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The histone deacetylase inhibitor valproic acid sensitizes diffuse large B-cell lymphoma cell lines to CHOP-induced cell death

Malin Ageberg¹, Karin Rydström², Thomas Relander², Kristina Drott¹,²

¹Division of Hematology and Transfusion Medicine, Lund University, BMC B13, Klinikg. 26, S-22184 Lund, Sweden; ²Skåne Department of Oncology, Lund University Hospital, S-22185 Lund, Sweden

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Abstract: Epigenetic code modifications by histone deacetylase inhibitors (HDACis) have recently been proposed as potential new therapies for hematological malignancies. Diffuse large B-cell lymphoma (DLBCL) is the most common form of aggressive lymphoma. At present, standard first line treatment for DLBCL patients is the antracycline-based chemotherapy regimen CHOP (cyclophosphamide, doxorubicin, vincristine and prednisone) combined with the monoclonal anti-CD20 antibody rituximab (R-CHOP). Since only 50-60% of patients reach a long-time cure by this treatment, there is an urgent need for novel treatment strategies to increase the response and long-term remission to initial R-CHOP therapy. In this study, we investigated the effect of the HDAC inhibitor valproic acid (VPA) on DLBCL cell lines. To elucidate the effects of VPA on chemo-sensitivity, we used a cell-line based model of CHOP-refractory DLBCL. All five DLBCL cell lines treated with VPA alone or in combination with CHOP showed decreased viability and proliferation. The VPA-induced sensitization of DLBCL cells to cytotoxic treatment resulted in increased number of apoptotic cell as judged by annexin V-positivity and the presence of cleaved caspase-3. In addition, pretreatment with VPA resulted in a significantly increased DNA-damage as compared to CHOP alone. In summary, HDAC inhibitors such as VPA, are promising therapeutic agents in combination with R-CHOP for patients with DLBCL.

Keywords: Non-Hodgkins lymphoma, valproic acid, valproate, HDAC

Introduction

Diffuse large B-cell lymphoma (DLBCL) is the most common subgroup of non-Hodgkin lymphoma, accounting for approximately 30% of all lymphomas. DLBCLs belong to the group of aggressive lymphoma, and usually result in death within a couple months if left untreated. DLBCL is most commonly diagnosed in middle-aged and elderly individuals. Antracycline-based chemotherapy comprised of cyclophosphamide, doxorubicin, vincristine and prednisone (CHOP) in combination with rituximab, a monoclonal antibody against CD20, is considered to be the standard treatment regimen for patients with DLBCL. Despite the improved prognosis after introduction of anti-CD20 therapy to the conventional CHOP treatment, the 5-year progression-free survival rate remains around 60% [1-3]. Hence, the need for new therapeutic agents that augment sensitiv-
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should aim at restoring physiologic acetylation levels, and the use of inhibitors of histone acetylation could have a rational basis in DLBCL.

Several histone deacetylase inhibitors (HDACis) are shown to have effect on specific tumor types as single agent drugs and hematological malignancies seem to be particularly sensitive to HDAC inhibitors. Accordingly, vorinostat (Zolinza® or SAHA) and romidepsin (Istodax®) were approved by the FDA in 2006 and 2009, respectively, for the treatment of cutaneous T-cell lymphoma (CTCL) [6]. Also, in 2011, FDA approved romidepsin for the treatment of patients with peripheral T-cell lymphoma following at least one prior therapy [7]. Vorinostat and the HDAC class I specific inhibitor, MGCD01103, has been tested as a monotherapy for the treatment of relapsed and refractory DLBCL but with limited activity [8]. Several other HDAC inhibitors are under evaluation in clinical trials both as single agents and in combination with chemotherapeutic drugs [9].

In 2001, valproic acid (VPA), a GABA agonist with a long history of clinical use for treatment of epilepsy and mood disorders (reviewed in [10]), was identified having HDAC inhibitory activity [11]. VPA is a short-chain fatty acid that has been shown to inhibit the class I and II HDAC enzymes [10, 12]. VPA was recently shown to bind with high affinity to the hydrophobic active site channel of HDAC8 by van der Waals interactions [12]. Since its identification as an HDAC inhibitor, VPA has been suggested to regulate several mechanisms involved in malignant transformation such as cell cycle control, differentiation, DNA repair and apoptosis (reviewed in [10]). In addition, VPA has also been associated with DNA methylation, as VPA can induce downregulation of chromatin maintenance proteins but also induce direct methylation of lysine 4 on histone 3 [13, 14]. The anti-tumor activity of VPA has been observed in both solid and hematological malignancies such as thyroid cancer [15], neuroblastoma [16], glioma [17], breast cancer [18], and hematological malignancies [19-21].

