

Design, synthesis, and antiviral activity of new 1*H*-1,2,3-triazole nucleoside ribavirin analogs

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Abstract Ribavirin is a broad antiviral compound with demonstrated activity against herpes simplex virus (HSV), human immunodeficiency virus HIV-1, influenza virus, respiratory syncytial virus, and hepatitis C virus, among other viruses. However, routine clinical use of ribavirin is limited because this compound is considerably cytotoxic. Herein, we describe the design, synthesis, and antiviral activity of new nucleoside ribavirin analogs based on the following: (1) ring bioisosterism of a 1,2,4-triazole for a 1,2,3-triazole; (2) amide group exchange for other substituents, such as *c*-propyl, methyl carboxylate, or trifluoromethyl groups; and (3) the ribofuranose remained linked to the triazole ring. Compounds **5a–c** were obtained with yields of 65–36 % and tested against Influenza A and HSV-1 replication as well as reverse transcriptase (RT) from human immunodeficiency virus type 1 (HIV-1 RT). Compound **5b** (R = CO₂CH₃) was the most effective analog, with IC₅₀ values 14 and 3.8 μM for Influenza A and HIV-1 RT, respectively.

Keywords Antiviral · Influenza A · Ribavirin · Reverse transcriptase · 1,2,3-Triazole

Introduction

Ribavirin is a broad antiviral compound that targets viral DNA/RNA polymerases and cellular enzymes (De Clercq, 2004; Magden *et al.*, 2005). Its activity has been demonstrated for herpes simplex virus (HSV) (De Clercq, 2004), human immunodeficiency virus (HIV-1) (De Clercq, 2004), influenza virus (Saladino *et al.*, 2010), respiratory syncytial virus (RSV) (Krilov, 2011), and hepatitis C virus (HCV) (Sarrazin *et al.*, 2012), among other viruses. However, routine clinical use of ribavirin is limited because it is considerably cytotoxic (De Clercq, 2004; Magden *et al.*, 2005). This drug is primarily used in a combinatory therapy for treating HCV infection with α-interferon (Sarrazin *et al.*, 2012) and in severe cases of acute respiratory infection from RSV (Krilov, 2011). More recently, given the emergence of multidrug-resistant influenza strains since the 2009 pandemics, clinical trials have evaluated the potency of ribavirin and neuraminidase inhibitor (NAIs) combinations (Kim *et al.*, 2011). Therefore, the ribavirin chemical structure can be an interesting prototype for developing novel antiviral compounds.

A series of 4*H*-1,2,4-triazole 3,4,5-trisubstituted derivatives was investigated for antiretroviral activity against HIV-1 reverse transcriptase enzyme (HIV-1 RT). The key interactions between the triazoles and HIV-1 RT were identified by X-ray crystallography (Kirschberg *et al.*, 2008). Girardet and co-workers described synthesis and the antiviral activities of 1,2,4-triazoles derivatives against several non-nucleoside reverse transcriptase inhibitor (NNRTI)-resistant strains of HIV-1. Several such

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compounds exhibited potent antiviral activities against efavirenz- and nevirapine-resistant viruses with K103N and/or Y181C mutations or a Y188L mutation (Rosa *et al.*, 2006).

The 1,2,4-triazolacetamide derivatives were evaluated for inhibitory effect on HIV-1 RT. This study identified a series of triazoles with inhibitory activity against wild-type HIV-1 and the viral strains with a K103N mutation (Zhan *et al.*, 2011). Chrysin and co-workers confirmed the bioisosteric relationship between the amide moiety and the 1,2,3-triazole ring (Chrysin *et al.*, 2009).

The 1,2,3-triazole and its derivatives can be synthesized using “click chemistry” and have been widely used in medicinal chemistry, such as in the pathogenesis of Alzheimer’s disease (as IDO inhibitors) (Huang *et al.*, 2011) as well as for antitubercular (Boechat *et al.*, 2011a; Jordão *et al.*, 2011a), antifungal (Aher *et al.*, 2009), antibacterial (Wang *et al.*, 2010), antileishmanial (Ferreira *et al.*, 2007; Coghi *et al.*, 2008), trypanocidal (Júnior *et al.*, 2009; Silva *et al.*, 2008), and anti-cancer (Yan *et al.*, 2010) activities.

Triazole scaffolds show significant biological potential for developing novel antiviral agents to treat various viral diseases (Kharb *et al.*, 2011; Jordão *et al.*, 2009, 2011b; Pérez-Castro *et al.*, 2007) in the future, including through in vitro inhibitory profiles against HIV-1 RT (Piotrowska *et al.*, 2012; Silva *et al.*, 2009).

Synthetic methodologies for preparing triazole and nucleoside derivatives have been described in the literature (Kharb *et al.*, 2011; Amblard *et al.*, 2009; Shawali, 2010). Preparing 1,2,3-triazoles through 1,3-dipolar cycloaddition using various acetylenes and an azido group connected to a sugar unit is a widely used method for synthesizing triazole nucleosides and nucleotides with biological activity (Chrysin *et al.*, 2009; Pérez-Castro *et al.*, 2007; El Akri *et al.*, 2007; Elayadi *et al.*, 2012; Rocha *et al.*, 2012; Ferreira *et al.*, 2010a, b; Hou *et al.*, 2011).

We have prepared bioactive triazoles using the standard medicinal chemistry such as isosteric replacement of functional groups and ring bioisosterism (Boechat *et al.*, 2011a, b; Ferreira *et al.*, 2007) (Fig. 1). Herein, we have designed new nucleosides **5a–c**, which are ribavirin analogs with potential antiviral activity. Such derivatives were based on ring bioisosterism wherein a 1,2,3-triazole substitutes for the 1,2,4-triazole. We also investigated the different groups bound to a 1,2,3-triazole ring instead of the ribavirin amide group. The selected substituents were *c*-propyl, methyl carboxylate, and trifluoromethyl.

To synthesize nucleosides **5a–c**, the respective triazoles were first synthesized and then coupled to the ribose sugar moiety. For the last step, we used the methodology described by de Souza and co-workers wherein the nitrogen base is coupled to the sugar group (Matta *et al.*, 1996, 1999).

