# Structural basis for activation of $\alpha$ -boranophosphate nucleotide analogues targeting drug-resistant reverse transcriptase

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AIDS chemotherapy is limited by inadequate intracellular concentrations of the active triphosphate form of nucleoside analogues, leading to incomplete inhibition of viral replication and the appearance of drug-resistant virus. Drug activation by nucleoside diphosphate kinase and inhibition of HIV-1 reverse transcriptase were studied comparatively. We synthesized analogues with a borano (BH<sub>3</sub>-) group on the α-phosphate, and found that they are substrates for both enzymes. X-ray structures of complexes with nucleotide diphosphate kinase provided a structural basis for their activation. The complex with d4T triphosphate displayed an intramolecular CH...O bond contributing to catalysis, and the  $R_p$  diastereoisomer of thymidine α-boranotriphosphate bound like a normal substrate. Using  $\alpha$ -( $R_p$ )-boranophosphate derivatives of the clinically relevant compounds AZT and d4T, the presence of the \alpha-borano group improved both phosphorylation by nucleotide diphosphate kinase and inhibition of reverse transcription. Moreover, repair of blocked DNA chains by pyrophosphorolysis was reduced significantly in variant reverse transcriptases bearing substitutions found in drug-resistant viruses. Thus, the \alpha-borano modification of analogues targeting reverse transcriptase may be of generic value in fighting viral drug resistance. Keywords: CH...O bond/crystallography/HIV/nucleoside

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#### Introduction

The fight against human immunodeficiency virus (HIV) relies on the association of drugs directed towards two viral enzymes, the reverse transcriptase (Lightfoote *et al.*, 1986) and the protease. Most drugs targeting reverse transcriptase are nucleoside analogues, such as 3'-azido 3'-deoxythymidine (AZT, Zidovudine) and 2',3'-didehydro 2',3'-dideoxythymidine (d4T, Stavudine).

Antiviral nucleoside analogues per se are neither substrates nor inhibitors of reverse transcriptase. They must first be converted into triphosphates and compete with natural 2'-deoxynucleotides (dNTPs) for incorporation into viral DNA by reverse transcriptase. Because they lack a 3'-OH group, DNA chain elongation terminates, which accounts for the observed antiviral effect (Mitsuya et al., 1985; Larder, 1992; Balzarini, 1999). Their potency as a drug depends on their uptake by the infected cell, their phosphorylation by cellular kinases and their ability to block viral DNA synthesis. A single enzyme, nucleoside diphosphate kinase (NDPK), produces the triphosphate derivatives. In vitro, NDPK is 10<sup>4</sup> and 10<sup>3</sup> times less efficient on AZT and d4T diphosphate, respectively, than on thymidine diphosphate (Bourdais et al., 1996; Schneider et al., 1998, 2000). The efficacy of incorporation is limited by the low concentration of the triphosphate analogues relative to dNTPs in the infected cell. The poor activation of the drugs has a dramatic consequence: incomplete suppression of viral DNA synthesis allows the selection of resistance mutations (Larder, 1992). The emergence of variant viruses is the most serious threat against the efficacy of currently used combinations of three anti-retroviral drugs (tritherapies). The advent of second-generation drugs is urgent: ~20 and 3% of HIV-1 isolates in western urban areas have nucleoside- and multidrug-resistant phenotypes, respectively (Wainberg and Friedland, 1998; Boden et al., 1999).

We conducted a structural and biochemical study of nucleoside analogue activation, which gives a basis to the rational design of novel drugs overcoming these limitations. Here we consider analogues where an oxygen of the  $\alpha$ -phosphate is replaced with a borano (BH<sub>3</sub><sup>-</sup>) group yielding  $\alpha$ -( $R_p$ )-boranophosphate. X-ray structures at 1.9 Å resolution show that the analogues bind NDPK like normal substrates. Biochemical data demonstrate that their activation by NDPK is enhanced, and that they provide reverse transcription inhibitors of increased efficiency. Moreover, repair of the analogue-terminated DNA chain by pyrophosphorolysis in drug-resistant reverse transcriptase mutants is greatly impeded.

#### Results

#### Synthesis of nucleotide analogues

We synthesized nucleotide analogues of thymidine, AZT and d4T containing a borano group  $(BH_3^-)$  on the  $\alpha$ -phosphate. The borano group is isoionic to oxygen, and it generates a chiral centre when present on the  $\alpha$ -phosphate of a dNTP (Figure 1). The two diastereo-isomers were separated by reverse-phase high-performance liquid chromatography (HPLC). The same single diastereoisomer of unknown absolute configuration acted

as an efficient substrate for both NDPK and HIV-1 reverse transcriptase.

## Stereochemistry of $\alpha$ -borano TDP in complex with NDPK

To determine the absolute configuration of the active diastereoisomer, we crystallized a complex of α-boranothymidine diphosphate (TDP) with NDPK. NDPK transfers the γ-phosphate of a nucleoside triphosphate onto a nucleoside diphosphate via a phospho-histidine intermediate. Eukaryotic NDPKs are closely related. Human NDPK has a very similar structure to the Dictyostelium enzyme, and essentially the same active site (Dumas et al., 1992; Moréra et al., 1995; Webb et al., 1995). The crystals of Dictyostelium NDPK in complex with α-borano-TDP were isomorphous to a previously determined complex with ADP and aluminium fluoride, where AlF<sub>3</sub> mimicks the y-phosphate of ATP undergoing transfer onto the catalytic histidine (Xu et al., 1997b). The 1.92 Å resolution X-ray structure (Table I) demonstrates that the α-borano analogue binds like ADP and makes the same interactions (Figure 2A). The presence of boron with five electrons instead of oxygen with eight electrons is apparent in the electron density, which is consistently weaker at the  $R_p$  than the  $S_p$  position of the  $\alpha$ -phosphate. Moreover, a  $Mg^{2+}$  ion ligates the  $\alpha$ - and  $\beta$ -phosphates of the analogue in the same way as for natural nucleotides. As only the  $R_{\rm p}$  diastereoisomer could be crystallized and the

