibit proliferation, migration, and tube formation of human umbilical vein endothelial cells both in vitro and in vivo [30, 31]. In our present study, VPA can significantly inhibit the in vitro angiogenic potential and VEGF expression of both HeLa and SiHa cells. This is consistent with previous study that VPA can impair the secretion of VEGF in colon and prostate cancer cells [32, 33]. Several studies also indicated that VPA can regulate the expression of VEGFR, FGF and other angiogenesis related factors in colon and myeloma cancer cells [18, 32, 34]. These effects of VPA on cervical cancer need further studies.

Various mechanisms have been reported to be involved in HDACI-induced anti-angiogenesis of cancer cells [12, 35]. In the present study, our results show that VPA can inhibit the HIF-1 α /VEGF signals and then suppress the angiogenic potential. Increasing evidences suggest that HIF-1 plays a critical role in tumor progression and acts as a key regulator of VEGF expression. It has been documented that HDACIs, including VPA, were able to repress HIF-1 function in cancer cells [9, 36]. HIF-1 α activates its



Fig. 6 Inhibition of PI3K/Akt and ERK1/2 cooperatively reinforces VPA-induced HIF-1 α and VEGF suppression. **a** HeLa or SiHa cells were treated with VPA (1 mM), LY294002 (10 μ M), or PD98059 (10 μ M) alone or together for 24 h, and then protein level of HIF-1 α was measured by use of Western blot analysis; HeLa (**b**) or SiHa (**c**) cells were treated with VPA (1 mM), LY294002 (10 μ M), or

target genes such as erythropoietin (EPO), VEGF, and phosphoglycerate kinase 1 (PGK1) through dimerization with HIF-1 β . In the present study, our data showed that VPA-treatment can significantly suppress the expression of HIF-1 α , while not HIF-1 β , in cervical cancer cells. This might be due to the fact that VPA can significantly inhibit the transcription of HIF-1 α . Further, overexpression of HIF-1 α abolishes VPA-induced VEGF down-regulation. It confirms the essential role of HIF-1 α in the suppression effects of VPA on angiogenic potential of cervical cancer cells.

The mechanisms responsible for HDACIs, including VPA, induced HIF-1 α down regulation are not fully illustrated. Some literature suggests that HDACIs decrease the stability of HIF-1 α through enhancing the acetylation of HIF-1 α and/or other proteins involved in modulating the degradation of HIF-1 α , thus accelerating the degradation of HIF-1 α [37]. Enhanced signaling through the PI3K/Akt and

PD98059 (10 μ M) alone or together for 24 h, the protein expression of VEGF was measured by ELISA; HeLa (**d**) or SiHa (**e**) cells were treated with VPA (1 mM), LY294002 (10 μ M), or PD98059 (10 μ M) alone or together for 12 h, the mRNA expression of VEGF was measured by qRT-PCR. Data represented the average of three independent experiments

MAPK pathways are a significant contributor to activate HIF- 1α /VEGF signals [26]. However, the pathways responsible for VEGF production may be dependent on the cell lines. It has been reported that activation of the Akt pathway, but not the ERK pathway, induced VEGF production in the PC3 human prostate cancer cell line [38]. Inversely, activation of the ERK pathway, but not the Akt pathway, promoted VEGF production in human HaCaT keratinocytes [39]. Our present study reveals that both the activation PI3K/Akt and ERK1/2 are inhibited by VPAtreatment; furthermore, the inhibitors of PI3K/Akt and ERK1/2 can cooperatively reinforce the inhibition effects of VPA on HIF-1 α and VEGF expression. It suggests that both PI3K/Akt and ERK1/2 are involved in the suppression effects of VPA on angiogenic potential of cervical cancer cells.

In conclusion, VPA inhibits angiogenic potential of cervical cancer cells via suppression of HIF-1 α /VEGF

signals. The suppression of PI3K/Akt and ERK1/2 signals are both involved in this process. Though the precise mechanism remains undetermined, these findings will help to better understand the roles and mechanisms of VPA as a potential anti-cancer drug for cervical cancer patients.

Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

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