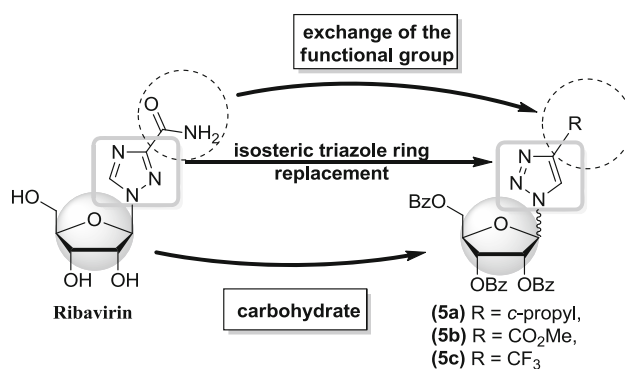


Fig. 1 Rational approach to design ribavirin-based nucleoside derivatives **5a–c**

Importantly, the ribofuranose moiety remains protected based on previous docking studies that described optimal intermolecular (docking) energies. Moreover, in a previous work we have identified a better performance of protected nucleoside when assayed against virus (Silva *et al.*, 2009).

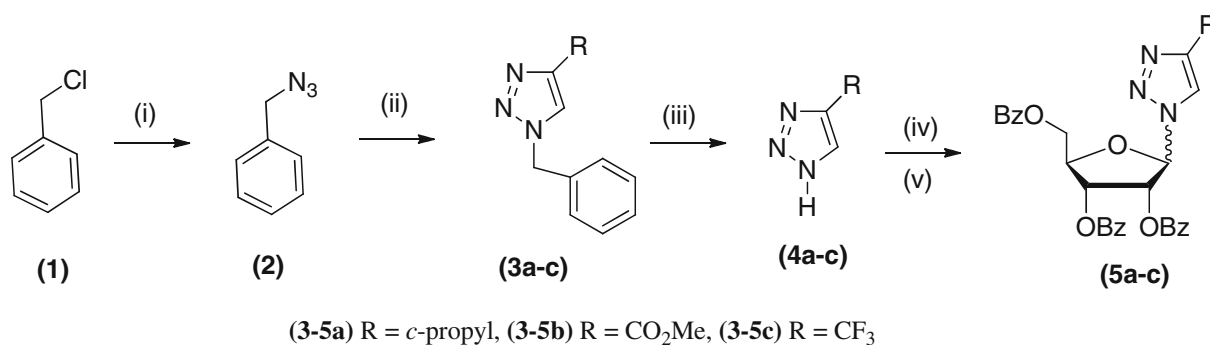
Results and discussion

Chemistry

To prepare the target compounds **5a–c** using the synthetic route described herein (Scheme 1), we began with (azidomethyl)benzene (**2**) that had previously been generated by nucleophilic substitution of (chloromethyl)benzene (**1**) with sodium azide and then reacted (**2**) with terminal alkynes to generate the 1-benzyl-1*H*-1,2,3-triazoles (**3a–c**) (Pagliai *et al.*, 2006). Click chemistry reaction conditions involved using $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$ as a catalyst that is fully controlled for reaction regioselectivity, and only the 1,4-regioisomers were generated (Lu and Gervay-Hague, 2007; Kolb *et al.*, 2001).

To synthesize the 1*H*-1,2,3-triazoles (**4a–c**), the respective benzyl-1*H*-1,2,3-triazoles (**3a–c**) were treated with palladium hydroxide on carbon under hydrogen flow, which generated the products (Esternam and Seebach, 1988; Kacprzak, 2005). The structure of compounds **3b** and **4a** were characterized using X-ray crystallography by our group (Boechat *et al.*, 2010).

Compounds **5a–c** were prepared in accordance with the methodology described in the literature (Matta *et al.*, 1996, 1999). The 1*H*-1,2,3-triazole (**4a–c**) derivatives were silylated with *N,O*-bis-(trimethylsilyl)trifluoroacetamide (BSTFA) containing trimethylchlorosilane (TMS-Cl). The corresponding trimethylsilyl derivatives formed were immediately condensed with 1-*O*-acetyl-2,3,5-tri-*O*-benzoyl- β -*D*-ribofuranose in a one-pot reaction under trimethylsilyl trifluoromethanesulfonate (TMSOTf). Good



Reagents and conditions: (i) NaN₃, CH₃CN/DMF, reflux, 20 h; (ii) appropriate acetylenes, L-ascorbic acid sodium salt, CuSO₄·5H₂O, H₂O/*t*-BuOH (1:1), 25 °C, 24 h; (iii) Pd(OH)₂/C, MeOH/H₂O, H₂, 25 °C, 72 h; (iv) BSTFA, TMS-Cl, CH₃CN, 70 °C, 6 h; (v) 1-*O*-acetyl-2,3,5-tri-*O*-benzoyl-β-D-ribofuranose, TMSOTf, CH₃CN, 25 °C, 24 h.

Scheme 1 Synthetic route used for preparing nucleosides **5a–c**

yields of the compounds were obtained and fully characterized by ¹H, ¹³C, and ¹⁹F nuclear magnetic resonance (NMR), IR, HRMS as well as elemental analysis (CHN).

The synthetic route was initiated by preparing a key intermediate via a reaction between (chloromethyl)benzene (**1**) and sodium azide to produce the (azidomethyl)benzene (**2**), as a yellow oil with a 99 % yield. Fourier transform infrared spectroscopy (FTIR) analysis showed a strong absorption band at 2113 cm⁻¹, which is characteristic of the stretching vibrations for the N₃ bond. The IR and the ¹H NMR confirm the chemical structure of the intermediate **2** as described in literature (Ankati and Biehl, 2009), (Campbell-Verduyn *et al.*, 2009). The azide was used in the next step without purification to prevent degradation.