Although VPA can induce histone acetylation within 30 minutes, events that regulate chromatin condensation status may take up to 48 hours to be completed. Consequently, pretreatment with VPA for 48 hours has been shown to alter chromatin structure by regulation of chromatin modulation proteins such as depletion of chromatin maintenance proteins DNA methyltransferase 1 and HP1. Hence, pretreatment with VPA for 48 hours has been shown to sensitize to cell death induced by the topoisomerase II inhibitor epirubicin in a mouse model, correlating to chromatin decondensation and increased DNA damage [13, 22, 23].

VPA is presently implicated in numerous clinical trials, both as a single agent and in association with other drugs, involving various pathologies such as mood disorders, auto-immune diseases and cancer. Interestingly, VPA has been evaluated in a sequence-specific combination with FEC100 (5-fluorouracil, epirubicin and cyclophosphamide) as a primary therapy in a phase I/II trial for locally advanced/metastatic breast cancer [24]. Results were encouraging, with no pharmacokinetic or pharmacodynamic interactions. Partial response was seen in 9 of 41 patients in phase I, and objective response in 9 of 14 patients in phase II.

In this study, we have used a cell line-based model of CHOP-resistant DLBCL to investigate the ability of VPA to sensitize diffuse large B-cell lymphoma cell lines to CHOP treatment. Our results demonstrate that VPA potentiates the cytotoxic effects of CHOP treatment by inducing apoptosis as determined by annexin V and an increased level of cleaved caspase-3. Rituximab-mediated cellular cytotoxicity is sustained in the presence of VPA. In addition, we demonstrate an increased formation of topoisomerase IIa-DNA complexes and also an increased level of γH2AX indicating higher amount of double-strand breaks (DSBs) in response to VPA. Our results support a possible novel treatment strategy of DLBCL, utilizing VPA in combination with the conventional R-CHOP protocol.

Materials and methods

Reagents

Cyclophosphamide monohydrate (C), vincristine sulfate (O), doxorubicin monohydrate (H), prednisolone (P), and valproic acid (VPA) was obtained from Sigma Aldrich (St Louis, MO). Prednisolone is the biologically active substance of prednisone. Rituximab was obtained from local pharmacy. 7-AAD was obtained from BD Pharmingen (San Diego, CA)
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Cells and culture conditions

The human diffuse large B-cell lymphoma (DLBCL) cell lines SU-DHL-5, Karpas-422, SU-DHL-8 and WSU-NHL were purchased from the German Collection of Microorganisms and Cell Cultures (DSMZ). The diffuse large B cell lymphoma cell line ULA [25] was kindly provided by Dr Berglund (Uppsala University, Uppsala, Sweden). Karpas-422, SU-DHL-5 and SU-DHL-8 was grown in RPMI 1640 (Invitrogen, Carlsbad, CA) supplemented with 20% fetal bovine serum (FBS) (Invitrogen). WSU-NHL was grown in RPMI 1640 supplemented with 10% FBS. ULA was grown in 45% Optimem (Invitrogen) and 45% IDEM (Invitrogen) supplemented with 10% FBS. All cell lines were cultured in a humidified atmosphere (37°C, 5% CO2).

Cell viability

Cells were seeded in a concentration of 0.8-1x10^6 cells/ml and treated with different combinations of substances as specified in figure legends. Cell viability was assessed after 24 h, 48 h and 72 h by trypan blue exclusion. The VPA pretreatment experiment was performed by a 24 h or a 48 h pretreatment of cells with 0.5 or 1.5 mM VPA alone or in combination with 20 µg/ml prednisolone followed by addition of CHOP. No additional prednisolone was added to cultures where prednisolone was included in the pretreatment (CHO). The CHOP regimen used consists of 10 µM cyclophosphamide monohydrate, 20 nM doxorubicin hydrochloride, 2 nM vincristine sulfate and 20 µg/ml prednisolone [25]. Viability was measured 48 h, 72 h, and 96 h after start of experiment using trypan blue exclusion.

Apoptosis analysis by flow cytometry

Labeling of cells with annexin V-PE (BD Pharmingen) was performed according to the manufacturer’s instructions. 7-AAD was added according to the manufacturer’s instructions. Apoptotic cells were defined as annexin V-positive, 7-AAD-positive and Annexin V-and 7-AAD-double positive cells.