**Fig. 1.** Chemical formula of the  $\alpha$ -( $R_p$ )-borano-d4T triphosphate diastereoisomer.

Table I. Summary of crystallographic data collection and refinement statistics

statistics					
d4T triphosphate	α-borano-TDP				
1.85	1.92				
37 405	33 469				
5.0	5.6				
5.6 (32.8)	10.5 (32.7)				
99.3	97.1				
30-1.85	30-1.92				
35 535	32 052				
3446	3446				
346	400				
84	75				
24.3	28				
0.012	0.010				
1.7	1.5				
20.9	22.8				
24.2	27.4				
	1.85 37 405 5.0 5.6 (32.8) 99.3 30–1.85 35 535 3446 346 84 24.3 0.012 1.7 20.9				

 $<sup>{}^</sup>aR_{\text{sym}} = \Sigma \text{hi } |I(\text{h})i - \langle I(\text{h}) \rangle |/\Sigma \text{hi } I(\text{h})i$ . Outer shell of resolution in parentheses

 $\alpha$ -borano group does not interact with the protein, metal ligation by oxygen atoms of the phosphates is likely to determine the stereochemistry of recognition of the  $\alpha$ -borano group by NDPK.

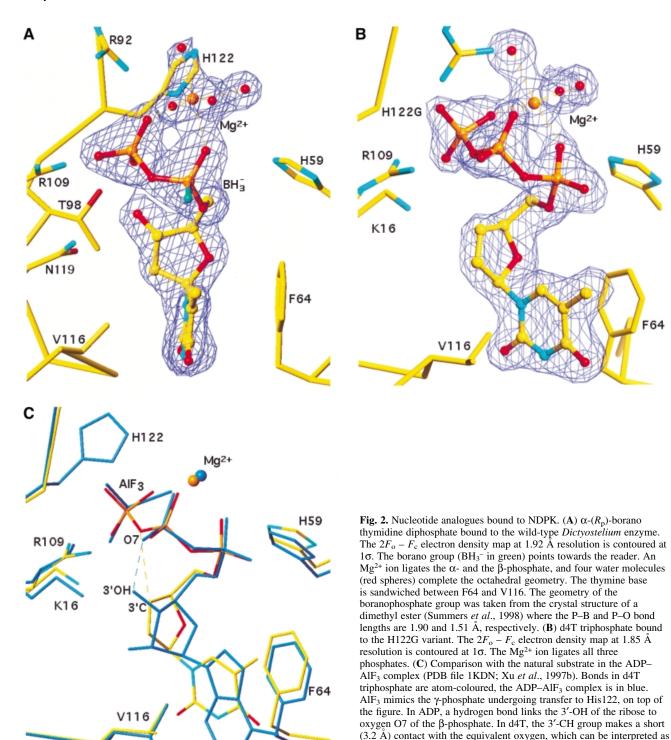
#### A CH...O bond in d4T triphosphate bound to NDPK

To check whether this structural model is valid for antiviral nucleoside analogues used in AIDS therapy, we prepared a complex of d4T triphosphate with NDPK and determined its crystal structure at 1.85 Å resolution. A similar analysis has been performed on AZT derivatives bound to thymidylate kinase (Lavie et al., 1997) and to NDPK (Xu et al., 1997a). The present study makes use of the H122G mutant of the *Dictyostelium* enzyme. This mutant lacks the catalytic histidine and cannot autophosphorylate, but it binds ATP and transfers its γ-phosphate efficiently onto externally supplied imidazole (Admiraal et al., 1999). Again, the complex with d4T triphosphate was isomorphous with the ADP-AlF<sub>3</sub> complex. The analogue binds like ADP, the additional γ-phosphate occupying the AlF<sub>3</sub> position, and the Mg<sup>2+</sup> ion interacts with all three phosphate groups (Figure 2B and C). Interactions with the protein are the same as for ADP-AlF<sub>3</sub>, with an important exception. In a natural substrate, the 3'-OH of ribose or deoxyribose receives two hydrogen bonds from protein groups, and donates one to the oxygen atom that is labelled O7 in Figure 2C and bridges the βand  $\gamma$ -phosphates. The latter bond is particularly important in catalysis (Xu et al., 1997b; Gonin et al., 1999).

Instead of a 3'-OH, d4T carries a double bond in the sugar ring. We find that polar protein groups pack around the double bond making short contacts (3.2–3.4 Å) with the C2' and C3' atoms. The shortest contact is between C3' and O7, the oxygen that bridges the  $\beta$ - and  $\gamma$ -phosphates. The comparison of our structural analysis with NDPK bound to AZT triphosphate suggests a reason why d4T derivatives are better substrates of NDPK than AZT and other analogues lacking a 3'-OH group. In AZT, the bulky 3'-azido substituent interferes with the proper positioning of a lysine side chain (K16 in Figure 2C) that is involved in catalysis (Xu et al., 1997a). With d4T triphosphate, this side chain and all other protein catalytic groups have the same position as in complexes with natural substrates such as ADP, GDP or TDP (Cherfils et al., 1994; Moréra et al., 1995).

Also, we noted the short distance between the modified sugar ring and the bridging oxygen O7. This oxygen is the leaving group when the γ-phosphate is transferred. In a natural substrate, the hydrogen bond to the sugar 3'-OH activates it (Figure 2C). In d4T triphosphate, the 3'-CH group points towards O7 in the plane of the double bond. The distance and geometry of the contact are consistent with a CH...O bond. CH...O bonds are observed in small molecule crystals, in proteins and in DNA (Neidle and Taylor, 1977; Derewenda et al., 1995; Mandel-Gutfreund et al., 1998). Although much weaker than the bond with the 3'-OH of a natural substrate, the CH...O7 bond may explain why d4T diphosphate is phosphorylated 10-fold faster than dideoxyTDP (Schneider et al., 1998, 2000). The dideoxy sugar has methylene groups instead of a double bond and their hydrogens are not oriented properly for an interaction with the bridging oxygen.

