



Antiviral activity of brincidofovir on parvovirus B19

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ABSTRACT

Parvovirus B19 (B19V), a single-stranded DNA virus in the family *Parvoviridae*, is a human pathogenic virus responsible for a wide range of clinical manifestations. Currently there is no approved antiviral therapy for parvovirus infection. The acyclic nucleoside phosphonate cidofovir (CDV) has been demonstrated to inhibit replication of B19V *in vitro*. The aim of the present study was to evaluate whether brincidofovir (BCV), a novel lipid conjugate of CDV, could also inhibit B19V replication. Experiments were carried out in erythroid progenitor cells (EPCs) and UT7/EpoS1 cells, infected with B19V and cultured in the presence of different concentrations of BCV and CDV for comparison. The dynamics of viral replication was evaluated by a qPCR-based assay and the extent of inhibition of viral replication exerted by the compounds determined, along with the effect of the compounds on cell viability and cell proliferation rates. Results confirmed that BCV showed significantly higher antiviral activity against B19V compared to CDV in both cell-based systems. For BCV, the calculated EC₅₀ values were in the range 6.6–14.3 μM in EPCs and 0.22–0.63 μM in UT7/EpoS1 cells. In comparison, the EC₅₀ values for CDV were > 300 μM in EPCs and 16.1 μM in UT7/EpoS1 cells. Concurrently, the effects on cell viability were observed at a much higher concentration of BCV, with calculated CC₅₀ values in the range 93.4–102.9 μM in EPCs and 59.9–66.8 μM in UT7/EpoS1. The antiviral activity was observed specifically with the metabolically active stereoisomer of BCV suggesting that CDV-diphosphate, the metabolite of both BCV and CDV, was the active antiviral. Our results support a selective role for BCV in the inhibition of B19 viral replication.

1. Introduction

Parvovirus B19 (B19V) is a single-stranded DNA virus in the family *Parvoviridae*, still in need of a specific antiviral (Gallinella, 2013; Qiu et al., 2017). The virus has a small genome, a 5.6 kb ssDNA molecule, with a compact organization and a relatively limited coding repertoire, comprising a non-structural protein, NS, whose function is essential for virus replication, and two structural proteins, VP1 and VP2, that form a T = 1, 22 nm icosahedral capsid. A pathogenic human virus, B19V is characterized by a selective but not exclusive tropism for erythroid progenitor cells (EPCs) in the bone marrow, and by a strict dependence on the cellular machinery and environment for its replication, so that the target cell types, their differentiation stage and proliferation rate are all critical to the diverse outcomes of infection. The virus is capable of long-term persistence in disparate tissues (Bua and Gallinella, 2017), and can establish a complex relationship with the immune system, whose efficacy in innate and adaptive responses can be critical to the course of infection and the development of pathological processes.

In the general population, infection is widespread and can be associated with a range of pathologies and clinical manifestations, from asymptomatic or mild, to severe and in some cases life-threatening, whose characteristics and outcomes depend on the interplay between the viral properties and the physiological and immune status of the infected individuals (Gallinella, 2013; Qiu et al., 2017). Cells of the erythropoietic lineage in the bone marrow constitute a primary target, but a large spectrum of secondary target tissues and organs can be involved (Adamson-Small et al., 2014). The selective tropism for erythroid progenitor cells in the bone marrow can cause a partial block in erythropoiesis, that may manifest as transient or persistent erythroid aplasia (Kerr, 2015). Common systemic manifestations of infection are erythema infectiosum in children and post-infection arthropathies mainly affecting adults; moreover, the virus has been implicated in a growing collection of other different pathologies, among them myocarditis (Verdonschot et al., 2016), connective tissue diseases and autoimmune processes (Kerr, 2016). Notably, infection in pregnancy may be transmitted to the fetus, posing a risk of fetal death, fetal hydrops

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and possible adverse perinatal outcome (Bonvicini et al., 2017a; Bascietto et al., 2018).

The gap in the development of antiviral strategies directed against B19V is striking (De Clercq and Li, 2016). A vaccine against B19V is technically feasible, but still under development (Chandramouli et al., 2013; Penkert et al., 2017), and no specific antiviral therapy has been investigated or clinically evaluated for B19V. Currently, treatment options are limited and generally supportive. Blood transfusions are required to overcome acute or chronic anemia, while intravenous immunoglobulin (IVIG) is considered the only available option to neutralize infectious virus and to control infection in cases of individuals with an impaired immune system (Mouthon and Lortholary, 2003; Crabol et al., 2013; Mouthon et al., 2015). However, the beneficial effects of supportive treatments are limited, and even IVIG treatments are seldom able to clear infection unless a patient's own antiviral immune response develops and becomes effective.

An effective antiviral would have profound implications in the treatment of hematological complications encountered during the acute or chronic phase of the infection, especially in subjects with stressed erythropoiesis or with immune system deficits, or to reduce the inflammatory or systemic aspects of infections in atypical cases, or potentially could be used for prophylaxis in selected cases.

The incomplete characterization of the viral proteome and of the molecular mechanisms involved in viral replication have hampered the rational design of specifically targeted drugs, while the demanding *in vitro* cell culture conditions have limited the feasibility of a high-throughput screen against available chemical libraries. To overcome these barriers, our group previously took two alternative approaches, based on a drug repositioning strategy, and on investigating known antiviral compounds for a possible activity against B19V. The first approach yielded antiviral activity provided by the cell-proliferation inhibitor hydroxyurea (HU) (Bonvicini et al., 2017b), and the second approach yielded the acyclic nucleoside phosphonate cidofovir (CDV) though with suboptimal activity (Bonvicini et al., 2015, 2016). The aim of the present research project was to evaluate whether brincidofovir (BCV), a novel lipid conjugate of CDV possessing enhanced antiviral activity against dsDNA viruses (Aldern et al., 2003; Williams-Aziz et al., 2005; Hostetler, 2009), also exerts an inhibitory effect against B19V replication, and in particular whether it might perform more favorably compared to CDV.