Western blot analysis

Cells (0.8x10^6/ml) were incubated with VPA alone or in combination with CHOP. Cells were harvested after 24 h, 48 h and 72 h and washed once with PBS and resuspended in Laemml buffer (62.5 mM Tris-HCl, pH 6.8, 25% glycerol, 2% SDS, 0.01% Bromophenol Blue; Bio-Rad Laboratories, Hercules, CA). Primary antibodies used were anti-cleaved caspase-3 (Asp175)(5A1) from Cell Signaling Technology; anti-GAPDH (6C5), anti-p27 (C-19), anti-p21 (187), anti-actin (C-2) and anti-topol- alpha (3F6) from Santa Cruz Biotechnology (Santa Cruz Biotechnology, Heidelberg, Germany); anti-γH2AX(pS139)(N1-431) from BD Pharmingen; anti-acetylated histone H3 from Millipore (Millipore, Bedford, MA); anti-histone 3 (ab1791) from Abcam (Abcam, Cambridge, United Kingdom). After incubation with horseradish peroxidase (HRP)-conjugated secondary antibody, antibody binding was visualized with enhanced chemiluminescence (ECL, Biological industries, Beit, Israel) followed by detection with hyperfilm ECL (GE Healthcare, Buckinghamshire, United Kingdom).

Cell cycle analysis

0.5-1x10^6 cells was washed with PBS and fixed in 70% EtOH and stored at -20°C for 1-7 days. Labeling of cells for cell cycle analysis was performed as follows. Cells were washed and stained in propidium iodide (PI)-staining solution (50 µg/ml PI, 0.05% Triton X-100, 0.1 mg/ml Rnase A). Cells were incubated in the dark at room temperature for 1 hour, thereafter analyzed on a FACS Canto II flow cytometer (Becton Dickinson, San Jose, CA). Markers were set to determine the percentage of hypodiploid cells (sub-G0/G1), and cells in the G0/G1, S and G2/M phase of the cell cycle.

ADCC assay

One day prior to start of assay, WSU-NHL cells were labeled with PKH26 red fluorescent cell linker kit for general cell membrane labeling (Sigma Aldrich) according to the manufacturer’s instructions. Heat-inactivated FBS was used throughout the experiment and obtained by heating FBS to 56°C for 30 min followed by filtration through 0.22 µm filter. At day one, the PKH26-labelled cells were plated on a round-bottom 96-well plate at a density of 10 000 cells/well. Cells were treated with or without VPA 1.5 mM and CHOP (C: 10 µM cyclophosphamide monohydrate; H: 20 nM doxorubicin; O: 2 nM vincristine; P: 20 µg/ml prednisolone)
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followed by overnight incubation at 37°C. At day two, rituximab was added to the cells at concentrations of 0.01-10 µg/ml followed by 20 min incubation at 37°C. NK cells were isolated from peripheral blood using NK cell isolation kit from MACS (Miltenyi Biotec). Briefly, isolation of MNC from peripheral blood was performed by density gradient centrifugation with lymphoprep (Axis- Shield, Oslo, Norway) thereafter MNC was labeled according to the manufacturer’s instructions and applied to MACS columns for negative selection of the NK cell population. The purity of CD56-positive NK cells was verified by FACS. NK cells were used as effector cells and added at an effector to target cell ratio of 10:1. Cells were incubated overnight thereafter the amount of dead cells was visualized by staining with 7-AAD (BD) followed by FACS analysis. Dead target cells were identified as double-positive for PKH26 and 7-AAD and used as readout of the assay.

Band depletion assay

To monitor the amount of topoisomerase II-DNA covalent complexes, treated cells were lysed with an alkaline lysis solution (200 mM NaOH, 2 mM EDTA) for 30 min on ice followed by a neutralization of lysate with 2M HCl and 1.2 M Tris (pH 8). The lysate was mixed with 3x SDS sample buffer (150 mM Tris/HCl pH 6.8, 6 mM EDTA, 45% sucrose, 9% DSD, 10% b-mercaptoethanol, 0.03% bromophenol blue). The lysate was passed through a 23 G needle 5 times, boiled and resolved by SDS-PAGE and analyzed by western blotting.