Synthesis of 1-benzyl-1*H*-1,2,3-triazoles (**3a–c**) through a 1,3-dipolar cycloaddition reaction between the appropriate acetylenes, benzyl azide (**2**), and sodium ascorbate and catalyzed by Cu(I) was performed at 25 °C and the compounds protected from light because the aromatic azides are photosensitive. After purification in a flash-type column, compounds **3a–c** were produced as white or yellow crystals with yields from 86 to 31 %. FTIR of **3a–c** demonstrated the absence of stretching vibrations from an azide group. In the ¹H NMR spectra, signals from the respective protons were observed as a singlet at 8.99–7.86 ppm, which corresponds to the triazole ring, whereas signals that correspond to protons in the phenyl moiety appear as a multiplet at 7.41–7.35 ppm, and the benzylic CH₂ group was observed at 5.69–5.50 ppm. The **3b** derivative was described by our group previously and showed characteristic ¹H and ¹³C NMR signals at 3.82 and 160.50 ppm due to the methyl group and ester carbonyl, respectively, and the melting points (m.p.) value is in accordance with the literature (Boechat *et al.*, 2010).

The ¹⁹F NMR spectrum for compound **3c** showed a characteristic signal at –59.5 ppm that confirms the trifluoromethyl group.

The 1-benzyl-1*H*-1,2,3-triazole (**3a–c**) derivatives were reduced to the respective 1*H*-1,2,3-triazoles (**4a–c**) with palladium hydroxide on carbon under hydrogen flow in PARR Reactor hydrogenator for 72 h at 25 °C. The compounds were purified by silica gel flash-type column chromatography, and the triazoles (**4a–c**) were produced as white or light-yellow crystals at yields from 76 to 40 %.

The structures were confirmed by ¹H, and ¹³C NMR, which did not exhibit signals related to a benzyl group. The **4a** derivative was described by our group previously and the m.p. value is in accordance with the literature (Boechat *et al.*, 2010). Compound **4c** was directly used in the next step without purification to prevent degradation and was characterized by ¹H NMR and mass spectroscopy (MS).

To prepare the nucleosides (**5a–c**), a modified procedure was used that included the 1*H*-1,2,3-triazoles (**4a–c**) with BSTFA/TMS-Cl under an argon atmosphere at 70 °C for 6 h. This procedure yielded the corresponding trimethylsilyl derivatives, which were immediately condensed with ribofuranose in a *one-pot* reaction under TMSOTf. This procedure generates acceptable yields (65–35 %) of the corresponding protected nucleosides. Spectroscopic analysis confirmed the formation of such products.

The anomeric protons of nucleosides **5a–b** were identified by the peaks at 6.65 and 6.79 ppm, respectively (*d*, *J* = 3.1 Hz), and for **5c** a doublet at 5.76 ppm (*t*, *J* = 5.9 Hz) was observed because of H–F couples. The ¹⁹F NMR spectrum for compound **5c** showed a characteristic signal at –74.6 ppm that confirmed the trifluoromethyl group.

Molecular modeling

The target compounds **5a–c** were modeled in silico, and energy minimization was performed over 1,000 steps using the steepest descent method, Gasteiger–Hückel charges, a dielectric constant 80, and the Tripos force field. The structures were further optimized using the conjugated gradient method.

The crystal structure of HIV-1 RT (PDB entry: 3V4I) (Das *et al.*, 2012) was used as the molecular target for ligand–enzyme docking simulations (Dias and Azebedo, 2008) by applying the MolDock algorithm (Thomsen and Christensen, 2006) implemented in Molegro Virtual Docker software (Molegro, 2009).

The optimum docking protocol was constructed by adding flexibility to the enzyme-binding pocket (“induced fit”). Each compound was docked with softened potentials (steric, hydrogen bonding, and electrostatic forces); the enzyme residues remained rigid in the default conformation. Next, the residue side chains that were sufficiently close to the compound to interact were energy minimized. The final step was energy minimization of the compounds.

Importantly, the docking simulations were performed with the protected form of the compounds i.e., the ribofuranose ring remained bound to the triazole ring through C–N coupling (Fig. 1); this was the form with the best intermolecular (docking) energies.

Antiviral activity of novel ribavirin analogs

Considering the broad range of viruses targeted by ribavirin (De Clercq, 2004), we tested the effects of the novel analogs of this compound against Influenza A and HSV-1 replication as well as the activity of HIV-1 RT. Shown in Table 1, our analogs inhibited Influenza A replication and RT activity with different potencies; compound **5b** (R = CO₂CH₃) was the most effective analog, with IC₅₀ values 14 and 3.8 μM for the two targets, respectively.

When we compared the antiviral data for ribavirin from the literature using assay conditions similar to ours

(Markland *et al.*, 2000; Kornev *et al.*, 2011; Fernandez-Larsson and Patterson, 1990), we found that our molecules are generally less cytotoxic and more potent against influenza A and RT. Ribavirin inhibits influenza A replication with an IC₅₀ at approximately 30 μM (Markland *et al.*, 2000; Kornev *et al.*, 2011), which is comparable to our observations for compound **5a** but 2.0- and 1.5-fold higher than the IC₅₀ values for compounds **5b** and **5c**, respectively.

In regard to HIV-1, in its unphosphorylated, diphosphorylated, and triphosphorylated forms, ribavirin inhibits the enzyme RT with IC₅₀ of 615, 81, and 112 μM (Fernandez-Larsson and Patterson, 1990), respectively. Thus, compound **5b** is ~20-fold more potent against HIV-1 RT than the most active form of ribavirin (Fernandez-Larsson and Patterson, 1990) (Table 1).

It is described in the literature that ribavirin cytotoxicity varies for different cell types from 70 to ~500 μM (Kornev *et al.*, 2011), whereas the CC₅₀ values for our compounds were in the 550 μM range. Thus, our compounds tend to be in the low cytotoxicity range compared with their counterpart, ribavirin (Kornev *et al.*, 2011).