 $<sup>{}^{\</sup>dot{b}}R_{\rm crys} = \Sigma h ||F_o| - |F_c||/\Sigma h |F_o|$  calculated with no  $I/\sigma$  cut-off;  $R_{\rm free}$  is derived from 4% of the data set excluded from refinement.



## Activation of $\alpha$ -borano nucleotide analogues by

We examined pre-steady-state kinetic parameters of the phosphorylation of the  $\alpha$ -borano diphosphate derivatives of AZT and d4T by Dictyostelium NDPK. Protein fluorescence is quenched upon phosphorylation of the protein by an NTP substrate and enhanced upon its dephosphorylation by NDP (Deville-Bonne et al., 1996). With AZT diphosphate as the phosphate acceptor, the reaction is 10<sup>4</sup>-fold slower than with TDP (Schneider et al., 1998). When α-borano-AZT diphosphate is tested in this way, the  $S_p$  isomer is inactive but the  $R_p$  isomer is a better phosphate acceptor than AZT diphosphate (Figure 3A). Its phosphorylation follows an exponential time course with a rate constant  $k_{obs}$  that increases with nucleotide concentration. The initial slope yields the catalytic efficiency of the reaction (Schneider et al., 1998), which is 10-fold higher for the α-borano compound (Figure 3B). A 10-fold enhancement of catalytic efficiency is also observed for the  $R_{\rm p}$  isomer of  $\alpha$ -borano TDP relative to TDP (Table II).

a CH...O bond (dashes). The superposition is based on all  $C_{\alpha}$  positions. The figure was made with TURBO (Roussel and Cambillau, 1991).

H59

F64

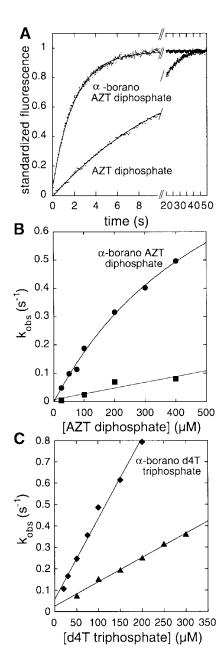


Fig. 3. Enhancement of AZT and d4T activation by NDPK in the presence of the  $R_p$   $\alpha$ -borano group. (A) Kinetics of phosphorylation of AZT diphosphate and α-borano-AZT diphosphate by phosphorylated human NDPK A. The phosphorylated enzyme (1 µM), prepared as described (Deville-Bonne et al., 1996), is mixed with 400 µM AZT diphosphate or α-borano-AZT diphosphate in 50 mM Tris-HCl pH 7.5, 5 mM MgCl<sub>2</sub>, 75 mM KCl, 1 mM DTT and 5% glycerol (final concentrations) at 20°C. The decrease in fluorescence is monitored with a Hi-Tech SF-61DX2 stopped flow device ( $\lambda_{exc}$  = 304 nm, excitation slit = 2 nm, emission filter <320 nm). The solid lines represent the best fit of each curve to a monoexponential. (B) Concentration dependence of the pseudo-first order phosphorylation rate constant  $k_{obs}$ :  $\alpha$ -borano-AZT diphosphate (circles) and AZT diphosphate (squares). Data were analysed according to the classical model of a fast binding reaction followed by a slow, rate-limiting phosphotransfer (Schneider et al., 1998). The catalytic efficiency is  $k_{\text{max}}/K_{\text{D}} = 2000 \text{ M}^{-1}\text{s}^{-1}$  for  $\alpha$ -borano-AZT diphosphate and 200 M<sup>-1</sup>s<sup>-1</sup> for AZT diphosphate. As no saturation is reached, the maximum rate of phosphotransfer,  $k_{max}$ , is not measurable in this experiment. (C) Concentration dependence of the pseudo-first order rate constant  $k_{\rm obs}$  for NDPK A phosphorylation by α-borano-d4T triphosphate (diamonds) and d4T triphosphate (triangles). The catalytic efficiency is  $k_{\text{max}}/K_{\text{D}} = 6000 \text{ M}^{-1}\text{s}^{-1}$  for  $\alpha$ -borano-d4T triphosphate and 800  $M^{-1}s^{-1}$  for d4T triphosphate.

**Table II.** NDPK binding and activation of  $\alpha$ - $(R_p)$ -borano thymidine nucleotide analogues

Nucleotide		Catalytic efficiency $(M^{-1}s^{-1})^a$	$K_{\rm D}~(\mu{\rm M})^{\rm b}$
TTP	_	$1 \times 10^{6}$	1.2
AZT triphosphate	–	80	30.0
	α-borano	375	2.0
d4T triphosphate	–	800	2.0
	α-borano	6000	0.25
TDP	–	$2 \times 10^{6}$	_,
	α-borano	$27 \times 10^{6}$	_
AZT diphosphate	–	200	_
	α-borano	2000	_
d4T diphosphate	–	2600	_
	α-borano	ND	_

 $^{a}$ The phosphotransfer reaction between human NDPK A and thymidine derivatives was studied in both directions as described in Figure 3. Note that the α-borano group increases the catalytic efficiency 4- to 10-fold

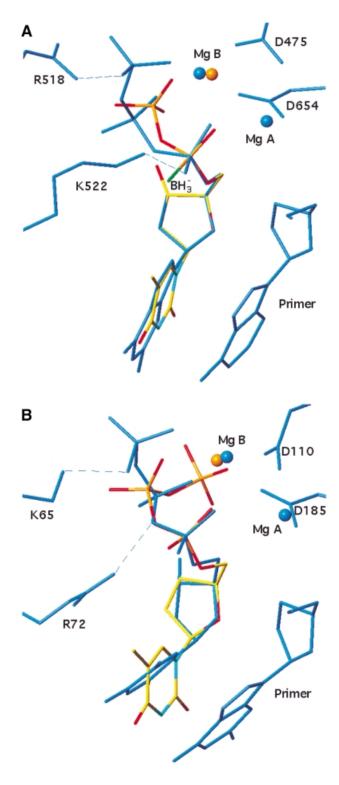
<sup>b</sup>The F64W-H122G variant of *Dictyostelium* NDPK is designed to measure NTP binding in the absence of phosphotransfer. Protein fluorescence (λ<sub>exc</sub> = 310 nm, λ<sub>em</sub> = 330 nm) increases by 50% upon binding saturating amounts of NTP at 20°C in 50 mM Tris–HCl, 5 mM MgCl<sub>2</sub> and 75 mM KCl pH 7.5. K<sub>D</sub> values are estimated by fitting the binding curve to a quadratic equation.

