This article was downloaded by:

On: 26 January 2011

Access details: Access Details: Free Access

Publisher Taylor & Francis

Informa Ltd Registered in England and Wales Registered Number: 1072954 Registered office: Mortimer House, 37-

41 Mortimer Street, London W1T 3JH, UK



Nucleosides, Nucleotides and Nucleic Acids

Publication details, including instructions for authors and subscription information: http://www.informaworld.com/smpp/title~content=t713597286

Borano-Nucleotides: New Analogues to Circumvent HIV-1 RT-Mediated Nucleoside Drug-Resistance

Karine Alvarez^a; Jérôme Deval^a; Boulbaba Selmi^a; Karine Barral^a; Joelle Boretto^a; Catherine Guerreiro^b; Laurence Mulard^b; Robert Sarfati^b; Bruno Canard^a

^a CNRS, UMR 6098, Architecture et Fonction des Macromolécules Biologiques, Marseille Cedex 9, France ^b Institut Pasteur, Unité de Chimie Organique, Paris Cedex 15, France

To cite this Article Alvarez, Karine , Deval, Jérôme , Selmi, Boulbaba , Barral, Karine , Boretto, Joelle , Guerreiro, Catherine , Mulard, Laurence , Sarfati, Robert and Canard, Bruno(2005) 'Borano-Nucleotides: New Analogues to Circumvent HIV-1 RT-Mediated Nucleoside Drug-Resistance', Nucleosides, Nucleotides and Nucleic Acids, 24: 5, 419 $-422\,$

To link to this Article: DOI: 10.1081/NCN-200059951 URL: http://dx.doi.org/10.1081/NCN-200059951

PLEASE SCROLL DOWN FOR ARTICLE

Full terms and conditions of use: http://www.informaworld.com/terms-and-conditions-of-access.pdf

This article may be used for research, teaching and private study purposes. Any substantial or systematic reproduction, re-distribution, re-selling, loan or sub-licensing, systematic supply or distribution in any form to anyone is expressly forbidden.

The publisher does not give any warranty express or implied or make any representation that the contents will be complete or accurate or up to date. The accuracy of any instructions, formulae and drug doses should be independently verified with primary sources. The publisher shall not be liable for any loss, actions, claims, proceedings, demand or costs or damages whatsoever or howsoever caused arising directly or indirectly in connection with or arising out of the use of this material.

Nucleosides, Nucleotides, and Nucleic Acids, 24 (5-7):419-422, (2005)

DOI: 10.1081/NCN-200059951



BORANO-NUCLEOTIDES: NEW ANALOGUES TO CIRCUMVENT HIV-1 RT-MEDIATED NUCLEOSIDE DRUG-RESISTANCE

Karine Alvarez, Jerôme Deval, Boulbaba Selmi, Karine Barral, and Joelle Boretto

CNRS, UMR 6098, Architecture et Fonction des Macromolécules Biologiques, Marseille Cedex 9, France

Catherine Guerreiro, Laurence Mulard, and Robert Sarfati • Institut Pasteur, Unité de Chimie Organique, Paris Cedex 15, France

Bruno Canard - CNRS, UMR 6098, Architecture et Fonction des Macromolécules Biologiques, Marseille Cedex 9, France

 $^{-}$ α -Boranophosphates suppress RT-mediated resistance when the catalytic rate of incorporation (k_{pol}) of the analogue 5'-triphosphate is responsable for drug resistance, such as in the case of K65R mutant and ddNTPs, and Q151M toward AZTTP and ddNTPs. This suppression is also observed with BH₃-d4T and BH₃-3TC toward their clinically relevant mutants Q151M and M184V. Moreover, the presence of the borano (BH_3^-) group renders the incorporation of the analogue independent from aminoacid substitutions in RT. To our knowledge, this is the first example of rescue of polymerase activity by means of a nucleotide analogue.