Although the ribavirin analogs developed herein are superior to ribavirin in antiviral activity against influenza A and HIV-1 RT, other compounds with routine clinical use for such viruses, including oseltamivir carboxylate for influenza A and AZT-TP for HIV-1 RT, are more potent than our ribavirin analogs. Nevertheless, our molecules are

Table 2 Energy docking results

| Compounds | E_{docking} (kcal/mol) | E_{HBond} (kcal/mol) | E_{Steric} (kcal/mol) | E_{vdw} (kcal/mol) |
|-------------|------------------------------------|----------------------------------|-----------------------------------|--------------------------------|
| 5a | −137.12 | −13.93 | −102.36 | −26.73 |
| 5b | −130.77 | −13.21 | −129.96 | 18.68 |
| 5c | −118.58 | −11.40 | −106.71 | 297.93 |
| AZT | −114.73 | −3.52 | −111.21 | −39.19 |
| Ribavirin | −110.60 | −6.97 | −103.63 | −38.34 |
| Zalcitabine | −99.88 | −3.57 | −96.31 | −31.10 |

Table 1 Antiviral activity and cytotoxicity of novel ribavirin analogs and reference compounds

| Compounds | IC ₅₀ (mean ± SEM; μM) | | | CC ₅₀ (mean ± SEM; μM) | | |
|-------------------------|-----------------------------------|--------------|-------------|-----------------------------------|-----------------------|----------|
| | Influenza A | HIV-1 RT | HSV-1 | MDCK | Supt1 | Vero |
| 5a | 33 ± 5.0 | 30 ± 9.0 | >50 | 512 ± 23 | 623 ± 16 | 586 ± 12 |
| 5b | 14 ± 1.6 | 3.8 ± 0.23 | >50 | 533 ± 14 | 598 ± 32 | 594 ± 56 |
| 5c | 19 ± 3.0 | 14 ± 6.0 | >50 | 518 ± 12 | 555 ± 45 | 532 ± 24 |
| Oseltamivir carboxylate | 0.03 ± 0.002 | NE | NE | >2000 | NE | NE |
| ACV | NE | NE | 0.95 ± 0.12 | NE | NE | 835 ± 62 |
| AZT-TP | NE | 0.01 ± 0.002 | NE | NE | 126 ± 15 ^a | NE |

^a For cytotoxicity assays AZT, in its non-phosphorylated form, was used

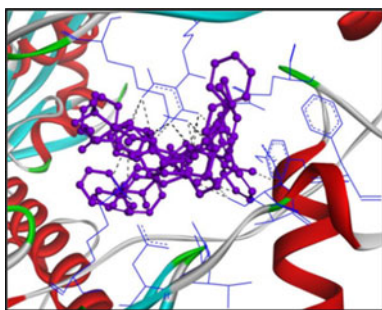


Fig. 2 Detailed depiction of the interactions between the compounds and HIV-1 RT

less cytotoxic than AZT. As shown, HSV-1 replication was not affected by our ribavirin analogs, which may suggest that our compounds are more specific (Table 1).

The development of novel strategies to control HIV-1 and influenza A replication is an important task given the large economic and public health burden from such agents

(Pasman, 2012). In particular, the emergence of multidrug-resistant strains of such agents further motivates the search for novel compounds (Pasman, 2012). The ribavirin mechanism of action may vary with inhibition of different viral agents, including inhibition of inosine monophosphate dehydrogenase (which would reduce GTP levels in the cells) and viral RNA polymerase and/or generation of an error catastrophe through its incorporation by viral RNA polymerase (De Clercq, 2004; Magden *et al.*, 2005). Thus, our analogs are interesting for the development of novel anti-influenza and antiretroviral agents, which may have a mechanism of action mentioned above or another one. Considering the fact that the tested compounds were potent against HIV-1 RT, we performed *in silico* studies to further characterize the docking sites for our compound on HIV-1 RT.

For the interactions between HIV-1 RT and compounds **5a–c**, docking scores, hydrogen bonds, and steric and van der Waals energy contributions are shown in Table 2.