Two *in vitro* cellular systems established for B19V were used for this study. *In vitro* derived EPCs constitute a cellular population analogous to the primary target cells in the bone marrow and present full permissiveness to viral replication depending on differentiation stage and proliferation rate (Wong et al., 2008; Bua et al., 2016); the human myeloblastoid cell line UT7/EpoS1 presents a restricted pattern of permissiveness (Bonvicini et al., 2006, 2008), allowing viral replication along with a limited yield of infectious virus production. In our study, both the EPC and UT7/EpoS1 cells were infected with B19V and then cultured for a single round of infection in the presence of test compounds, BCV and CDV for comparison. The dynamics of viral replication was evaluated by a qPCR-based assay, and the extent of inhibition of viral replication exerted by the compounds determined, along with the extent of inhibition on cell viability and cell proliferations rates. Experimental evidence indicated that BCV can be effective in inhibiting B19V replication, and demonstrated an enhanced inhibitory activity compared to CDV.

2. Materials and methods

2.1. Chemicals

Brincidofovir (BCV) was provided by Chimerix (Chimerix, Inc., Durham, USA). Unless stated, BCV refers to the S-enantiomer (CMX001) as opposed to the R-enantiomer (CMX029). Cidofovir (CDV) was purchased from Sigma at a purity of > 98%. In all cases, dry powder was

resuspended in phosphate buffered saline (PBS), pH 11.6, at 10 mM, then aliquots of the stock solution were stored at -70°C until use.

2.2. Cells

Erythroid progenitor cells (EPCs) were generated *in vitro* from peripheral blood mononuclear cells (PBMC), obtained from the leukocyte-enriched buffy coats of anonymous blood donors available for institutional research purposes from the Immunohematology and Transfusion Service, S. Orsola-Malpighi University Hospital, Bologna (<http://www.aosp.bo.it/content/immunoematologia-e-trasfusionale>; authorization 0070755/1980/2014, issued by Head of Service). Availability was granted under conditions complying with Italian privacy law. Neither specific ethics committee approval nor written consent from donors was required for this research project. PBMC, isolated by centrifugation in Ficoll-Paque Plus (GE Healthcare Bio-Sciences AB), were cultured in IMDM (Gibco) supplemented with 20% serum substitute BIT 9500 (StemCell Technologies), and enriched with erythropoietic growth factors as described (Bua et al., 2016; Filippone et al., 2010). The cells were maintained at 37°C in 5% CO_2 and used for infection experiments at day 9 ± 1 , when permissiveness to B19V infection is maximal. UT7/EpoS1 cells were cultured in IMDM, supplemented with 10% FCS and 2 U/mL rhu erythropoietin, at 37°C and 5% CO_2 (Wong and Brown, 2006).

2.3. Infection

A B19V viremic serum sample, identified in our laboratory in the course of institutional diagnostic service and available for research purposes according to Italian privacy law, was used as source of virus for the infection experiments. The viremic serum contained 10^{12} B19V (genotype 1) genome copies (geq)/mL, as determined by quantitative PCR analysis (Gallinella et al., 2004; Bonvicini et al., 2013a), and tested negative by routine diagnostic assays to other viruses, including HIV, HBV, HCV, HSV, VZV, EBV, CMV, HHV8, AdV, BKV.

For infection, cells were incubated in PBS at a density of 10^7 cell/mL, in the presence of the B19V viremic serum, diluted in PBS in order to obtain the desired multiplicity of infection (moi, expressed as geq/cell). EPCs were infected with different moi, from 10^1 geq/cell to 10^4 geq/cell, while UT7/EpoS1 were infected with at the highest moi of 10^4 geq/cell. Following 2 h at 37°C , the inoculum virus was washed twice in PBS and the cells were incubated at 37°C in 5% CO_2 in the respective growth medium, at the different concentrations of tested compounds, at an initial density of 10^6 cells/mL.

2.4. Nucleic acids purification

Equal amounts of cell cultures, corresponding to 1.5×10^5 cells, were collected as appropriate at 2 or 48 h post-infection (hpi) and processed by using the Maxwell Viral Total Nucleic Acid kit on a Maxwell MDx platform (Promega), following the manufacturer's instructions, in order to obtain a total nucleic acid fraction in elution volumes of 150 μL .

2.5. Quantitative real-time PCR

For quantitative analysis of viral DNA, an aliquot of the eluted nucleic acids, corresponding to ~ 500 cells, was amplified by qPCR using the primer pair R2210-R2355, located in the central exon of B19V genome (Bua et al., 2016; Bonvicini et al., 2006, 2008), and quantified using an external calibration curve, in a RotorQ system (Qiagen). As a control, a target sequence in the region of genomic DNA coding for 5.8S rRNA (rDNA) was amplified in parallel reactions. Amplification reactions were performed by using Maxima SYBR Green qPCR Master Mix (Thermo Scientific), including primers (obtained from Eurofins Genomics) at a final concentrations of 0.3 μM . Quantitation of viral DNA was