As  $\alpha$ -borano-d4T diphosphate was not available, we used  $\alpha$ -borano-d4T triphosphate to phosphorylate NDPK and study phosphate transfer in the reverse direction. Again, the presence of the  $\alpha$ -borano group in the  $R_p$  position results in an enhancement of the catalytic efficiency of the reaction relative to d4T triphosphate (Figure 3C; Table II).

The binding affinity for the analogues was measured using a variant of *Dictyostelium* NDPK that lacks the catalytic histidine and where a tryptophan replaces the phenylalanine stacking on the base in the active site (F64 in Figure 2B). In this variant, protein fluorescence changes upon nucleotide binding whilst no phosphorylation takes place (Deville-Bonne *et al.*, 1996; Schneider *et al.*, 2000). Binding isotherms show a 10-fold decrease in  $K_D$  for the  $\alpha$ -borano analogue relative to d4T triphosphate (not shown). Similar experiments with TTP, AZT triphosphate and their  $\alpha$ -borano analogues also indicate a 10-fold decrease in  $K_D$  when the borano group is present in the  $R_p$  position (Table II). We conclude that the increase in catalytic efficiency of  $\alpha$ -boranophosphate derivatives results from an improvement in the affinity for the enzyme.

## Inhibition of reverse transcription by $\alpha$ -borano nucleotide analogues

Examination of crystal structures of nucleotide substrates bound either to NDPK (this work) or to DNA polymerases such as reverse transcriptase (Huang *et al.*, 1998) and bacteriophage T7 DNA polymerase (Doublié *et al.*, 1998) shows that their NDP moieties have similar conformations. This is illustrated in Figure 4 by comparing d4T triphosphate from Figure 2B with dideoxyGTP bound to T7 DNA polymerase (Figure 4A; Doublié *et al.*, 1998) and with deoxyTTP bound to reverse transcriptase (Figure 4B;



**Fig. 4.** Conformation of the nucleotide substrate in NDPK, T7 DNA polymerase and HIV reverse transcriptase. Thymidine diphosphate and d4T triphosphate from the complexes in Figure 2 are shown in atom-type coloured bonds superimposed onto, respectively, (**A**) dideoxyGTP in the ternary complex with bacteriophage T7 DNA polymerase–DNA (PDB file 1T7P) and (**B**) deoxyTTP in the ternary complex with HIV reverse transcriptase–DNA (PDB file 1RTD). Least-squares fitting was performed on atom N1 of the base and common atoms in the sugar and the α-phosphate. In T7 polymerase and reverse transcriptase, relevant active site residues, DNA and the ligand are in blue bonds, and blue spheres represent two Mg<sup>2+</sup> ions bound. In NDPK, the red sphere is the single Mg<sup>2+</sup> ion bound to d4T triphosphate. It is located <1 Å away from one of the two Mg<sup>2+</sup> of the polymerases.

**Table III.** Kinetic parameters of wild-type reverse transcriptase for thymidine nucleotide analogues

	TTP	AZT triphosphate		d4T triphosphate	
		_	α-borano	-	α-borano
$K_{\rm m} (\mu M)^{\rm a}$	1.7	2.4	0.2	1.2	0.44
$k_{\rm cat} \ ({\rm s}^{-1})^{\rm a}$	0.05	0.08	0.06	0.055	0.056
$k_{\rm cat}/K_{\rm m}~(\mu{\rm M}^{-1}{\rm s}^{-1})$	0.029	0.033	0.30	0.046	0.127
$K_{\rm D} (\mu \rm M)^{\rm b}$	11 <sup>c</sup>	7.1	7.6	21.3	18.7
$k_{\text{pol}} (s^{-1})^{\text{b}}$	5.4 <sup>c</sup>	12.8	18.4	10.8	16
$k_{\rm pol}/K_{\rm D}~(\mu{\rm M}^{-1}{\rm s}^{-1})$	$0.49^{c}$	1.8	2.4	0.51	0.85

<sup>a</sup>A single-nucleotide incorporation and gel assay was used as described (Reardon and Miller, 1990) except that incubations were at 37°C, and the heteropolymeric DNA primer–template system as described (Canard *et al.*, 1998).  $K_{\rm m}$  and  $k_{\rm cat}$  were obtained using Lineweaver–Burk or Eadie–Hofstee plots in which values of correlation coefficients were >0.98

 $^{\rm b}K_{\rm D}$  and  $k_{\rm pol}$  were obtained as described in Figure 5 (Kati *et al.*, 1992). Standard deviations were <16%.

<sup>e</sup>Determinations carried out at 25°C and taken from Reardon and Miller (1990).

Huang *et al.*, 1998). Common features are the conformation of the  $\alpha$ - and  $\beta$ -phosphates and the position of one of the bound Mg<sup>2+</sup> ions. The similarity is especially striking in the case of T7 polymerase, which is a higher resolution (2 Å) structure than for reverse transcriptase, making details of the nucleotide conformation and the Mg<sup>2+</sup> binding mode more reliable. In all three enzymes, the same oxygen of the α-phosphate ligates the metal, explaining why the other α-phosphate oxygen can be modified without interfering with activation by NDPK or incorporation in DNA by reverse transcriptase.