INTRODUCTION

Nucleoside analogues (ddN, AZT, d4T, or 3TC) have been extensively used as antiviral drugs targeting HIV reverse transcriptase (RT).^[1] Each of 5'-triphosphate analogue compete for DNA incorporation with its natural counterpart. The lack of a 3' hydroxyl group promote viral DNA chain termination. However, under therapeutic pressure, the viral RT gene mutates and specifies enzymes bearing substitutions responsible for the loss of drug efficacy. Resistance-associated mutations map at the RT active site and resistance mechanisms are now well characterized.^[2,3] Three major mechanisms have been described: repair of the analogue-terminated DNA, discrimination of the analogue by a decreased affinity

This work was supported by the Agence Nationale de la Recherche contre le SIDA and by Ensemble Contre le SIDA (SIDACTION).

Address correspondence to Bruno Canard, CNRS, UMR 6098, Architecture et Fonction des Macromolécules Biologiques, ESIL-Case 925, 163 Ave., de Luminy, 13288 Marseille Cedex 9, France; E-mail: bruno@afmb.cnrs-mrs.fr

 (K_d) for RT, or decreased polymerization rate (k_{pol}) . One strategy to circumvent resistance is to develop nucleoside analogues that are still potent against mutant RTs.

 α -Boranophosphate nucleosides (α -BH₃-dNTPs) are nucleotide analogues on which one non-bridging oxygen on the α phosphate is replaced by a borano BH₃⁻ group. Here, we summarize results obtained with BH₃-nucleotide analogues as inhibitors of HIV-1 RT, emphasizing their capacity to overcome RT-mediated drug resistance. At α -10

RESULTS AND DISCUSSION

(Rp)- α -BH₃-AZT and (Rp)- α -BH₃-d4T act as DNA chain terminators for RT.^[7] The BH₃⁻ group improves both phosphorylation by nucleotide diphosphate kinase and chain termination efficiency.^[7] We investigated whether α -boranophosphate nucleosides have the capability to affect nucleotide drug resistance through their affinity (K_d) or catalytic rate of incorporation (k_{pol}). We have extented our studies to other interesting available α -boranophosphate nucleosides.

We evaluated the capability of resistance suppression of $\alpha\text{-BH}_3\text{-ddATP}$ and $\alpha\text{-BH}_3\text{-AZTTP}$ towards Q151M_{complex} RT, $^{[8]}$ and the $\alpha\text{-BH}_3\text{-ddATP}$ toward K65R RT. $^{[9]}$ We also evaluated the capability of resistance suppression of $\alpha\text{-BH}_3\text{-d4TTP}$ and $\alpha\text{-BH}_3\text{-3TCTP}$ toward Q151M and M184V RTs, respectively (Table 1). $^{[10]}$

Incorporation of these analogues into DNA by purified RTs were measured comparatively using pre-steady-state kinetics. The nucleotide affinity K_d is calculated as the nucleotide concentration giving half of the maximum incorporation rate k_{pol} . The incorporation efficiency of the nucleotide into DNA (k_{pol}/K_d) is used to calculate the selectivity factor: $(k_{pol}/K_d)_{dNTP}/(k_{pol}/K_d)_{analogue}$. A selectivity factor greater than one means that the enzyme discriminates the analogue over the natural nucleotide. Finally, the resistance of RT to the inhibitor is the ratio between the selectivity of the mutant over the selectivity of the wild-type enzyme.

Data presented in Table 1 indicate that the impact of the BH₃ modification is larger on k_{pol} than on K_d . Using Q151M_{complex} RT, k_{pol} increases up to 70- and 13-fold using α -BH₃-ddATP and α -BH₃-AZTTP, respectively, but K_d remains unchanged and suppression of resistance is observed. α -BH₃-ddATP is a 2-fold better substrate than dATP and inhibits DNA synthesis by K65R RT 153-fold better than ddATP. This complete suppression of resistance is due to an important increase of the catalytic rate constant k_{pol} (20-fold) associated with a better binding affinity K_d (7-fold). α -BH₃-d4TTP is well incorporated (high k_{pol} values) regardless of the enzyme used and there is no discrimination of α -BH₃-d4TTP relative to dTTP. As a consequence, there is no resistance arising from the Q151M_{complex} RT toward borano-phosphate analogues. Similarly, there is a limited resistance to α -BH₃-3TCTP by K65R and M184V. Is it quite remarkable that the incorporation rate of α -BH₃-3TCTP is > 11-fold higher than that of 3TCTP in the case of wild-type RT. The presence of the BH₃- α group on 3TCTP compensates the initial