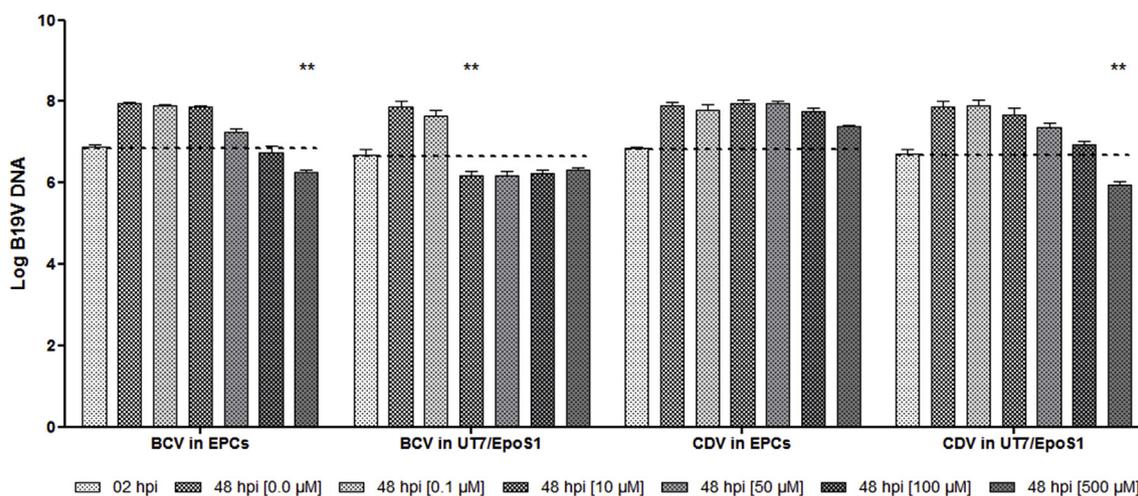


Fig. 1. Inhibition of B19V by BCV and CDV, in EPCs and UT7/EpoS1 cells. The amounts of B19V DNA were determined by qPCR, at 2 hpi and 48 hpi, at the different BCV or CDV concentrations, and expressed as Log B19V DNA geq/1000 cells. Histograms indicate mean values \pm standard error of means obtained from 3 (for EPCs) or 5 (for UT7/EpoS1) independent experiments. Analysis of variances were followed by Bonferroni post-comparison test to indicate a significant reduction in B19V replication at 48 hpi compared to the respective 2 hpi samples (dashed lines, ** $p < 0.01$).

obtained by the absolute quantitation algorithm, converting quantification cycle (Cq) values to geq number using external calibration curves obtained from standard targets. For the rDNA target, a coefficient of variation of quantification cycle values for the different samples $\leq 5\%$ was required, so that this parameter could be considered invariant and the normalization by rDNA not necessary for the following quantitative determination of viral targets.

2.6. Fluorescent in situ hybridization (FISH) assay

Aliquots of cell cultures were processed by on slide FISH assay for the detection of viral nucleic acids, as described (Manaresi et al., 2015) with minor modifications. Cells were spotted on glass slides, fixed in PBS-paraformaldehyde 0.5% at 4 °C for 30 min and permeabilized for 45 min in PBS-saponin 0.2%. Hybridization reaction was carried out in 25 μ l of a hybridization solution containing 25 ng of a digoxigenin labelled, random-primed full-length genomic probe (Dig High Prime, Roche). Specimens and hybridization mixture were denatured together by heating at 95 °C for 5 min and then incubated at 37 °C for 12 h. Following hybridization, the slides were washed twice at 37 °C with 50% formamide - 2 \times SSC (300 mM NaCl, 30 mM sodium citrate, pH 7.0) buffer and twice at room temperature in 2 \times SSC buffer, 10 min each. Detection of the hybrids was performed with a FITC-conjugate anti-digoxigenin antibody (Roche) diluted 1:20 in PBS-BSA 1% and incubated for 1 h; after washing in PBS, slides were stained with Evans blue and observed on a fluorescence microscopy (EX 450–490 nm, BA 520 nm).

2.7. Cell viability and proliferation assays

The effects of tested compounds on cell viability and proliferation were monitored by a WST-8 based assay, a water-soluble salt reagent that produces a formazan dye upon reduction in response to metabolic activity (CCK8 assay, Dojindo Molecular Technologies), and by evaluation of 5-bromo-2'-deoxyuridine (BrdU) incorporation into newly synthesized DNA (Cell proliferation ELISA BrdU Assay, Roche Diagnostics), following established and optimized protocols (Bonvicini et al., 2017b). For experiments, 5×10^4 cells were seeded in 100 μ l volumes in a 96-well culture microplate, and cultured in the absence, as control, or in the presence of different concentration of compounds added to medium for 48 h, paralleling the time course of infection. For the cell viability assay, WST-8 reagent was added for the last 6 h for EPCs, or 2 h for UT7/EpoS1. For the cell proliferation assay, BrdU at

10 μ M was added for the last 24 h for both cell types. The amounts of formazan dye and of incorporated BrdU were measured as absorbance (OD) values according to manufacturers' instructions. Replicate net OD values were normalized with respect to the control samples and expressed as mean percentage values for, respectively, cell viability and proliferation.

2.8. Statistical analysis

Experiments were carried out in duplicate (or more) series unless stated and, for each sample, quantitative determinations were carried out in duplicate (or more) values. Statistical analysis was carried out using GraphPad Prism version 5.00 for Windows (GraphPad Software, San Diego California, USA). Two-way ANOVA (analysis of variance) followed by the Bonferroni test was used to compare data obtained in different experimental conditions. Linear regression analysis was carried out to correlate input moi values and calculated amounts of viral nucleic acids. EC_{50} and CC_{50} values (mean and 95% C.I.) for the different experimental series were determined by non-linear regression analysis on percentage residual viral replicative activity and/or cell viability or proliferation, for each different tested concentration of compounds relative to control samples.

3. Results

3.1. Effect of BCV and CDV on B19V replication

As a first set of experiments, a comparative investigation of the ability of BCV as compared to CDV to inhibit B19V replication in both EPCs and UT7/EpoS1 cell systems was carried out. PBMC derived, *in vitro* differentiated EPCs and UT7/EpoS1 cells were infected with B19V at a moi of 10^4 geq/cell. Following incubation to allow attachment and penetration of virus (2 hpi samples), infected cells were divided and cultured in the absence, as a control, and in the presence of different concentrations of BCV or CDV, in the range 0.1–500 μ M, for a complete course of infection (48 hpi samples). Equivalent cell fractions were collected at 2 hpi and at 48 hpi for the different conditions, and quantitative determination of viral DNA to determine viral replicative activity, and of cellular rDNA for control and normalization, was carried out by qPCR.