When the α-borano analogues of AZT and d4T triphosphate were tested for their ability to inhibit reverse transcription (Ueno et al., 1995), potent inhibition was observed only with the  $R_p$  analogue. The fact that only one diastereoisomer is a DNA polymerase substrate is consistent with results obtained by others using DNAdependent DNA polymerases (Eckstein and Thomson, 1995). Chain terminator analogues produce a pattern of DNA polymerization inhibition similar to that of true competitive inhibitors (Reardon and Miller, 1990). The apparent inhibition constant  $K_i$  was determined for each analogue in triplicate. We used Lineweaver–Burk and IC<sub>50</sub> analysis (Copeland, 1996). For AZT triphosphate, the two methods gave  $K_i$  values of 28 and 22 nM; they gave 2.8 and 8.2 nM for the α-borano analogue. For d4T triphosphate, we obtained 35 and 49 nM, and 15 and 20 nM for the  $\alpha$ -borano analogue. Thus,  $K_i$  decreased by a factor of 2.2–10 in the presence of the  $\alpha$ -borano group.

Michaelis–Menten parameters were measured using a primer extension assay and synthetic DNA-primed DNA template (Reardon and Miller, 1990) (Table III). The catalytic efficiency of incorporation ( $k_{\rm cat}/K_{\rm m}$ ) of the α-borano analogues increases 9-fold for α-borano-AZT triphosphate and 3-fold for α-borano-d4T triphosphate. This results from a decrease in the  $K_{\rm m}$  values while  $k_{\rm cat}$  is unchanged (Table III). Using this primer–template system, we performed pre-steady-state kinetics that allow the measurement of  $K_{\rm D}$  for the analogues and their incorporation rate constant  $k_{\rm pol}$  (Kati *et al.*, 1992; Hsieh *et al.*, 1993) (Figure 5). The catalytic efficiency  $k_{\rm pol}/K_{\rm D}$  is ~50% higher when the borano group is present (Table III).

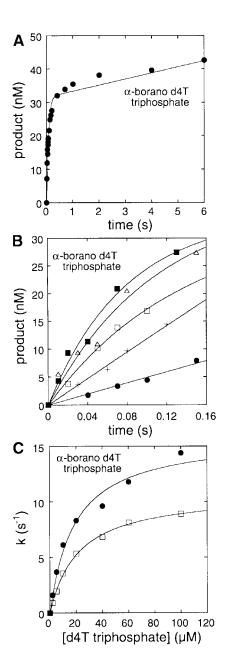


Fig. 5. Pre-steady-state kinetics of α-borano-d4T monophosphate and d4T monophosphate incorporation into DNA as described (Kati et al., 1992). (A) Pre-steady-state kinetics of a single α-borano-d4T monophosphate addition. A 5'-32P-labelled 21mer was annealed to a 31mer DNA template (100 nM) specifying a single thymidine insertion site immediately adjacent to the 3' end of the primer, and reverse transcriptase (30 nM active sites) was allowed to bind (Canard et al., 1998). The reaction was initiated by the addition of 100  $\mu$ M  $\alpha$ -boranod4T triphosphate, 6 mM MgCl<sub>2</sub> in a rapid quench apparatus (Kin Tek, State college, PA) at 37°C, and quenched at various times with 0.3 M EDTA. Products were analysed by denaturing gel electrophoresis and quantitated using photostimulatable plates and a Fuji Imager. Data were fitted to the burst equation: product =  $A(1 - e^{-k}t) + k_{ss}t$ , where A is the amplitude, k is the apparent pre-steady-state rate and  $k_{\rm ss}$  is the steadystate rate. (B) Partial data set of pre-steady-state rates determined as described above using 2.5 (filled circles), 5 (crosses), 10 (open circles), 20 (open triangles) and 60  $\mu M$  (filled squares)  $\alpha$ -borano-d4T triphosphate, 6 mM MgCl<sub>2</sub>. (C) First-order rates determined as described above of  $\alpha$ -borano-d4T triphosphate and d4T triphosphate plotted against nucleotide analogue concentration to determine  $K_D$ ( $\mu$ M) and  $k_{pol}$  (per second) using a hyperbolic fit of the data (Kati et al., 1992).

Unlike NDPK, this is due to an increase in  $k_{pol}$  whereas the affinity is not significantly changed.

## Targeting drug-resistant reverse transcriptase using $\alpha$ -borano analogues

In the crystal structure of reverse transcriptase in complex with DNA and deoxyTTP, the positive charges of R72 and K65 side chains interact with  $\alpha$ - and  $\gamma$ -phosphate groups of the nucleotide, respectively (Figure 4B). As these side chains would be closest to the borano group in an  $R_{\rm p}$ α-borano substrate analogue, the R72A and K65A substitutions were made in order to characterize putative interactions with the  $\alpha$ -borano group. Unexpectedly and unlike wild-type reverse transcriptase, R72A reverse transcriptase discriminates against AZT triphosphate and d4T triphosphate 350- and 13-fold, respectively (Table IV). The activity of R72A reverse transcriptase is at least 25-fold lower than that of wild type (Table IV; Sarafianos et al., 1995), and mutations at position 72 are never observed in viral isolates. K65A reverse transcriptase also discriminates 6- to 7-fold against analogues. Therefore, both R72A and K65A confer resistance to AZT triphosphate and d4T triphosphate in vitro. In contrast, neither variant reverse transcriptase discriminates against α-borano analogues (Table IV). We conclude that an intricate pattern of interactions between residues 72 and 65, the phosphate groups and the sugar 3' position of the substrate nucleotide mediates the AZT triphosphate and d4T triphosphate resistance observed here in vitro.

This prompted us to examine the K65R variant, a substitution found in HIV-1 strains cross-resistant to deoxynucleosides (Gu *et al.*, 1994; Foli *et al.*, 1996). Introduction of an  $\alpha$ -borano group in AZT triphosphate and d4T triphosphate enhances the affinity of the analogue for K65R reverse transcriptase by a factor of 2 and 3, respectively (Table IV), indicating that the  $\alpha$ -borano substitution might also be of interest on other non-thymine nucleoside analogues.