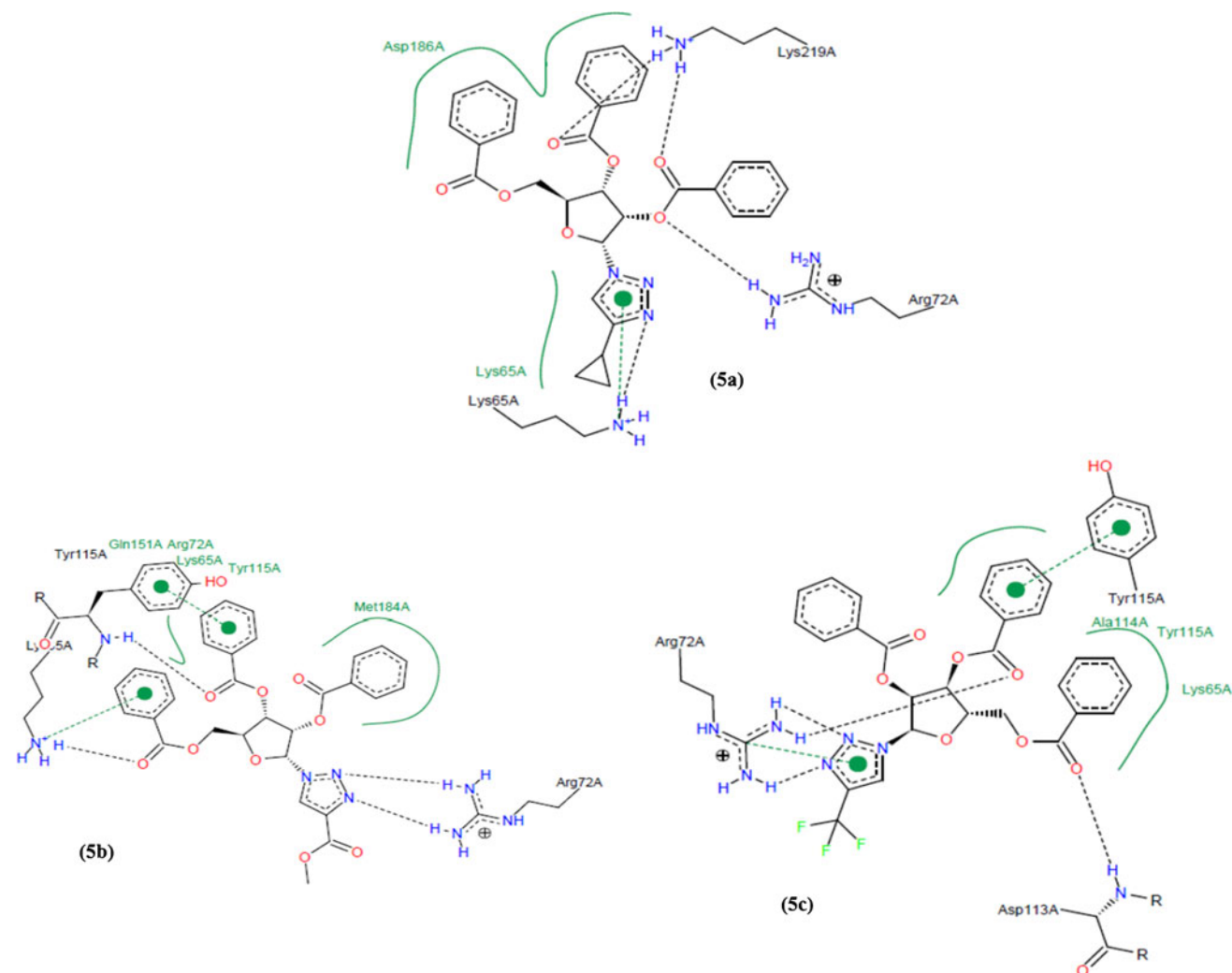


Fig. 3 2D representation of the interactions between the compounds and enzyme

Three additional compounds were included: AZT, ribavirin, and zalcitabine. The first was used as the standard molecule in the experimental assays. Ribavirin was included because the work herein includes design of its analogs. Consequently, ribavirin is the primary reference compound. Zalcitabine is also an example of a known reverse transcriptase inhibitor. According to the relevant values, both compounds **5a** and **5b** interact better with the enzyme than compound **5c**. Moreover, the correlation between the docking score, hydrogen bonds, and van der Waals energies is clear. This correlation was confirmed by the calculated cross-correlation matrix of the energy values. Figure 2 shows a detailed depiction of the interactions between the compounds and HIV-1 RT generated from docking calculations. Figure 3 shows a 2D-representation of the interactions between the compounds and enzyme (Stierand and Rarey, 2010). Notably, the intermolecular hydrogen bonds, which are shown as black dashed lines, are important and operate as “molecular anchors” to bind the compounds to the enzyme active site.

Conclusions

Three novel nucleosides of 1,2,3-triazole analogs were successfully synthesized via cyclization of azides and terminal alkyne using “click chemistry” with a 65–35 % yield. The nucleoside structures are supported by IR; ^1H , ^{19}F , and ^{13}C NMR [DEPT, ^1H - ^1H -COSY, HETCOR] spectroscopy; CHN and MS data.

Compound **5b** was superior to ribavirin in terms of antiviral activity against influenza A replication and HIV-1 RT activity; however, it was less potent than other drugs, such as oseltamivir carboxylate and AZT-TP. Remarkably, the most effective compound tested against HIV-1 RT, molecule **5b**, required Tyr115 in this enzyme structure for docking. This characteristic is consistent with another potent antiretroviral and anti-HSV-1 molecule which we previously described (Souza *et al.*, 2009).

Finally, our results indicate that the chemical structure of our synthesized 1,2,3-triazole analogs can be interesting prototypes for development of novel antiretroviral and anti-influenza drugs.

Experimental section

The ^1H , ^{13}C , and ^{19}F nuclear magnetic resonance (NMR) spectra were obtained generated at 400.00, 100.00, and 376.00 MHz, respectively, on a BRUKER Avance instrument equipped with a 5-mm probe and using tetramethylsilane as an internal standard.

The chemical shifts (δ) are reported in ppm and the coupling constants (J) in Hertz. The FTIR absorption spectra were recorded on a Shimadzu mode IR Prestige-21 spectrophotometer through KBr reflectance (cm^{-1}). The electron ionization mass spectrometry (EI-MS, scan ES + Capilar (3.0 kV)/cone (30 V)/extractor (1 V)/RF lens (1.0 V)/source temperature (150 °C)/desolvation temperature (300 °C)) were recorded using a Micromass/Waters Spectrometer (model: ZQ-4000). The HRMS data were obtained using LC-MS—Bruker Daltonics MicroTOF (analyzer time of flight). The CHN data were obtained using a Perkin-Elmer 2400 CHN Analyzer. The (m.p.) were determined using a Büchi model B-545 apparatus. The Mini Bench Top Reactor PARR Model 4842 hydrogenator was used for the reactions for obtaining the compounds (**4a–c**). Thin layer chromatography (TLC) was performed using a silica gel F-254 Glass Plate (20 × 20 cm^2). Column chromatography was performed using a silica gel 60 (0.040–0.063 mm). The remaining reagents and solvents used were analytical grade.

Procedure for preparing (azidomethyl)benzene (**2**)

These compounds were prepared by the reaction of (chloromethyl)benzene (**1**) (7.9 mmol) and NaN_3 (11.8 mmol) in CH_3CN (50 mL) and DMF (3 mL). The reaction mixture was kept under reflux with stirring for 24 h. The development of the reaction was followed by TLC. The mixture was diluted with 50 mL of H_2O and extracted with CH_2Cl_2 (3 × 30 mL), and the combined organic extracts were washed with water (3 × 30 mL), dried over anhydrous MgSO_4 , and filtered; the solvent was removed by vacuum. The product was obtained as yellow oil with 99 % yield.

IR (KBr): 2113 (N_3). ^1H NMR (DMSO-d_6 , 400 MHz, δ in ppm): 7.31–7.41 (m, 5H), 4.34 (s, 2H). [lit. IR (KBr): 2098.3 (N_3) (Ankati and Biehl, 2009). ^1H NMR 7.25–7.43 (m, 5H), 4.35 (s, 2H) (Campbell-Verduyn *et al.*, 2009)].

General procedure for preparing the compounds **3a–c**

These compounds were prepared by the reaction of (azidomethyl)benzene (**2**) (22.5 mmol), with the appropriate acetylene (33.7 mmol) and L-ascorbic acid sodium salt (2.25 mmol), and of $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$ (0.225) mmol in a mixture of $\text{H}_2\text{O}/t\text{-BuOH}$ (1:1) (40 mL). The reaction mixture was kept under vigorous stirring at 25 °C for 24 h. The development of the reaction was followed by TLC. After cooling, the reaction mixture was poured into ice-cold water (50 mL). The precipitate was filtered and dried. The residual crude product was purified via silica gel column chromatography using a gradient mixture of

hexane/ethyl acetate. The compounds **3a–c** were obtained as a white solid with 86–31 % yield.