The extent of viral replication, measured as the variation in viral DNA detected after a course of infection, varied in function of the cell type, the compound, and the concentration of compounds added to

infected cell cultures (Fig. 1). By 2-way ANOVA, all variables as well as their interaction contributed significantly as a source of variation to the observed effects (at $p < 0.001$), as the cell/compound combination accounted for 18.7% of total variance, compound concentration for 39.2%, and interaction of variables for 20.2%. The amount of DNA measured from 2 to 48 hpi in control samples increased on average 1.1 ± 0.1 Log, indicating productive viral replication. For all cell/compound combination series, increasing the concentration of compounds progressively decreased the amount of viral DNA detected at 48 hpi, thus indicating an inhibitory effect of the compounds on the extent of viral replication. Complete inhibition, as indicated by a significant (at $p < 0.01$) net reduction in the amount of viral DNA from 2 hpi to 48 hpi, could be observed: for BCV, -0.6 Log at $\geq 500 \mu\text{M}$ in EPCs and -0.5 Log at $\geq 10 \mu\text{M}$ in UT7/EpoS1; for CDV, not in the tested range in EPCs and -0.7 Log at $\geq 500 \mu\text{M}$ in UT7/EpoS1. At 48 hpi, the observed variations compared to the respective control samples were: for BCV, -1.7 Log at $\geq 500 \mu\text{M}$ in EPCs and -1.7 Log at $\geq 10 \mu\text{M}$ in UT7/EpoS1; for CDV, -0.5 Log at $\geq 500 \mu\text{M}$ in EPCs and -0.7 Log at $\geq 500 \mu\text{M}$ in UT7/EpoS1.

The inhibitory effects of BCV and CDV on B19V replication were also assessed by FISH analysis (Fig. 2). FISH-positive cells, corresponding to permissive cells actually supporting B19V replication within a restrictive cell population, were evident as clearly stained above a low intensity Evans blue counterstain, and showed a localized, mainly nuclear, distribution of target. Although qualitative, this result is consistent with qPCR assay data (Fig. 1), confirming that productively infected cells could not be detected where BCV or CDV concentrations were sufficient to completely inhibit B19V replication.

Thus, experimental data obtained from both qPCR and FISH assays confirmed that BCV is effective against B19V, and proved that BCV was

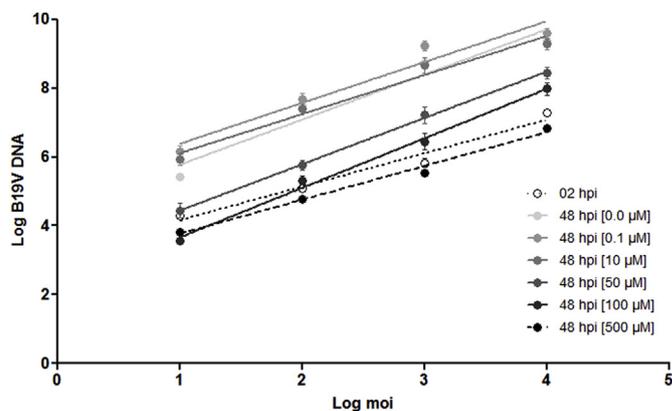


Fig. 3. B19V DNA in EPCs, as a function of moi and BCV concentration. Amounts of viral DNA (Log B19V DNA geq/1000 cells) plotted as a function of the Log of moi used (10^1 – 10^4), for samples series collected at 2 hpi and at 48 hpi for all the different concentrations of BCV added to infected EPCs cell cultures. Values are mean of duplicate independent experiments and bars indicate the standard error of means. Lines are linear regression analysis for the respective sample series (R^2 0.91–0.98).

a much more potent inhibitor of B19V replication in both EPCs and UT7/EpoS1 cell systems. In particular, in EPCs, only BCV but not CDV led to a complete inhibition of viral replication, although only at the highest concentration tested.