**Table IV.** Inhibition of wild-type and drug-resistant HIV-1 reverse transcriptase by nucleotide analogues

Reverse transcriptase	Wild type	R72A	K65A	K65R
Relative catalytic efficiency <sup>a</sup>	1	0.04	0.46	0.96
AZT triphosphate $K_i$ (nM) <sup>b</sup> $\alpha$ -borano $K_i$ (nM) <sup>b</sup> Ratio	21.8	6500	125	20.5
	8.25	32.5	9	9
	2.6	200	13.9	2.2
d4T triphosphate $K_i$ (nM) <sup>b</sup> $\alpha$ -borano $K_i$ (nM) <sup>b</sup> Ratio	49.2	500	275	60
	20.3	5	27.5	19.5
	2.4	100	10	3

 $^{a}$ Oligo(dT)<sub>12–18</sub>–poly(rA) primer–template, TTP as the nucleotide substrate (Ueno *et al.*, 1995). The catalytic efficiency is measured relative to wild-type reverse transcriptase (0.63  $\mu$ M<sup>-1</sup>s<sup>-1</sup>).  $^{b}$ These  $K_{i}$  values were determined as described (Ueno *et al.*, 1995), from three and two independent experiments for wild-type and variant reverse transcriptases, respectively.

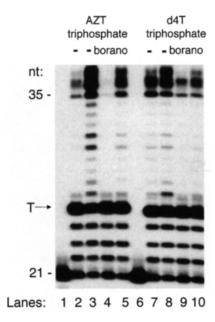
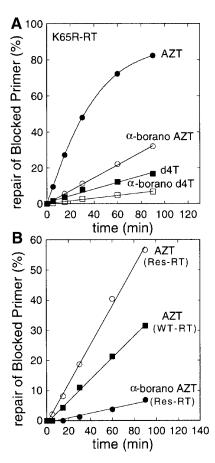


Fig. 6. Pyrophosphorolytic repair of incorporated  $\alpha$ -borano thymidine 5'-monophosphate analogues by drug-resistant reverse transcriptase. Incorporation of α-borano thymidine 5'-triphosphosphate analogues in the presence of pyrophosphate. A 5'-32P-labelled 21mer was annealed to a 35mer DNA template (50 nM) specifying a single thymidine insertion site ('T' arrow) four bases away from the 3' end of the primer. Polymerization was initiated by the addition of K65R reverse transcriptase (100 nM) and nucleotides (50 µM each of dATP, dCTP, dGTP and analogue-TP, and 5 µM TTP) for 15 min at 37°C. The pyrophosphorolytic repair reaction was initiated by the addition of 1 mM pyrophosphate (PP<sub>i</sub>). Aliquots were withdrawn during the time course of the reaction, and products analysed by denaturing gel electrophoresis. This autoradiograph of the gel shows the unextended primer (control, lane 1), and products at zero (even lanes) and 60 min (odd lanes). Over time, the amount of products >25 nucleotides does not change when PP<sub>i</sub> is omitted (not shown), but increases in the presence of PPi due to pyrophosphorolytic repair of the analogueterminated DNA chain followed by DNA polymerization. Lanes 2, 3, 6 and 7, unsubstituted analogues; lanes 4, 5, 8 and 9, α-borano analogues.

## Reverse transcriptase-catalysed pyrophosphorolytic repair of blocked DNA chains

Drug resistance to nucleoside analogues by HIV sometimes correlates poorly with the loss of affinity of the corresponding nucleotide for reverse transcriptase in vitro (Lacey et al., 1992; Larder, 1992; Gu et al., 1994; Arion et al., 1996). For instance, AZT resistance has been shown to involve a reverse transcriptase-catalysed repair mechanism by excision of the analogue from the viral DNA chain (Arion et al., 1998; Meyer et al., 1999). The DNA 3' end can be unblocked by the nucleophilic attack of either pyrophosphate or ATP. Repair of AZT-terminated DNA by the two drug-resistant reverse transcriptases used above was examined using a gel assay (Figure 6). During DNA polymerization in the presence of added pyrophosphate, the appearance of full-length extension products past the analogue insertion site indicates nucleotide incorporation made possible through pyrophosphorolysis of the terminal AZT nucleotide. Strikingly, the rate of the excision reaction by K65R reverse transcriptase decreases 5-fold in the presence of an  $\alpha$ -borano group (Figure 7A). A similar effect is observed when comparing d4T triphosphate and its α-borano counterpart. When the variant reverse transcriptase carrying the AZT resistance muta-



**Fig. 7.** Time course of the pyrophosphorolytic repair reaction under reaction conditions shown above. (**A**) Variant K65R reverse transcriptase. Polymerization products were quantitated from an experiment similar to that in Figure 6 using photostimulatable plates and a Fuji Imager. The percentage repair of blocked primer is the ratio of extension products >25 nucleotides over those >24 nucleotides. (**B**) Same as in (A) but using wild-type and AZT-resistant (D67N/K70R/T215F/Y/K219Q) reverse transcriptase in conjunction with either AZT triphosphate or α-borano-AZT triphosphate as indicated.

tions D67N/K70R/T215F/K219Q is used, increased pyrophosphorolysis of the unsubstituted analogue (~2-fold) is observed (Figure 7B). Again, incorporation of the α-borano analogue reduces pyrophosphorolysis 9-fold.

We conclude that the presence of the  $\alpha$ -borano group confers increased stability towards pyrophosphorolytic repair to the blocked DNA chain.