Figure 1. VPA sensitizes DLBCL cell lines to CHOP treatment. The DLBCL cell lines ULA, WSU-NHL, Karpas-422, SU-DHL-8 and SU-DHL-5 were treated for 72 h with 0.1, 2 and 10 mM VPA in the absence or presence of CHOP. The cell and cell proliferation in millions of cells viability and cell proliferation in millions of cells was assessed after 24 h, 48 h, 72 h by trypan blue exclusion and normalized to untreated control cells at 0 h. Data are presented as mean ± SEM, n=3.
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Statistics

Significant differences were evaluated using the Student’s unpaired t-test. All tests were two-sided. An effect was considered to statistically significant at p<0.05 (*), p<0.01 (**) or p<0.001(***). Data analysis was performed with the GraphPad Prism 5.0a (GraphPad software, Inc, La Jolla, CA) or with Microsoft Excel. Data are plotted as means ± standard error of the mean (SEM).

Results

Effect of VPA on viability and proliferation of large diffuse B-cell lymphoma (DLBCL) cell lines

We have previously established a cell line-based model of CHOP refractory DLBCL [25]. Although relapsed or refractory cases of DLBCL have not shown a pronounced response to monotherapy with HDAC inhibitors like vorinostat or MGCD01103 [8], still several pre-clinical and clinical studies indicate that combination therapy with HDAC inhibitors and DNA-damaging chemotherapy could be an effective treatment [26]. To assess whether the chemo-resistance of DLBCL cells can be reversed, we treated the DLBCL cell lines Karpas-422, WSU-NHL, ULA, SU-DHL-8, SU-DHL-5 with increasing concentrations of the HDAC inhibitor VPA, alone or in combination with CHOP. The two most CHOP sensitive cell lines SU-DHL-8 and SU-DHL-5 showed highest sensitivity to VPA treatment both with VPA alone and in combination with CHOP (Figure 1). The three cell lines that are most resistant to CHOP treatment, Karpas-422, WSU-NHL and ULA showed decreased viability and proliferation in the presence of VPA at the higher concentrations of 2 mM and 10 mM (Figure 1). To conclude, the addition of VPA significantly increases CHOP-sensitivity of DLBCL cell lines.

Clinically relevant concentrations of VPA sensitize DLBCL cells to CHOP treatment

To further characterize the effects of VPA on DLBCL cell lines, we continued all experiments with the CHOP-resistant cell line WSU-NHL and the CHOP-sensitive cell line SU-DHL-8. VPA is used clinically in the treatment of epilepsy, and is well tolerated at continuous serum-concentrations up to 0.7 mM. Moreover, the maximal tolerated dose during 3-day treatment periods in combination with FEC in a phase I/II study by Münster et al, was 140 mg/kg/day, which corresponds to approximately 1.5 mM of total serum VPA (at day 3) [24]. Therefore, we continued to characterize the effects of 0.5 mM and 1.5 mM VPA alone or in combination with CHOP.
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in WSU-NHL and SU-DHL-8. VPA treatment alone at a concentration of 1.5 mM resulted in decreased viability of both WSU-NHL and SU-DHL-8 cells (Figure 2A and 2B). The presence of 0.5 mM VPA alone did not considerably affect viability, but in combination with CHOP, a sensitizing effect of VPA after 72 h could be noticed as the viability decreased to 60% for WSU-NHL and to 50% for SU-DHL-8 as compared to 85% and 65%, respectively, for CHOP alone (Figure 2A and 2B). Most striking was the additive effect of 1.5 mM VPA to CHOP, that resulted in a viability of 25% and 15% after 72 h, compared to the viability cells treated with of 1.5 mM VPA alone, that resulted in 40% and 60% viability in WSU-NHL and SU-DHL-8, respectively (Figure 2A and 2B). The proliferation of WSU-NHL and SU-DHL-8 was reduced in a dose-dependent manner in the presence of VPA (Figure 2A and 2B). Interestingly, 0.5 mM VPA initially showed a pro-proliferative effect especially in SU-DHL-8 (Figure 2B). Treatment with CHOP resulted in a proliferation arrest, which was not altered by the presence of VPA (Figure 2A and 2B). In conclusion, clinically relevant concentrations of VPA are enough for sensitizing diffuse large B cell lymphoma cells to CHOP treatment.