3.2. Enhanced antiviral activity of BCV in EPCs

To further characterize the antiviral activity of BCV, *in vitro*

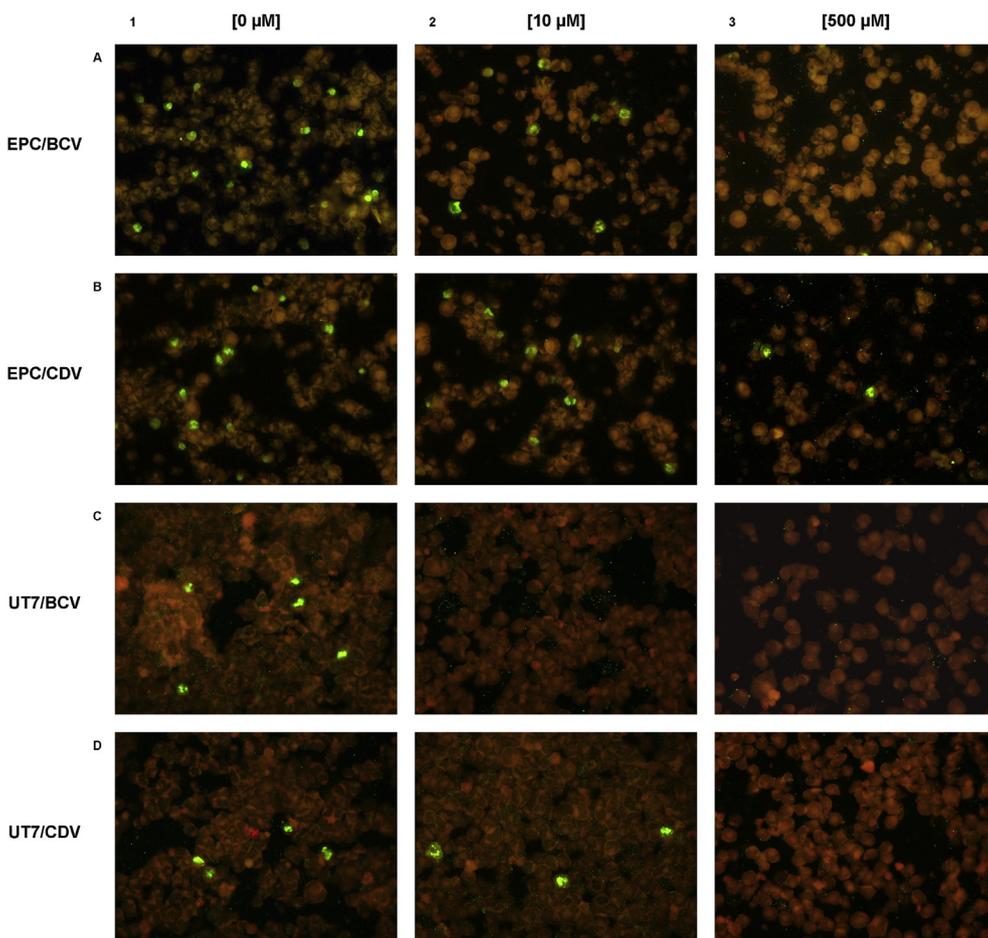


Fig. 2. FISH assay for the detection of viral nucleic acids. EPCs (A, B) and UT7/EpoS1 (C, D) were infected with B19V at the moi of 10^4 geq/cell, then cultured for a 48 h time course of infection in the presence of BCV (A, C) or CDV (B, D) at 0.0 (as control), 10, 500 μM (columns 1, 2, 3). Cells were analyzed by FISH assay for the detection of B19V nucleic acids. A whole length genomic probe for B19V, digoxigenin-labelled, was used and hybrids detected by anti-DIG Fab, FITC conjugated. Where present, positive cells are clearly stained above a low intensity background and show a localized, mainly nuclear, distribution of target, indicating productively infected cells.

differentiated EPCs were infected with B19V at different multiplicities of infection (moi), in the range of 10^1 - 10^4 geq/cell, and cultured in the absence and presence of different concentrations of BCV (0.1–500 μM). The extent of viral replication in the different experimental conditions was then assessed by qPCR analysis according to the previous experimental scheme.

The amount of viral DNA detected after a course of infection varied as a function of both the moi used and the concentration of BCV added to infected cell cultures. Amounts of viral DNA were plotted as a function of the moi used, for samples series collected at 2 hpi and at 48 hpi for all the different concentrations of BCV added to infected cell cultures (Fig. 3). For any given moi and BCV concentration, the extent of viral replicative activity resulted in a different increase in the amount of viral DNA from 2 hpi to 48 hpi, excepting for the highest BCV concentration. Linear regression analysis showed a good correlation between calculated amounts of viral DNA and the moi used, for all sample series in the whole test range (R^2 in the range 0.91–0.98). The linear correlation evident for the 2 hpi sample series indicated that virus attachment and penetration was driven by concentration of virus. For the different 48 hpi sample series, with BCV at concentrations ≤ 100 μM , the slopes of regression curves were statistically equivalent ($p = 0.12$), thus indicating a constant relative biological activity of B19V within the EPC cell system independent of the moi used. The shift observed in the regression curves showed statistically significant differences ($p < 0.0001$), with decrements that might be attributed to the progressive inhibitory effects of BCV on viral DNA replication. For the 48 hpi sample series, with BCV at 500 μM , the slope of the regression curve was different from the other 48 hpi samples and similar to the 2 hpi sample series ($p = 0.88$), and in addition the shift of the curve was lower ($p < 0.0001$) than the 2 hpi sample series. This suggests both a block in cell proliferation and a complete inhibition of viral replication at this high BCV concentration.

Based on these linear relationships, the inhibitory effect of BCV on B19V replication for any given moi could be expressed and quantified as a dose-dependent reduction in the amount of viral DNA detected at 48 hpi. By expressing a normalized dose-dependent relationship between BCV concentration and percentage replicative activity with respect to the control (Fig. 4), non-linear regression curve allowed determining EC_{50} values for BCV for all the different moi tested. Values in this experiment ranged between 6.6 and 11.5 μM for the different moi used, yielding a value of 9.35 μM [C.I. 8.84–9.89, R^2 0.99] for the pooled data. Activity of BCV and CDV were compared at a moi of 10^4 geq/cell; the calculated EC_{50} values were: for BCV, 9.65 μM [C.I.

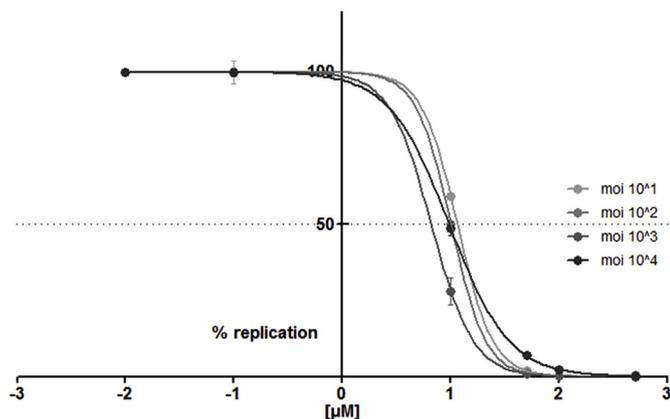


Fig. 4. Viral replication, normalized response curve to BCV in B19V-infected EPCs. Normalized dose-response curve, calculated from qPCR data obtained at 48 hpi for the different experimental series (B19V in EPCs, moi 10^1 – 10^4). Log B19V DNA as a function of BCV concentration, expressed as percentage replication compared to control samples. EC_{50} values derived from this plot are reported in the text.

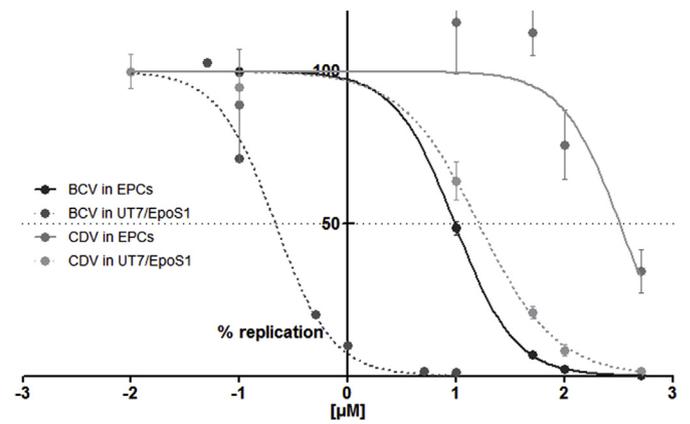


Fig. 5. Viral replication, normalized response to BCV and CDV in EPCs and UT7/EpoS1. Normalized dose-response curve, calculated from qPCR data obtained at 48 hpi for the different experimental series (B19V in EPCs and UT7/EpoS1). Log B19V DNA as a function of BCV or CDV concentration, expressed as percentage replication compared to control samples. EC_{50} values derived from this plot are reported in the text.