#### **Discussion**

The present study shows that the mode of binding of the nucleotide substrate has common features in NDPK and in DNA polymerases such as that of bacteriophage T7 and HIV reverse transcriptase. The two classes of enzymes, which are otherwise very different, catalyse a nucleophilic attack on either the  $\gamma$ -phosphate (NDPK) or the  $\alpha$ -phosphate (the polymerases) of the substrate nucleotide. The common features of the bound substrate observed in X-ray structures make a modification of the  $\alpha$ -phosphate compatible with both the activation into a triphosphate and its incorporation into the growing DNA chain. The substitution of a BH<sub>3</sub>- group for an oxygen yieding an  $\alpha$ -( $R_p$ )-phosphate actually enhances the rate of phosphorylation by NDPK and makes the product a more

efficient inhibitor of reverse transcriptase. In contrast, the  $S_p$  derivative is inactive.

Our study shows that stereospecificity originates from the requirement that the nucleotide analogue interacts with a Mg<sup>2+</sup> ion, which has the same ligation geometry in NDPK as in reverse transcriptase. This requirement applies to natural substrates such as thymidine nucleotides, and also to the  $\alpha$ -boranophosphate derivatives of AZT and d4T, two thymidine analogues that are major drugs against AIDS. In addition, we find that the sugar modification in d4T makes the diphosphate derivative a better substrate for NDPK than AZT or dideoxy analogues. Whilst all these analogues lack a 3'-OH group involved in catalysis by NDPK, the geometry of the C2'-C3' double bond in d4T allows the formation of an intramolecular CH...O hydrogen bond that partly substitutes for the missing 3'-OH...O. The 10-fold rate enhancement is highly significant, and may suffice to change the rate-limiting step for activation, i.e. the formation of the monophosphate form for d4T and the di- and triphosphate form for AZT (Lavie et al., 1997; Schneider et al., 2000). Moreover, the implication of a CH...O bond in enzymatic catalysis is a rare observation deserving notice in this context.

In the NDPK reaction, the  $\alpha$ -phosphate is involved only through its contribution to Mg<sup>2+</sup> ligation. We find that its modification into a boranophosphate enhances catalysis if it does not prevent metal binding, i.e. for the  $R_p$  diastereoisomer. The affinity of the enzyme for  $\alpha$ -( $R_p$ )-boranophosphate derivatives is improved by an order of magnitude, increasing the catalytic efficiency for the natural substrate TDP, and also for AZT and d4T analogues. Taking place *in vivo*, this 10-fold increase in efficiency would be of great value in the case of AZT, which is converted very poorly into a triphosphate by cellular enzymes.

The better activation by NDPK is only one of the elements making α-boranophosphate nucleotide analogues attractive. The major limitation of current therapeutic protocols against AIDS is the emergence of resistant reverse transcriptase variants in viral isolates. Novel nucleoside analogues should either elicit less resistance or retain activity on reverse transcriptase variants. To design such second-generation drugs, nucleoside activation and reverse transcription inhibition should be taken as a whole, as the low concentration of the active form of the drugs in HIV-infected cells contributes to the emergence of drug resistance. Due to incomplete inhibition of viral replication, escape mutants are selected quickly, taking over wild-type viral populations, and conferring a drug resistance phenotype to the virus. It is shown here that the presence of a borano group on the α-phosphate of both AZT and d4T derivatives results in an enhanced activation by NDPK and an enhanced incorporation into DNA by reverse transcriptase.

Moreover, α-borano derivatives exhibit increased stability towards repair mechanisms that contribute to drug resistance. Resistance observed *in vivo* cannot be attributed simply to a loss of affinity of reverse transcriptase for nucleotide analogues. Mutant reverse transcriptase isolates from AZT-resistant viruses often display little *in vitro* resistance to inhibition of DNA synthesis. Rather, the hallmark of AZT resistance is an increased pyrophosphorolysis allowing repair of the blocked DNA chain. The

reason may be a longer lifetime of the reverse transcriptase–DNA complex allowing pyrophosphorolysis, and also polyphosphate synthesis, to proceed at the active site (Arion *et al.*, 1998; Canard *et al.*, 1998; Meyer *et al.*, 1999). Consistent with these findings, AZT-resistant reverse transcriptase (D67N/K70R/T215F/K219Q) exhibits an increased processivity of DNA synthesis (Caliendo *et al.*, 1996). Our assay demonstrates that pyrophosphorolysis increases 2-fold in AZT-resistant reverse transcriptase, and that it is greatly reduced after incorporation of the  $\alpha$ -borano analogue. Therefore, compounds of this class are good candidates to target selectively reverse transcriptases made drug resistant by mutations enhancing repair of analogue-terminated DNA chains.

In contrast, mutations such as M184V or L74V leading to resistance to dideoxynucleotides, d4T or 3TC tend to reduce processivity (Back *et al.*, 1996; Sharma and Crumpacker, 1999). Nevertheless, α-borano analogues may still prove useful in that case. Resistance may be due to the analogue-terminated DNA chain being released more readily from the enzyme and made available for repair by cellular exonucleases. As the 3′–5′ boranophosphodiester bond is resistant to cleavage by nucleases (Eckstein and Thomson, 1995; He *et al.*, 1999), the α-borano group should have a protective effect against repair.

These experiments suggest that  $\alpha$ -borano analogues may be of use *in vivo* against variant viruses that have become resistant by several repair mechanisms. The design of an  $\alpha$ -borano d4T diphosphate pro-drug using nanoparticle-based biovectors and/or phosphotriester nucleotide analogues is being explored. It would bypass two steps of intracellular phosphorylation, including the step catalysed by thymidine kinase that appears to be rate limiting in forming d4T triphosphate. More generally, boranophosphate nucleotide analogues should be of interest in the study of all enzymes carrying out phosphotransfer reactions, of which NDPK and reverse transcriptase are two examples.

#### Materials and methods

#### Synthesis of nucleotide analogues

AZT di- and triphosphate were prepared from AZT monophosphate as described (Hoard and Ott, 1965). α-borano-AZT 5'-diphosphate was synthesized according to the procedure described (Tomasz et al., 1992), but the phosphitylation step of the nucleotide was performed as described (Meier et al., 1992). α-borano-AZT 5'-triphosphate and α-borano-d4T 5'-triphosphate were synthesized as described (He et al., 1998). NMR assignments are available upon request. Diastereoisomers were purified (>99%) by reverse-phase HPLC on a C<sub>18</sub> column using a 0–15% acctonitrile gradient in 10 mM triethyl ammonium acetate buffer pH 6.8.