1-Benzyl-4-cyclopropyl-1H-1,2,3-triazole (3a)

Yield: 84 %. m.p. 61.2–62.4 °C. IR (KBr): 3462; 3076; 1635; 1215; 1047; 1028; 1012; 719. ¹H NMR (DMSO-d₆, 400 MHz): 0.67–0.71 (m, 2H, CH₂ cyclopropyl); 0.89–0.85 (m, 2H, CH₂ cyclopropyl); 1.88–1.95 (m, 1H, CH cyclopropyl); 5.50 (s, 2H, CH₂); 7.28–7.29 (m, 2H, H-Ph); 7.28–7.38 (m, 3H, H-Ph); 7.86 (s, 1H, CH triazole). ¹³C NMR (DMSO-d₆, 100 MHz): 6.48 (CH cyclopropyl); 7.58 (2C, CH₂ cyclopropyl); 52.6 (CH₂); 120.8; 127.7; 127.9; 128.6; (all C-Ph); 136.1 (CH triazole); 149.2 (Cq triazole).

Methyl 1-benzyl-1H-1,2,3-triazole-4-carboxylate (3b)

Yield: 86 %. m.p. 104.1–106.6 °C (Lit. m.p. 104.0–107.0 °C). IR (KBr): 3431; 3111; 1724; 1541; 1230; 1047; 1020; 713. ¹H NMR (DMSO-d₆, 400 MHz): 3.82 (s, 3H, CO₂CH₃); 5.66 (s, 2H, CH₂); 7.33–7.38 (m, 5H, H-Ph); 8.89 (s, 1H, CH triazole). ¹³C NMR (DMSO-d₆, 100 MHz): 51.6 (CO₂CH₃); 53.0 (CH₂); 127.9; 128.2; 128.7; 129.1; (all C-Ph); 135.3 (CH triazole); 138.6 (Cq triazole); 160.5 (CO₂CH₃).

1-Benzyl-4-(trifluoromethyl)-1H-1,2,3-triazole (3c)

Yield: 31 %. m.p. 73.0–75.0 °C. IR (KBr): 3458; 3089; 1570; 1382; 1217; 1151; 1051; 997; 709. ¹H NMR (DMSO-d₆, 400 MHz): 5.69 (s, 2H, CH₂); 7.35–7.41 (m, 5H, H-Ph); 8.99 (s, 1H, CH triazole). ¹³C NMR (DMSO-d₆, 100 MHz): 120.75 (q, *J* = 265 Hz, CF₃); 125.6 (d, *J* = 2.5 Hz, CH triazole); 128.0; 128.3; 128.8; 135.0 (all C-Ph); 136.7 (CH triazole); 137.0 (Cq triazole). ¹⁹F NMR (DMSO-d₆, 376 MHz): –59.5 (s, 3F, CF₃).

General procedure for preparing the compounds **4a–c**

These compounds were prepared by the mixture of appropriate 1-benzyl-1H-1,2,3-triazole (**3a–c**) (18.6 mmol) and palladium hydroxide on carbon (20 wt.% loading, dry basis, matrix carbon, and wet support) (14.2 mmol) in a mixture of H₂O/MeOH (1:1; 100 mL). The reaction remained under hydrogen flow for 72 h at 25 °C. The reaction mixture was filtered on Celite and extracted with CHCl₃ (3 × 50 mL); the combined organic extracts were washed with water (3 × 30 mL), dried over anhydrous MgSO₄, and filtered; the solvent was removed by vacuum. The residual crude product was purified via silica gel column chromatography using a gradient mixture of

hexane/ethyl acetate. The compounds **4a–c** were obtained as a white solid with 76–40 % yield.

4-Cyclopropyl-1H-1,2,3-triazole (4a)

Yield: 76 %. m.p. 54.8–55.2 °C, (Lit. m.p. 54.8–55.2 °C). IR (KBr): 3446; 2983; 1637; 1230; 1006; 995. ¹H NMR (DMSO-d₆, 400 MHz): 0.67–0.74 (m, 2H, CH₂ cyclopropyl); 0.90–0.91 (m, 2H, CH₂ cyclopropyl); 1.92–1.98 (m, 1H, CH cyclopropyl); 7.55 (s, 1H, CH triazole); 14.63 (s, 1H, NH). ¹³C NMR (DMSO-d₆, 100 MHz): 6.31 (CH cyclopropyl); 7.79 (2C, CH₂ cyclopropyl); 130.3 (CH triazole); 149.1 (Cq triazole).

Methyl 1H-1,2,3-triazole-4-carboxylate (4b)

Yield: 40 %. m.p. 125.7–127.1 °C. IR (KBr): 3446; 3136; 1714; 1354; 1240; 1211; 779. ¹H NMR (DMSO-d₆, 400 MHz): 3.82 (s, 3H, CO₂CH₃); 8.52 (s, 1H, CH triazole); 15.7 (s, 1H, NH). ¹³C NMR (DMSO-d₆, 100 MHz): 53.0 (CO₂CH₃); 138.6 (Cq triazole); 160.5 (CO₂CH₃).

4-(Trifluoromethyl)-1H-1,2,3-triazole (4c)

Yield: 48 %. IR (KBr): 3444; 3091; 1635; 1217; 1153; 709. ¹H NMR (DMSO-d₆, 400 MHz): 8.50 (s, 1H, CH triazole). ¹⁹F NMR (DMSO-d₆, 376 MHz): –59.1 (s, 3F, CF₃). MS *m/z* [M + 1]⁺: 138.0.

General procedure for preparing the compounds **5a–c**

These compounds were prepared by the reaction of appropriate 1H-1,2,3-triazoles (**4a–c**) (1.83 mmol), *N,O*-bis(trimethylsilyl)trifluoroacetamide (BSTFA) (6.86 mmol), and 4.6 mmol of chlorotrimethylsilane (TMS-Cl) in CH₃CN (4 mL). The reaction mixture was kept under argon atmosphere at 70 °C for 6 h. Thereafter, the reaction mixture was cooled to 25 °C, and a solution of 1-*O*-acetyl-2,3,5-tri-*O*-benzoyl-β-D-ribofuranose (1.83 mmol) and trimethylsilyl trifluoromethanesulfonate (TMSOTf) (0.915 mmol) in CH₃CN (5 mL) was slowly added and remained under vigorous stirring for 24 h. The development of the reaction was followed by TLC. The resulting mixture was poured into ice-cold water (10 g) and neutralized with a saturated aqueous NaHCO₃ solution. The solution was extracted with CH₂Cl₂ (3 × 50 mL), the combined organic layers were washed with water (3 × 20 mL), and then dried over anhydrous MgSO₄. The solvent was removed under reduced pressure. The residual crude product was purified via silica gel column chromatography using a gradient mixture of hexane/ethyl acetate. The compounds **5a–c** were obtained as a white solid with 65–35 % yield.