Pretreatment of DLBCL cell lines with VPA

An interesting clinical study has been performed, assessing the use of sequential administration of VPA and chemotherapy for patients with solid malignancies [24]. Therefore, we investigated whether pretreatment with VPA 48 h before addition of the cytotoxic combination of CHOP had the same sensitizing effect as seen for simultaneous treatment of VPA and CHOP. As seen in Tables 1 and 2, both SU-DHL-8 and WSU-NHL show significantly decreased viability for cells pretreated with 1.5 mM VPA in comparison with cells treated with VPA or CHOP alone. Taken together, sequential or simultaneous treatment of VPA and CHOP has similar effects on cell viability. Because VPA is a well-known tranquilizer, with documented sedative effects, it could be advantageous to combine it with prednisolone, which is known to have strong invigorating effects. Moreover, prednisolone is part of the CHOP regimen, and could easily be administered together with VPA without major changes in the CHOP protocol. Therefore, pretreatment with VPA and prednisolone (20 µg/ml) for 48 h was performed before the remaining cytotoxic drugs comprising CHOP i.e. cyclophosphamide, doxorubicin and vincristine (CHO) were added. Table 1 and 2 show a significant decrease in viability of WSU-NHL and SU-DHL-8 pretreated with 1.5 mM VPA and prednisolone compared to cells pretreated with prednisolone alone. In conclusion, pretreatment with VPA alone or VPA in combination with prednisolone before addition of cytotoxic drugs has a significant negative effect on the viability of DLBCL cells.

Table 1. Pretreatment of SU-DHL-8 cells with VPA in combination with prednisolone and CHOP

<table>
<thead>
<tr>
<th>SU-DHL-8</th>
<th>48 h (mean viability/SEM)</th>
<th>72 h (mean viability/SEM)</th>
<th>96 h (mean viability/SEM)</th>
<th>P-value (at 96 h compared to CHOP alone)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>100/3</td>
<td>99/2</td>
<td>98/2</td>
<td>0.057</td>
</tr>
<tr>
<td>0.5 mM VPA</td>
<td>98/3</td>
<td>100/2</td>
<td>88/5</td>
<td>0.780</td>
</tr>
<tr>
<td>1.5 mM VPA</td>
<td>80/5</td>
<td>78/5</td>
<td>69/11</td>
<td>0.262</td>
</tr>
<tr>
<td>Prednisolone (P)</td>
<td>94/7</td>
<td>100/3</td>
<td>95/3</td>
<td>0.150</td>
</tr>
<tr>
<td>CHOP</td>
<td>96/3</td>
<td>98/3</td>
<td>86/4</td>
<td>-</td>
</tr>
<tr>
<td>0.5 mM VPA + CHOP</td>
<td>98/3</td>
<td>97/1</td>
<td>76/5</td>
<td>0.175</td>
</tr>
<tr>
<td>1.5 mM VPA + CHOP</td>
<td>87/3</td>
<td>69/4</td>
<td>49/7</td>
<td>0.015*</td>
</tr>
<tr>
<td>P + CHO</td>
<td>99/0</td>
<td>89/3</td>
<td>48/4</td>
<td>-</td>
</tr>
<tr>
<td>0.5 mM VPA + P + CHO</td>
<td>93/6</td>
<td>92/8</td>
<td>70/5</td>
<td>0.028*</td>
</tr>
<tr>
<td>1.5 mM VPA + P + CHO</td>
<td>60/11</td>
<td>49/6</td>
<td>25/6</td>
<td>0.0486*</td>
</tr>
</tbody>
</table>

SU-DHL-8 cells were treated with different combinations of VPA, prednisolone (P, 20 µg/ml)) and CHOP/CHO as illustrated as in Table 2. Cell viability was assessed at 48, 72, 96 h by trypan blue exclusion. Standard error of the mean (SEM) of 3 independent experiments is shown. A two-sided t-test was performed. *P<0.05.
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VPA induces apoptosis in CHOP resistant DLBCL cells

To assess if the observed cytotoxicity after VPA treatment reflects an induction of apoptosis, WSU-NHL and SU-DHL8 cells were treated with 0.5 mM or 1.5 mM VPA alone or in combination with CHOP for 72 h, followed by FACS analysis of annexin V and 7-AAD positivity. In WSU-NHL cells, 1.5 mM VPA but not 0.5 mM VPA induced a prominent annexin V/7AAD-positivity of about 80% (Figure 3A) confirming the viability data in Figure 2A. A significant additive effect of 1.5 mM VPA to CHOP in WSU-NHL resulted in 85% viability after 72 h as judged by trypan blue exclusion (Figure 2A). Treatment with CHOP alone in WSU-NHL resulted in 85% viability after 72 h as judged by trypan blue exclusion (Figure 2A) but when analyzing annexin V and 7-AAD positivity, the initiation of an apoptotic program is demonstrated by the 40% of annexin V and 7-AAD positive cells (Figure 3A). The SU-DHL-8 cell line is most likely more responsive to VPA than WSU-NHL as judged by higher number of annexin V and 7-AAD positive cells with increasing concentration of VPA as compared to WSU-NHL (Figure 3A). An evident additive effect of VPA to CHOP is observed also in SU-DHL-8 cells (Figure 3A) as judged by the amount of apoptotic annexin V and 7-AAD positive cells.