#### Enzyme purification

Human NDPK A was purified as a recombinant protein overexpressed in *Escherichia coli* (BL21-DE3). Plasmid pJC20 was a kind gift from M.Konrad (Göttingen, Germany). After cell breakage in a French press, and centrifugation (10 000 g, 20 min), the cell extract was treated with 0.15% polymin P (BASF) in order to precipitate most of the nucleic acids, then NDPK A was purified by two successive steps of ion-exchange chromatography on a POROS HQ column (PerSeptive Biosystem; 2.6 × 12 cm) with a linear KCl gradient (0.02–1 M) in 50 mM Tris–HCl pH 8.1, 1 mM EDTA and 1 mM dithiothreitol (DTT), followed by a ceramic hydroxyapatite column (American International Chemical; 1.6 × 10 cm) with a linear potassium phosphate gradient (0.01–1 M) in 10 mM potassium phosphate pH 6.5 and 1 mM DTT. After concentration and desalting, the enzyme was dialysed extensively against

50 mM Tris-HCl pH 7.5, 1 mM DTT, 20 mM KCl and 50% glycerol, then stored at -20°C. NDPK A was >97% pure as judged by SDS-PAGE.

The double mutant F64W-H122G NDPK from *Dictyostelium* was overexpressed in *E.coli* (XL1-Blue) and purified as previously described for the single mutant F64W NDPK. Enzyme concentration expressed as 17 kDa subunits was determined either colorimetrically or using an absorbance coefficient of  $\Delta A_{280} = 0.85$  for a 1 mg/ml solution. F64W-H122G mutant NDPK had no measurable NDPK activity and a very weak ATPase activity ( $k_{cal}/k_m = 450 \text{ M}^{-1}\text{s}^{-1}$ ). The affinity of NDP and NTP derivatives for NDPK was determined by following the variation of the intrinsic fluorescence upon nucleotide binding as described (Schneider *et al.*, 1998). The correction for the contribution of ATPase activity was <2%.

HIV-1 reverse transcriptases were expressed in *E.coli*, and p66–p66 homodimers were purified as described (Canard *et al.*, 1999). Wild-type reverse transcriptase used in pre-steady-state kinetic experiments was purified as p66–p51 heterodimers as described (Canard *et al.*, 1998).

#### Crystallographic methods

Crystals of the H122G variant of *Dictyostelium* NDPK (Admiraal *et al.*, 1999) grew in drops containing 19% PEG 400, 50 mM Tris–HCl pH 7.5, 20 mM MgCl<sub>2</sub>, 6 mg/ml protein and 7.5 mM d4T triphosphate over wells containing 38% PEG 400. They belong to space group  $P3_121$  with a=b=70.02 Å and c=152.10 Å and a trimer in the asymmetric unit. Diffraction data were collected from a cryocooled crystal at  $\lambda=1.364$  Å on station D41 of the LURE-DCI synchrotron radiation source (Orsay, France) using an MAResearch Image plate detector, and processed using the HKL program package (Otwinowski and Minor, 1997). The initial model was the protein moiety of the ADP–AIF<sub>3</sub> complex (PDB file 1KDN). It was refined with CNS using anisotropic overall *B*-factors and a bulk solvent mask (Brünger *et al.*, 1998). Statistics on data collection and the refined model are given in Table I.

Co-crystals of α-borano-TDP with wild-type Dictyostelium NDPK were obtained by seeding with crystals of the ADP-AlF3 complex (Xu et al., 1997b). The drops contained 20% PEG 550 monomethylester, 50 mM Tris-HCl pH 7.5, 25 mM MgCl<sub>2</sub>, 7.5 mg/ml protein and 11 mM  $\alpha$ -borano-TDP over wells containing 28% PEG 550 in the same buffer. The first crystals were then used for macro-seeding under the same conditions. They belong to space group  $P3_121$  with a = b = 70.36 Å and c = 153.79 Å and a trimer in the asymmetric unit. Before flash-freezing in liquid propane, PEG 550 was replaced with 39% PEG 400. A flash-frozen crystal was maintained at 100 K in an MSC cryostream while diffraction data were collected on a rotating anode source (Rigaku) using the R-Axis IIc image plate system. Diffracted amplitudes were indexed and integrated using MOSFLM (Leslie, 1992) and scaled with SCALA of the CCP4 package. All subsequent steps were the same as for the  $\alpha$ -borano-TDP complex. For model refinement with CNS, the geometry of the boranophosphate group was taken from the crystal structure of a dimethyl ester (Summers et al., 1998) where the P-B and P-O bond lengths are 1.90 and 1.51 Å, respectively. Statistics on data collection and the refined model are given in Table I.

### Nucleotide analogue incorporation by reverse transcriptase and pyrophosphorolysis assays

Standard reverse transcriptase and nucleotide selectivity assays were as described (Canard et al., 1999). Analysis of data was performed according to published procedures (Henderson, 1972; Cheng and Prusoff, 1973; Copeland, 1996) as described in the text. The primer-template system used for primer extension assays has been described (Canard et al., 1998). It was used for both steady-state and pre-steady-state kinetic assays. Experimental conditions are given in the figure legends. The primer-template system used for pyrophosphorolysis assays was a 21mer (5'-ATACTTTAACCATATGTACC-3') annealed to a 35mer (5'-GGT-CCGTTGCATGCGGGATACATATGGTTAAAGTAT-3') DNA template. The single template 'A' specifying a single thymidine insertion site four bases away from the 3' end of the primer is shown in bold. Experimental conditions are given in the figure legends.

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#### Note added in proof

Co-ordinates have been deposited at the Protein Data Bank (accession Nos 1F6T and 1F3F).