(2*R*,3*R*,4*R*)-2-((benzoyloxy)methyl)-5-(4-cyclopropyl-1*H*-1,2,3-triazol-1-yl)tetrahydrofuran-3,4-diyl dibenzoate (**5a**)

Yield: 65 %. m.p. 142.7–143.5 °C. IR (KBr): 3446; 2981; 1637; 1724; 1274; 1126; 1093; 709. ¹H NMR (DMSO-*d*₆, 400 MHz): 0.68–0.72 (m, 2H, CH₂ cyclopropyl); 0.88–0.92 (m, 2H, CH₂ cyclopropyl); 1.92–1.99 (m, 1H, CH cyclopropyl); 4.55 (dd, *J* = 4.6, 12.2 Hz, 1H, CH₂); 4.66 (dd, *J* = 4.6, 12.2 Hz, 1H, CH₂); 4.91–4.95 (m, 1H, H-4'); 6.10 (t, *J* = 6.0 Hz, H-3'); 6.22 (dd, *J* = 3.1, 5.2 Hz, 1H, H-2'); 6.65 (d, *J* = 3.1 Hz, H-1'); 7.44–7.52 (m, 2H, H-3'', H-5''); 7.63–7.69 (m, 1H, H-4'); 7.90–7.95 (m, 2H, H-2'', H-6''); 8.12 (s, 1H, CH triazole). ¹³C NMR (DMSO-*d*₆, 100 MHz): 6.27 (CH); 7.52 (CH₂); 7.57 (CH₂); 63.3 (2C, CH₂); 71.0 (C-3'); 74.2 (C-2'); 79.5 (C-4'); 88.9 (C-1'); 120.7 (CH triazole); 128.2, 128.4, 129.0 (C-1''); 128.6, 128.6, 128.7 (C-3'', C-5''); 129.2, 129.2, 129.3 (C-2'', C-6''); 133.4, 133.8, 133.9 (C-4''); 149.4 (Cq triazole); 164.3 (C=O); 164.5 (C=O); 165.3 (C=O). Anal. Calcd. for C₃₁H₂₇N₃O₇: C, 67.26; H, 4.92; N, 7.59; Found: C, 67.52; H, 5.20; N, 7.64. HRMS (ESI) *m/z* [M + Na] Calcd.: 576.1747; Found: 576.1745

(2*R*,3*R*,4*R*)-2-((benzoyloxy)methyl)-5-(4-methoxycarbonyl)-1*H*-1,2,3-triazol-1-yl)tetrahydrofuran-3,4-diyl dibenzoate (**5b**)

Yield: 35 %. m.p. 190.0–192.0 °C. IR (KBr): 3429; 3124; 1720, 1600; 1274; 1126; 1103; 711. ¹H NMR (DMSO-*d*₆, 400 MHz): 3.85 (s, 3H, CO₂CH₃); 4.60 (dd, *J* = 4.7, 12.3 Hz, 1H, CH₂); 4.72 (dd, *J* = 4.7, 12.3 Hz, 1H, CH₂); 4.98–5.01 (m, 1H, H-4'); 6.15 (t, *J* = 5.8 Hz, H-3'); 6.31 (dd, *J* = 3.1, 5.2 Hz, 1H, H-2'); 6.79 (d, *J* = 3.1 Hz, H-1'); 7.46–7.52 (m, 1H, H-4'', H-5''); 7.65–7.70 (m, 1H, H-3''); 7.91–7.95 (m, 2H, H-2'', H-6''); 9.11 (s, 1H, CH triazole). ¹³C NMR (DMSO-*d*₆, 100 MHz): 51.9 (CO₂CH₃); 63.2 (2C, CH₂); 70.9 (C-3'); 74.5 (C-2'); 80.1 (C-4'); 89.6 (C-1'); 128.7 (CH triazole); 129.2, 129.3, 129.4 (C-1''); 128.6, 128.8, 129.0 (C-2''); 128.4, 128.2, (C-4''); 133.5, 133.9, 134.1 (C-3''); 139.0 (Cq triazole); 160.3 (CO₂CH₃), 164.3 (C=O); 164.6 (C=O); 165.3 (C=O). Anal. Calcd. for C₃₀H₂₅N₃O₉: C, 63.04; H, 4.41; N, 7.35; Found: C, 62.80; H, 4.58; N, 7.54. HRMS (ESI) *m/z* [M + Na] Calcd.: 594.1489; Found: 594.1483.

(2*R*,3*R*,4*R*)-2-((benzoyloxy)methyl)-5-(4-(trifluoromethyl)-1*H*-1,2,3-triazol-1-yl)tetrahydrofuran-3,4-diyl dibenzoate (**5c**)

Yield: 35 %. m.p. decomp. IR (KBr): 3435; 3109; 1680; 1123; 1093; 1130; 709. ¹H NMR (DMSO-*d*₆, 400 MHz): 4.51–4.63 (m, 2H, CH₂), 4.66–4.69 (m, 1H, H-2' or H-3' or H-4'), 5.75–5.77 (m, 1H, H-2' or H-3' or H-4'), 5.82–5.84

(m, 1H, H-2' or H-3' or H-4'), 5.89 (t, *J* = 5.9 Hz, 1H, H-1'), 7.40–7.53 (m, 6H, OBz), 7.61–7.68 (m, 3H, OBz), 7.85 (d, *J* = 8.0 Hz, 2H, OBz), 7.93 (d, *J* = 4.0 Hz, 2H, OBz), 8.02 (d, *J* = 8.0 Hz, 2H, OBz), 10.62 (d, *J* = 6.1 Hz, 1H, CH triazole). ¹³C NMR (DMSO-*d*₆, 100 MHz): 63.47 (2C, CH₂), 70.50 (C-1'), 73.69 (C-2'), 78.03 (C-3'), 82.60 (C-4'), 115.42 (q, *J* = 286.4 Hz, CF₃), 128.41 (d, *J* = 5.8 Hz, CH triazole), 128.59, 128.74, (C-4''), 129.26, 129.19, 129.16 (C-1''), 133.72, 133.44 (C-3''), 133.85 (Cq triazole), 156.79 (C=O), 164.54 (C=O), 165.37 (C=O). ¹⁹F NMR (DMSO-*d*₆, 376 MHz): –74.6 (s, 3F, CF₃).