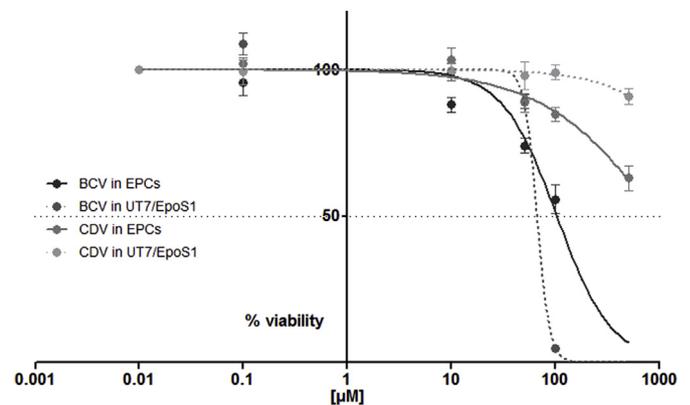


Fig. 6. Cell viability, normalized response to BCV and CDV in EPCs and UT7/EpoS1. Normalized dose-response curve, calculated from CCK8 assay data obtained at 48 hpi for the different experimental series (B19V in EPCs and UT7/EpoS1). Cell viability as a function of BCV or CDV concentration, expressed as percentage value compared to control samples. CC_{50} values derived from this plot are reported in the text.

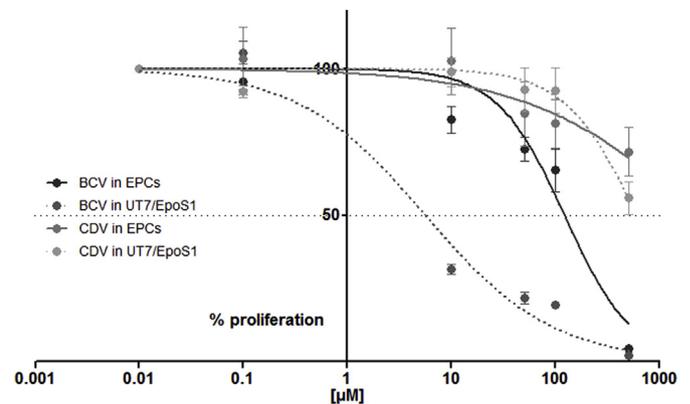


Fig. 7. Cell proliferation, normalized response to BCV and CDV in EPCs and UT7/EpoS1. Normalized dose-response curve, calculated from BrdU incorporation assay data obtained at 48 hpi for the different experimental series (B19V in EPCs and UT7/EpoS1). Cell proliferation as a function of BCV or CDV concentration, expressed as percentage value compared to control samples. CC_{50} values derived from this plot are reported in the text.

9.15–10.20, R^2 0.99]; for CDV, 320.5 μM [C.I. 173.9–590.7, R^2 0.41] (Fig. 5).

In parallel, non-linear regression was performed to determine CC_{50} values for cell viability (Fig. 6) and proliferation (Fig. 7). Regarding cell viability, calculated CC_{50} values were: for BCV, 102.9 μM [C.I. 87.6–120.8, R^2 0.93]; for CDV, > 500 μM . Regarding cell proliferation, calculated CC_{50} values were: for BCV, 121.6 μM [C.I. 91.7–161.3, R^2 0.89]; for CDV, > 500 μM . Comparing antiviral effects (data from Fig. 4) and effects on cell viability and cell proliferation (data from Figs. 5 and 6), BCV yielded selectivity indexes (SI) values of 11.0 and 13.0 for viability and cell proliferation respectively.

3.3. Enhanced antiviral activity of BCV in UT7/EpoS1

The inhibition of B19V replication by BCV in UT7/EpoS1 cells was more pronounced than in EPCs. For an accurate characterization of the antiviral activity of BCV under conditions of high viral load, UT7/EpoS1 cells were infected with B19V at the moi of 10^4 geq/cell in the presence of 0.01–10 μM BCV, or 0.1–500 μM CDV under the same experimental scheme that were used for EPCs. A dose-dependent progressive reduction in the amount of viral DNA detected at 48 hpi samples could be observed for both BCV and CDV. Non-linear regression curve analysis allowed determining EC_{50} value of 0.22 μM [C.I. 0.19–0.25, R^2 0.99] for BCV, and 16.1 μM [C.I. 12.9–20.2, R^2 0.95] for CDV (Fig. 5).

In parallel, non-linear regression was performed to determine CC_{50} values for cell viability (Fig. 6) and proliferation (Fig. 7). Regarding cell viability, calculated CC_{50} values were: for BCV, 66.8 μM [C.I. 62.0–72.9, R^2 0.98]; for CDV, > 500 μM . Regarding cell proliferation, calculated CC_{50} values were: for BCV, 6.9 μM [C.I. 5.7–8.4, R^2 0.97]; for CDV, > 500 μM . Comparing antiviral potency (data from Fig. 4) and effects on cell viability and cell proliferation (data from Figs. 5 and 6), BCV yielded selectivity indexes (SI) values of 303.6 and 31.5, respectively.

3.4. Enantiomer-specific antiviral activity in EPCs and UT7/EpoS1

To confirm that the observed inhibitory effects on the virus were specifically exerted by the metabolite, CDV-PP, the activity of the functional stereoisomer of BCV, BCV (S), was compared with that of its cognate enantiomer BCV (R). The latter is not converted to CDV-PP within cells. Infection was carried out following the established experimental scheme with both enantiomers tested in the range of 0.01–100 μM in both EPCs and UT7/EpoS1 cell systems. For each sample series, the amounts of viral DNA were assessed by qPCR and EC_{50} values calculated as described previously (Fig. 8). Cell viability was assessed in the same assay format by a WST-8 based assay and cytotoxicity CC_{50} values determined (Fig. 9).