 $\label{eq:table 1} \textbf{TABLE 1} \ \ Pre-Steady \ State \ Kinetic \ Constants \ of \ dATP/ddATP/BH_3-ddATP/dTTP/AZTTP/\\ BH_3-AZTTP \ \ Incorporation \ by \ WT, \ Q151M_{complex} \ \ and \ K65R \ \ Mutant \ RTs, \ of \ dTTP/d4TTP/BH_3-d4TTP \ \ Incorporation \ by \ WT, \ Q151M \ \ and \ Q151M_{complex} \ \ RT \ \ mutants, \ of \ dCTP/3TCTP/BH_3-3TCTP \ \ \ Incorporation \ by \ WT, \ M184V \ \ and \ K65R/M184V \ \ RT \ \ Mutants$

RT	Nucleotide	Kd (μ M)	$\mathrm{Kpol}\ (\mathrm{s}^{-1})$	Kpol/Kd	${\bf Selectivity}^f$	Resistancegg
WT	dATP^a	7.5	50	6.7	7.4	
	$\mathrm{ddATP}^{b,h}$	8.0	7.2	0.91		
	BH ₃ -ddATP ^a	29.9	22.9	0.75	8.94	
$\rm Q151M_{complex}$	dATP^b	41	99	2.3	63.8	8.62
	ddATP^b	11	0.38	0.036		
	BH_3 -dd ATP^b	7.6	28	3.7	0.62	0.084
WT	dTTP^a	17	13	0.75	0.41	
	$AZTTP^b$	7.1	13	1.8		
	BH_3 -AZTTP e	7.6	18.4	2.4	0.31	
$\rm Q151M_{complex}$	dTTP^b	9.7	7.6	0.79	7.18	17.51
	AZTP^b	9.9	1.1	0.11		
	BH_3 -AZTP b	13	14	1.1	0.72	1.76
WT	$dATP^a$	7.47	50.16	6.71	34.95	
	$\mathrm{ddATP}^{a,h}$	33.8	6.49	0.192		
	BH_3 -dd ATP^a	29.9	22.9	0.75	8.94	
K65R	dATP^a	6.89	11.63	1.69	112.7	3.2
	ddATP^a	47.54	0.71	0.015		
	BH3-ddATPa	6.5	14.9	2.3	0.73	0.0816
WT	dTTP^a	17	13	0.75	1.5	
	$d4TTP^d$	21	11	0.51		
	$\mathrm{BH}_3\text{-}\mathrm{d}4\mathrm{TTP}^d$	19	16	0.85	0.88	
Q151M	dTTP^b	14	17	1.2	4.1	2.8
	$d4TTP^d$	23	6.7	0.29		
	BH_3 -d4 TTP^d	14	20	1.4	0.86	1.0
$\rm Q151M_{complex}$	dTTP^b	9.7	7.6	0.79	7.0	4.7
	$d4TTP^d$	19	2.2	0.12		
	$\mathrm{BH}_3\text{-}\mathrm{d}4\mathrm{TTP}^d$	14	10	0.77	1.0	1.2
WT	dCTP^b	7.9	7.3	0.93	50	
	$3TCTP^c$	2.5	0.047	0.019		
	BH_3 - $3TCTP^d$	4.1	0.54	0.13	7	
M184V	$dCTP^c$	21	9.5	0.45	1500	30
	$3TCTP^c$	88	0.026	0.0003		
	BH_3 - $3TCTP^d$	46	0.36	0.0078	58	8.3
K65R/M184V	$dCTP^c$	19	5.7	0.3	8900	180
	3TCTP ^c	89	0.003	0.000034		
	BH_3 - $3TCTP^d$	82	0.49	0.006	50	6.8