Compounds

Compounds **5a–c** were diluted in 100 % dimethylsulfoxide (DMSO) and stored at –20 °C. The concentration used for the assays was below 0.01 %

Cells and virus

Vero (African green monkey kidney cells) and MDCK (Mardin–Darby canine kidney cells) cells were cultured with Dulbecco's modified Eagle's medium (DMEM; GIBCO, Grand Island, NY, USA), and a T-lymphoid cell line (Supt1) was cultured in RPMI1640 (GIBICO). The cultures were supplemented with 10 % fetal bovine serum (FBS; HyClone, Logan, UT, USA), 100 U/mL penicillin as well as 100 µg/mL streptomycin and incubated at 37 °C in 5 % CO₂.

The viral stocks were prepared as we have described previously (Souza *et al.*, 2010; Denizot and Lang, 1986). The Vero cells were infected with HSV-1 (AR-29 strain) in FBS-free DMEM, and the MDCKs were infected with influenza A (A/England/42/1972 strain) in DMEM with 0.2 % serum albumin and trypsin at 4 µg/mL (inoculation medium). After a 1-h inoculation period, the cells were washed using PBS, and DMEM with 2 % FBS or inoculation medium was added to HSV-1- or influenza A-infected cells, respectively. At 48-h post-infection (p.i.), the cells were lysed using three cycles of freezing and thawing then centrifuged at 1.500×*g* and 4 °C for 20 min; the supernatant was collected and stored at –70 °C for further studies.

Cytotoxicity assay

In 96-multiwell plates (1 × 10⁴/well), Vero, MDCK, or Supt1 cells were treated with different concentrations of compounds **5a–c** for 72 h; then 50 µL of a 1 mg/mL solution comprising 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT, Sigma) was added to cells diluted in DMEM without serum. MTT was removed after 3 h, 50 µL of acid-isopropanol (0.04 N HCl in isopropanol)

was added and the optical density (OD) was determined using an automatic plate reader with a 570 nm test wavelength and 690 nm reference wavelength (Kuo *et al.*, 2001). The cytotoxic concentration in 50 % (CC₅₀) was calculated by linear regression analysis of the dose–response curves generated from the data.

Anti-HSV-1 assays

Yield reduction assay

In 24-well plates (1×10^5 /well), Vero cell monolayers were infected with HSV-1 at a MOI 5 for 1 h at 37 °C. The cells were washed and treated with different concentrations of compounds **5a–c** in DMEM with 2 % FBS. After 20 h, the cells were lysed, the cellular debris was cleared by centrifugation, and virus titers at the supernatant were determined using a plaque-forming assay and the Vero cells, which is described in the following subsection. The HSV-1 yield was also generated at different concentrations of acyclovir (ACV).

Plaque-forming assay

In 6-well plates (3×10^5 /well), Vero cell monolayers were exposed to the supernatant from the yield reduction assay for 1 h at 37 °C. Next, the residual viruses were washed out, and DMEM with 5 % FBS and 1 % methylcellulose (Fluka) (overlay medium) was added to cells. After 72 h at 37 °C, the monolayers were fixed using 10 % formaldehyde in PBS and stained with a 0.1 % crystal violet solution in 70 % methanol, and the virus titers were calculated by scoring the plaque-forming units (PFU) (Souza *et al.*, 2009). The inhibitory concentration in 50 % (CC₅₀) was calculated by linear regression analysis of the dose–response curves generated from the data.

Anti-HIV-1 RT inhibitory activity

The inhibitory effect of compounds **5a–c** on the RT_{HXB2} RNA-dependent DNA polymerase (RDDP) activity was evaluated using purified recombinant HIV-1 enzyme as reported (Souza *et al.*, 2009) with minor modifications. RDDP activity was assayed in 50 mM Tris HCl (pH 7.8), 6 mM MgCl₂, 1 mM dithiothreitol, 50 mM KCl, 5 μM dTTP, 80 μg/mL poly(rA) oligo(dT)_{12–18} template primer (Pharmacia, Piscataway, NJ, USA), and 3 U of enzyme (one unit is the enzyme concentration that incorporates 1 pmol of dTTP per minute per mg of enzyme at 37 °C under the standard assay conditions). The isotopic dilutions for the reactions were prepared at the ratio 2 μCi [³H]dTTP (49 Ci/mmol)/2.7 μM dTTP. The reactions were initiated at 37 °C, incubated for 30 min, and arrested with 0.5 M

EDTA. The precipitate was collected on a Whatman DE 81 filter and washed with 0.1 M sodium phosphate; the incorporated nucleotides were measured by liquid scintillation (Packard tri carb 2100). The inhibitory concentration in 50 % (CC₅₀) was calculated by linear regression analysis of the dose–response curves generated from the data. The polymerization reactions were performed with and without as well as various concentrations of the tested compounds and with and without AZT-TP.

Anti-influenza assay

To evaluate the effect of compounds **5a–c** on Influenza A replication, MDCK cells were seeded in a 6-well plate and grown to 80 % confluence in DMEM with 10 % FBS. Prior to infection, the cells were washed two times with inoculation medium. Then, Influenza A was added in inoculation medium at an MOI 5 for 1 h at 35 °C and 5 % CO₂. Thereafter, the cells were washed to remove unbound virus and treated with different concentrations of the compounds or with NAI oseltamivir carboxylate as a reference compound. Influenza A-infected cells were incubated with the compounds for 3 days in inoculation medium; the culture supernatant was harvested and RNA was extracted using the QIAmp Viral RNA mini kit (Qiagen, CA) in accordance with the manufacturer's instructions. The purified RNA was subjected to a one-step real-time RT-PCR to detect a viral matrix gene in accordance with the CDC–WHO protocol for detecting this pandemic virus (WHO, 2009).

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