For this experimental series, calculated EC_{50} and CC_{50} values for both EPCs and UT7/EpoS1 cell systems are reported in Table 1. In EPCs, the SI values (S.I.: $\text{CCK8 CC}_{50}/\text{EC}_{50}$) for BCV (S) and BCV (R) were 6.5 and 1.6 respectively, yielding an S/R ratio of 4.0. In UT7/EpoS1, SI values were 95.1 and 1.3, yielding an S/R ratio of 73.2. Altogether, experimental data indicated that the inhibitory activity on B19V replication was exerted by the active enantiomer, BCV (S), to a significantly higher extent with respect to its cognate enantiomer BCV (R); the observed differences in selectivity indexes and S/R activity ratios were dependent on the cellular environment.

4. Discussion

The present study was conducted with the aim of closing the gap in the development of an antiviral strategy against B19V, a virus commonly distributed in the population, and responsible for a wide spectrum of clinical manifestations. In the absence of a vaccine to prevent infection and an effective antiviral therapy, treatment of B19V

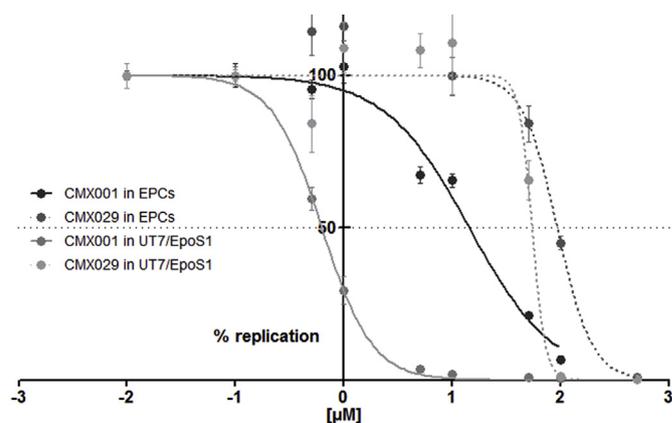


Fig. 8. Viral replication, normalized response to BCV (S) and BCV (R) in EPCs and UT7/EpoS1. Normalized dose-response curve, calculated from qPCR data obtained at 48 hpi for the different experimental series (B19V in EPCs and UT7/EpoS1). Log B19V DNA as a function of BCV (S) (CMX001) or BCV (R) (CMX029) concentration, expressed as percentage replication compared to control samples. EC_{50} values derived from this plot are reported in Table 1.

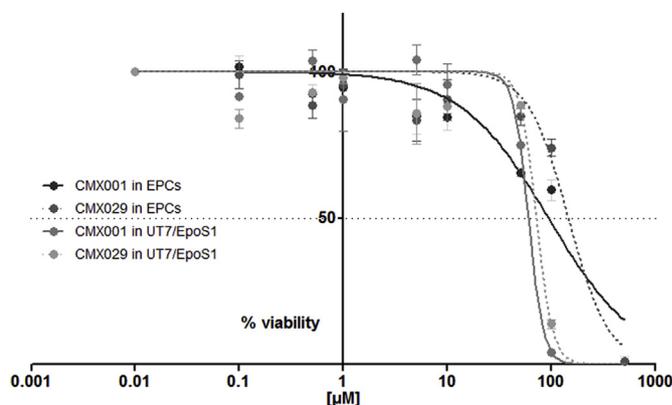


Fig. 9. Cell viability, normalized response to BCV-S and BCV-R in EPCs and UT7/EpoS1. Normalized dose-response curve, calculated from CCK8 assay data obtained at 48 hpi for the different experimental series (B19V in EPCs and UT7/EpoS1). Cell viability as a function of BCV (S) (CMX001) or BCV (R) (CMX029) concentration, expressed as percentage value compared to control samples. CC_{50} values derived from this plot are reported in Table 1.

Table 1

Effects of tested compounds on B19V replication, cell viability and cell proliferation.

Compound	Cells/moi	EC_{50} [C.I.]	CCK8 CC_{50} [C.I.]	S.I.
BCV (S)	EPC 10^4	14.3 [11.8–17.3]	93.4 [68.9–126.6]	6.5
CMX001	UT7 10^4	0.63 [0.58–0.68]	59.9 [52.7–68.1]	95.1
BCV (R)	EPC 10^4	93.0 [77.4–111.8]	146.2 [106.4–200]	1.6
CMX029	UT7 10^4	54.7 [42.7–69.9]	72.1 [62.5–83.1]	1.3

EC_{50} and CC_{50} and respective 95% C.I. values are expressed as μM .

infections remains limited to supportive and nonspecific (e.g., IVIG) interventions (Gallinella, 2013; Qiu et al., 2017). Therefore, we believe that the efforts in research and development of antiviral drugs with potent inhibitory activity against B19V should be considered of utmost importance so as to increase the available options for specific therapeutic and/or preventive treatments.

We have previously shown that CDV, an antiviral with a broad spectrum *in vitro* activity but with limited clinical effectiveness, inhibits B19V replication *in vitro* (Bonvicini et al., 2015, 2016). This was the first reported evidence of activity of an antiviral compound against B19V, and at the same time an expansion of the antiviral range of CDV

to encompass a ssDNA virus in addition to the more widely recognized activity against dsDNA viruses. The present set of experiments investigated the activity of the lipid conjugate of CDV, BCV, to assess its potential as an antiviral agent against B19V. BCV has been shown to be more potent than CDV against dsDNA viruses (Aldern et al., 2003; Williams-Aziz et al., 2005; Hostetler, 2009) and its administration has been shown to be safe, well-tolerated and to produce a low circulating plasma concentration of CDV (Painter et al., 2012), which greatly reduces the nephrotoxicity associated with intravenous CDV administration (Tippin et al., 2016).

Our experiments demonstrate that BCV, compared to CDV, exerts an enhanced inhibitory effect on replication of the ssDNA virus B19V. This is evident in both the EPC and UT7/EpoS1 cell system even at the highest multiplicities of infection tested, with a distinct dose-dependent reduction in the amount of viral DNA synthesized. This inhibitory activity was achieved at relatively lower compound concentrations in UT7/EpoS1 than in EPCs. In addition, in the EPC cells only BCV, and not CDV, yielded a significant EC_{50} value and achieved complete inhibition of viral replication. Accordingly, the effects of BCV on cell viability and proliferation were measurable at lower concentrations compared to CDV in both cellular systems. The effect on cell proliferation were more pronounced than those on cell viability, nevertheless a favorable selectivity index for their antiviral effect was demonstrated in both assay systems.