To further confirm that the decreased viability after VPA treatment is in fact due to apoptosis, we determined the presence of cleaved caspase-3 (Figure 3B). The increased amount of cleaved caspase-3 in WSU-NHL cells treated with 1.5 mM VPA alone and in combination with CHOP (Figure 3B) is in accordance with viability data and annexin V/7AAD data, supporting the observed apoptotic effect of VPA with and without CHOP. In SU-DHL-8 cells, a strong increase in cleaved caspase-3 in the presence of CHOP is observed, consistent with the increased CHOP sensitivity of this cell lines, while the effects of VPA alone are comparable to effects in WSU-NHL cells (Figure 3B).

VPA induces G1 arrest of DLBCL cells

HDAC inhibitors are reported to rapidly induce cell cycle arrest and induce tumor cell-selective apoptosis. VPA has been reported to induce
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Inhibition of HDACs results in disruption of cellular acetylation homeostasis and can induce hyperacetylation of both histone and non-histone proteins. To verify that the concentrations of VPA used in our experiments have effects on the acetylation of histones, we investigated the acetylation status of histone 3 in WSU-NHL and SU-DHL-8 after 72 h of VPA and CHOP treatment. As seen in Figure 3B, the acetylation of histone 3 is increased in cells treated with 0.5 mM and 1.5 mM VPA both alone and in combination with CHOP.

Figure 3. VPA induces apoptosis in WSU-NHL and SU-DHL cells. WSU-NHL and SU-DHL-8 cells were treated with with 0.5 and 1.5 mM VPA alone or in combination with CHOP for 72 h. A. Apoptosis was assessed by annexin V- and 7-AAD-labeling followed by FACS analysis. Cells positive for annexin V or 7-AAD were judged as apoptotic cells. Data are presented as mean ± S.E.M., n=3 (WSU-NHL) and range, n=2 (SU-DHL-8). Statistical analysis was performed using Student’s t test. p<0,05 (*), p<0,01 (**), p<0,001 (***).

B. WSU-NHL and SU-DHL-8 cells were harvested at 72 h after addition of substances. After protein isolation, detection cleaved caspase-3, p21, acetylated histone 3 (acH3) and histone 3 (H3) was performed by western blot analysis.

C. The amount of p27 protein in WSU-NHL and SU-DHL-8 was investigated by western blotting 48 h after addition of substances. H3 and GAPDH were used as equal loading controls. Non-specific bands are marked n.s. Data are representative of at least two independent experiments.
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VPA increases the level of covalent topoisomerase IIα-DNA complexes and potentiates CHOP-induced DSBs

As shown in Figure 3B, treatment with VPA leads to histone acetylation. It has previously been shown that this is followed by a modulation of genes and proteins essential for the maintenance of heterochromatin, resulting in chromatin decondensation [13]. Topoisomerase II (Topo II) is an enzyme that regulates DNA under- and overwinding. In order to carry out its critical physiological functions, topoisomerase II generates transient double-stranded breaks in DNA, binding covalently to the DNA [27]. Chromatin decondensation induced by HDACis is associated with an increased binding of topoisomerase II to the DNA substrate and in the presence of topoisomerase II inhibitors, it is also associated with increased DNA damage and cell death [13, 28]. Doxorubicin is a Topo II inhibitor that forms covalent complexes with Topo II–induced double-stranded DNA breaks (DSBs). We used a band depletion assay to control for possible differences in the extent of doxorubicin-induced Topo IIα-trapping in VPA treated cells compared to non-treated cells. Cells were treated with or without 1 mM VPA for 24 h followed by an incubation of CHOP for 24 hours. Thereafter the band depletion assay was performed to analyze the amount of Topo IIα that was trapped in covalent complexes with DNA and therefore unable to enter the gel. Hence, the proportion of Topo IIα trapped in cleavage complexes is proportional to the reduction in Topo IIα as assessed by western blotting. Both WSU-NHL and SU-DHL-8 cells treated with VPA alone and with VPA in combination with CHOP had reduced levels of Topo IIα compared to untreated or CHOP-treated cells, indicating that a higher amount of Topo IIα was trapped in DNA complexes in response to VPA (Figure 5B). Interestingly, treatment with VPA alone resulted in Topo IIα-trapping to a comparable extent to cells treated with a combination of VPA and CHOP. This may suggest that VPA-mediated HDAC inhibition alone leads to increased binding of Topo IIα to the DNA, even in the absence of topoisomerase II inhibitors like doxorubicin. In conclusion, our data support that VPA may increase the DNA binding of Topo IIα, which may contribute to sensitization to CHOP-treatment.