 $[^]a$ Values from, $^{[4]}$ b from, $^{[8]}$ c from, $^{[9]}$ d from, $^{[10]}$ e from, $^{[7]}$ f Selectivity = $[k_{pol}/K_d$ (nucleotide)]/ $[k_{pol}/K_d$ (analogue)], g Resistance = selectivity_{mutant}/selectivity_{WTRT}), b K_d values for ddATP differ for these two sets of experiments because the assay conditions are different. When slow incorporation rates are measured, an excess of enzyme is required to avoid the influence of the off-rate due to multiple turnovers. The latter gives artificially high K_d values such as for ddATP in ref. $^{[4]}$. However, In ref. $^{[4]}$ artificially high K_d(ddATP) values were measured for both WT and K65R, and thus, the general conclusion regarding the cause of resistance is still valid. In subsequent works, this artifact was eliminated using a 2-fold excess of RT over primer/template. $^{[8-10]}$

discrimination by K65R and M184V, mostly through an increase of the catalytic rate k_{pol} . Overall, resistance due to M184V or K65R/M184V is 2.5- to 25-fold reduced when α -BH₃-3TCTP replaces 3TCTP.

Until a counter-example is found, the borano modification acts as a general suppressor of resistance whenever k_{pol} promotes resistance. Indeed, the presence of the BH₃⁻ group does not influence K_d , but provides (or restores) a high k_{pol} value specifically. The α -boranophosphate group at the RT active site renders incorporation independent from the nature of catalytic amino acid side chains.^[10]

REFERENCES

- 1. De Clercq, E. Antiviral drugs: current state of the art. J. Clin. Virol. 2001, 22(1), 73-89.
- Arion, D.; Kaushik, N.; McCormick, S.; Borkow, G.; Parniak, M.A. Phenotypic mechanism of HIV-1 resistance to 3'-azido-3'-deoxythymidine (AZT): increased polymerization processivity and enhanced sensitivity to pyrophosphate of the mutant viral reverse transcriptase. Biochemistry 1998, 37(45), 15908–15917.
- Selmi, B.; Boretto, J.; Sarfati, S.R.; Guerreiro, C.; Canard, B. Mechanism-based suppression of dideoxynucleotide resistance by K65R human immunodeficiency virus reverse transcriptase using an α-boranophosphate nucleoside analogue. J. Biol. Chem. 2001, 276(51), 48466–48472.
- Meyer, P.R.; Matsuura, S.E.; Mian, A.M.; So, A.G.; Scott, W.A. A mechanism of AZT resistance: an increase in nucleotide-dependent primer unblocking by mutant HIV-1 reverse transcriptase. Mol. Cell. 1999, 4, 35– 43.
- Shaw, B.R.; Sergueev, D.; He, K.; Porter, K.; Summers, J.; Sergueeva, Z.; Rait, V. Boranophosphate backbone: a mimic of phosphodiesters, phosphorothioates, and methyl phosphonates. Methods Enzymol. 2000, 313, 226–257.
- Barth, R.F.; Soloway, A.H.; Fairchild, R.G.; Brugger, R.M. Boron neutron capture therapy for cancer, realities and prospects. Cancer 1992, 70, 2995–3007.
- Meyer, P.; Schneider, B.; Sarfati, S.; Deville-Bonne, D.; Guerreiro, C.; Boretto, J.; Janin, J.; Veron, M.; Canard, B. Structural basis for activation of α-boranophosphate nucleotide analogues targeting drug-resistant reverse transcriptase. EMBO J. 2000, 19(14), 3520–3529.
- Deval, J.; Selmi, B.; Boretto, J.; Egloff, M.P.; Guerreiro, C.; Sarfati, S.; Canard, B. The molecular mechanism
 of multidrug resistance by the Q151M human immunodeficiency virus type 1 reverse transcriptase and its
 suppression using α-boranophosphate nucleotide analogues. J. Biol. Chem. 2002, 277(44), 42097 42104.
- Deval, J.; White, K.L.; Miller, M.D.; Parkin, N.T.; Courcambeck, J.; Halfon, P.; Selmi, B.; Boretto, J.; Canard, B. Mechanistic basis for reduced viral and enzymatic fitness of HIV-1 reverse transcriptase containing both K65R and M184V mutations. J. Biol. Chem. 2004, 279(1), 509–516.
- Deval, J.; Alvarez, K.; Selmi, B.; Bermond, M.; Boretto, J.; Guerreiro, C.; Mulard, L.; Canard, B. Mechanistic
 insights into the suppression of drug-resistance by human immunodeficiency virus type-1 reverse transcriptase
 using α-boranophosphate nucleoside analogues. J. Biol. Chem. 2005, in press.