There are two enantiomeric forms of BCV, (S) and (R), but only BCV (S) is the active enantiomer. BCV (S) utilizes the natural lipid uptake pathway in cells that results in its rapid uptake, consequent increase in the active antiviral, CDV-PP, and hence an increase in antiviral potency (Hostetler, 2009; Painter et al., 2012). On the other hand, CDV-PP is undetectable in cells treated with the R-enantiomer (McMullan et al., 2016). We demonstrated here that the inhibition of viral replication is restricted to the S-enantiomer of BCV, thus supporting the hypothesis of an active and specific antiviral role of CDV-PP derived from BCV rather than an inhibition of viral replication related to a general, nonspecific cytotoxic effect.

The antiviral activity of BCV and CDV is directly correlated to CDV-diphosphate (CDV-PP) which acts as an alternate substrate for viral DNA synthesis (De Clercq and Holy, 2005; Magee and Evans, 2012). Both BCV and CDV have been reported to possess antiviral activity also against DNA viruses that do not encode their own polymerases (Johnson and Gangemi, 1999; Randhawa et al., 2006; Jiang et al., 2010), however in these cases the exact mechanism of inhibition has not been yet established. Possible mechanism could include the replication complex of these viruses being more conducive to incorporating CDV into viral DNA causing chain termination or impeding template activity, a decreased ability of the replication complex to excise incorporated CDV, or the possibility of inhibiting helicase activity, as encoded by LT in case of the polyomaviruses and E1 in the case of papillomaviruses (Andrei et al., 2015; Tsang et al., 2016). It should be noted that the B19V genome replicative intermediates are actually dsDNA forms that utilize the host replication machinery and metabolic environment, so the mechanisms of inhibition could also probably be similar to what is proposed for other dsDNA virus that do not encode their own DNA polymerase.

Replication of B19V *in vitro* is limited by a highly restrictive cellular environment, so that even at the high multiplicity of infection of 10^4 , only a minor fraction of cells can support viral replication (Fig. 2) (Manaresi et al., 2015). As a result, a relatively low (~ 1 – 2 Log) increase in the amount of viral DNA in the course of infection from 2 to 48 hpi is normally achieved (Figs. 1 and 3). Although this can be considered suboptimal, the data obtained allowed discrimination of the activities of the different compounds tested and calculation of the different EC_{50} values in different cellular environments. EPCs and UT7/EpoS1 cell systems possess different properties that are to be taken into consideration when considering our experimental results. EPCs are primary cells with a limited lifespan, and represent a cellular system

more closely resembling the natural target cells within bone marrow environment (Merryweather-Clarke et al., 2011). Circulating EPCs can be selectively induced to proliferate and differentiate along the erythroid lineage, progressively acquiring sensitivity to viral infection and full permissiveness to viral replication, in particular at a stage corresponding to the proerythroblast differentiation stage in the bone marrow (Bua et al., 2016). However, EPCs are cultured in conditions that approximate the conditions of the bone marrow environment, and constitute a population with heterogeneous characteristics related to the differentiation stage, proliferation rate and metabolic activity. The effective inhibitory activity of BCV observed in these cells is of crucial relevance, especially in comparison to the limited activity shown by CDV. The inherent heterogeneity and a slow overall metabolic rate of EPCs may hinder the accurate evaluation of the antiviral potency of BCV, as the conversion of BCV to the active antiviral CDV-PP is expected to be more pronounced in metabolically active cells, and hence the lower potency in EPCs observed in our experiments. On the other hand, UT7/EpoS1, a cell line of myeloblastoid origin, is the most permissive and commonly used cellular line when investigating B19V (Wong and Brown, 2006). In these cells, active viral replication is shown to occur only in a subset of cells (Manaresi et al., 2015; Bonvicini et al., 2013b), but at the population level the degree of replication of viral DNA shows relative increases comparable to the fully permissive EPCs (Bonvicini et al., 2006, 2008). Thus, UT7/EpoS1 may constitute a more suitable and appropriate experimental system, for evaluating the activity of BCV against B19V. We report here that the EC_{50} against B19V is in the same order of magnitude as compared to those determined for other dsDNA viruses, and more importantly, within the range of concentrations achieved with clinically relevant doses (Hostetler, 2009; Painter et al., 2012).

B19V is highly dependent on cellular machinery for its lifecycle (Luo and Qiu, 2015; Ganaie and Qiu, 2018) and offers few virus specific antiviral targets, but the observed capacity of broad spectrum antiviral compounds to interfere with its biological activity indicates the presence of pathways susceptible to potential inhibition in a complex and as yet not fully characterized virus-cell interaction network. We believe that the differences observed within the two cellular systems may reflect differences in the permeability of cells to the compounds, the overall metabolic rate of the cellular environment, or to different mechanisms of interference with the normal metabolism of nucleotides. Hence, in the present set of experiments, the observed lower efficacy of BCV in EPCs than in UT7/EpoS1 may reflect the low metabolic rate in these cells as playing an important role in the higher EC_{50} scored in our assays. Further experiments will be required to evaluate the kinetics of CDV-PP formation within the cells, whether the actual intracellular concentration of compounds effectively correlate with the antiviral activity, and if activity can be enhanced by an extended exposure of cells to non-toxic concentrations of drugs. The recent development of a reverse genetic system for B19V (Manaresi et al., 2017) will also provide a valuable tool to investigate in more detail the targets and mechanisms of activity of compounds in the observed inhibition of viral replication.

In conclusion, we demonstrate that the investigational compound, BCV, effectively inhibits B19V with highly favorable properties in terms of *in vitro* activity and selectivity, thus expanding its antiviral spectrum to encompass a ssDNA virus and establishing its potential as a B19V antiviral agent. Further progress in our knowledge of the biological characteristics of B19V and of its interaction with the cellular environment will lead to a better understanding of the viral target/s and mechanism/s appropriate for the continued development of an antiviral strategy directed against the human pathogenic virus B19V.

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