Figure 4. VPA treatment alone induces G1 arrest of WSU-NHL and SU-DHL-8 cells, and in combination with CHOP, VPA increases the number of cells in sub-G1/G0. A. WSU-NHL and B. SU-DHL-8 cells were treated with 0.5 or 1.5 mM VPA alone or in combination with CHOP for 72 h. Cells were harvested after 24 h, 48 h and 72 h and cell cycle analysis was performed using propidium iodide labeling. Error bars represent S.E.M., n=3.

Pretreatment with VPA increases the amount of DNA double-strand breaks in CHOP treated cells

Since VPA can increase the amount of DNA cleavage complexes with Topo IIα, we investigated if the presence of VPA also can compromise the repair of DNA double-strand breaks (DSBs). The accumulation of γH2AX (histone H2AX phosphorylated on S139), is an early marker of DNA DSBs. The level of γH2AX is proportional to the level of free DSBs [29]. Hence,
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we measured the levels of γH2AX as a marker to monitor the generation and repair of VPA-induced DSBs. As a positive control we used VM-16, also called etoposide, a drug reported to induce cell death with the preferential formation of Topo IIα-containing cleavable complexes. In SU-DHL-8 cells, treatment with both CHOP alone and 1 mM VPA alone resulted in an increased amount of γH2AX, and pretreatment with 1 mM VPA before addition of CHOP resulted in an even higher amount of γH2AX (Figure 5A). Interestingly, in WSU-NHL cells treated with 1 mM VPA and cells pretreated with 1 mM VPA for 24 h before addition of CHOP, an increased level of γH2AX was detected by western blot analysis as compared to cells treated with CHOP alone (Figure 5A). Taken together, our data suggest that pretreatment with VPA does indeed potentiate CHOP-induced DNA damage in DLBCL cell lines.

VPA does not counteract rituximab-induced cellular cytotoxicity

The standard therapy for patients diagnosed with DLBCL is CHOP in combination with the monoclonal anti-CD20-antibody rituximab. Since the addition of rituximab to CHOP therapy the overall survival has increased for DLBCL patients [30]. Therefore, it is of great importance that future additional drugs in the treatment regimen of these patients do not impede the function of the antibody-based therapy. Indeed, Shimizu et al recently reported that HDAC inhibitors including VPA increase CD20 expression on the cell surface, and augment rituximab-mediated complement dependent cytotoxicity (CDC) in lymphoma cell lines [31]. However, as the major part of the cytotoxic effect of rituximab is mediated by antibody dependent cellular cytotoxicity (ADCC) [32], we investigated the effect of VPA on rituximab-
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mediated ADCC. To that purpose, SU-DHL-8 and WSU-NHL cells were treated with 1.5 mM VPA alone or in combination with CHOP for 24 h followed by an incubation of cells with increasing concentration of rituximab. Thereafter NK cells from peripheral blood was added as effector to target cell ratio of 10:1 thereafter the cells were incubated for an additional 16 hours. Dead target cells were identified as double positive for PKH26 and 7-AAD and used as readout of the assay. Error bars represent S.E.M., n=3.

Figure 6. VPA does not interfere with rituximab-mediated cellular cytotoxicity. A. WSU-NHL and B. SU-DHL-8 cells were labeled with PKH26, either left untreated or incubated with 1.5 mM VPA for 24 h followed by addition of varying concentrations of rituximab. NK cells were added at an effector to target cell ratio of 10:1 thereafter the cells were incubated for an additional 16 hours. Dead target cells were identified as double positive for PKH26 and 7-AAD and used as readout of the assay. Error bars represent S.E.M., n=3.

Discussion

Anthracyclin-based therapy with CHOP has been the unchallenged treatment for diffuse large B-cell lymphoma since more than thirty years. Although addition of the monoclonal CD20 antibody rituximab (R-CHOP) during the last decade has led to improved overall and progression free survival, the prognosis for patients diagnosed with DLBCL is still far from satisfying, as approximately 40% of affected patients ultimately die from their disease.

Therefore, the clinical need of improved R-CHOP-based therapy is urgent.

Histone deacetylase inhibitors (HDACis) are a new class of therapeutic agents gaining growing attention within cancer treatment. Through modification of the structural components of chromatin, HDACis regulate expression of tumour suppressor genes and activities of transcription factors involved in cancer initiation and progression, and are also able to regulate chromatin condensation. Along these lines, several in vitro studies have suggested that HDACis can synergize with chemotherapy [26]. HDACis, at concentrations required for chromatin remodeling, have been shown to potentiate DNA damage induced by anthracyclins such as doxorubicin and epirubicin in breast cancer cell lines and in mouse models [22, 23]. Moreover, it has been suggested that pretreatment with HDACis for 48 hours before treatment with anthracyclins may give time for chromatin remodelling, resulting in maximal DNA damage [23].

In this study, we show that HDACis like valproic acid can be valuable in combination also with R-CHOP in diffuse large B-cell lymphoma cell lines. Both pretreatment and concomitant treatment with VPA sensitizes strongly to CHOP treatment at clinically relevant concentrations. Importantly, the sensitizing effect of VPA to chemotherapy was prominent also in cell lines resistant to treatment with CHOP, such as WSU-NHL. In contrast to its effects in breast cancer cell lines [23], VPA had pronounced effects in lymphoma cell lines also as a single agent.

We demonstrate anti-proliferative and pro-apoptotic activity of VPA as a single agent on DLBCL cell lines. VPA has previously been suggested to induce a G0/G1 cell cycle arrest by an
increased expression of p21 and p27 [33]. Our data demonstrate a dose-dependent G0/G1 arrest after 24 h of treatment, corresponding to an increased expression of p21 and p27. It could be argued that this cell cycle arrest should have protective effects to CHOP treatment. However, our data showing increased apoptosis both after treatment with VPA alone and with a combination of VPA and CHOP, suggest that the apoptotic response overrides a possible effect on cell cycle arrest. Interestingly, also levels of γH2AX and of Topo IIα cleavage complexes, indicative of DNA DSBs, increase after combination treatment with VPA and CHOP and also after VPA alone. This suggests that VPA increases DNA damage, which could contribute to the apoptotic response. Hence, VPA, both alone and in combination with CHOP, has pronounced effects on viability of lymphoma cells, which supports its use in clinical lymphoma treatment. In addition, combination of VPA with prednisone further increased cytotoxicity. In the light of the sedative effects of VPA, this could be a clinically relevant finding, given the well-established invigorating effects of prednisone.

A possible addition of histone deacetylase inhibitors to conventional R-CHOP therapy is dependent on the sustained effect of rituximab. VPA treatment has been reported to increase the mRNA and protein level of CD20 on B-cell lymphoma cell lines [31], resulting in increased CDC. However, in contrast to the stimulatory effects on CDC, we see no effects of VPA on the rituximab-mediated ADCC of on WSU-NHL and SU-DHL-8 cells after a 24 h incubation time. This is in agreement with data by van Meerten et al showing no clear correlation between the level of CD20 expression and Rituximab-induced ADCC [34]. Still, our data indicate that VPA does not negatively affect rituximab-mediated ADCC, and support that VPA could be combined with R-CHOP in a clinical setting.

Compared to newer “second generation” HDACis such as belinostat and romidepsin, VPA is a rather weak HDACi, with HDAC inhibitory activity in the millimolar range. Still, several clinical trials have reported that its utility in cancer treatment is dependent on its HDAC inhibitory activity [35, 36], and its synergistic effects together with epirubicin correlate to its histone acetylating capacity [23]. In the present study, the ability of VPA to inhibit HDACs, was verified by the increased expression of acetylated histone 3 after 24 h treatment with VPA at 1 mM, suggesting that VPA treatment had the desired effects.

Taken together, VPA in clinically relevant concentrations potentiates CHOP-induced apoptosis of lymphoma cell lines. Moreover, VPA does not interfere with rituximab-induced ADCC. Our results support VPA as a potential pretreatment therapy before R-CHOP in diffuse large B-cell lymphoma. Indeed, this concept is presently being evaluated in VALFRID, a clinical phase I trial at the Skåne Clinic of Oncology in Sweden (ClinicalTrials.gov ID NCT01622439).

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Address correspondence to: Dr. Kristina Drott, Department of Hematology and Transfusion Medicine, Lund University, BMC B13, Klinikg. 26, S-221 84 Lund, Sweden. Phone: +46-46-2220632; Fax: +46-46-184493; E-mail: Kristina.Drott@med.lu.se

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