## Endogenous Reverse Transcription Assays Reveal High-Level Resistance to the Triphosphate of  $(-)2'$ -Dideoxy-3'-Thiacytidine by Mutated M184V Human Immunodeficiency Virus Type 1

YUDONG QUAN,<sup>1</sup> ZHENGXIAN GU,<sup>1</sup> XUGUANG LI,<sup>1</sup> ZHUO LI,<sup>1</sup> CASEY D. MORROW,<sup>2</sup> AND MARK A. WAINBERG<sup>1,3\*</sup>

*McGill University AIDS Centre, Lady Davis Institute, Jewish General Hospital,*<sup>1</sup> *and Departments of Medicine and Microbiology, McGill University,*<sup>3</sup> *Montreal, Quebec, Canada, and Department of Microbiology, University of Alabama at Birmingham, Birmingham, Alabama 35294-0007*<sup>2</sup>

Received 9 October 1995/Accepted 4 April 1996

**Kinetic analysis showed that the** *Ki* **values and the** *Ki /Km* **ratios for mutated, recombinant M184V human immunodeficiency virus type 1 reverse transcriptase (RT) for (**2**)2**\***-dideoxy-3**\***-thiacytidine triphosphate (3TCTP) were 35-fold higher than the equivalent values for wild-type RT but only about twice as high as the equivalent values for each of the triphosphates of ddC (ddCTP) and ddA (ddATP). Fully endogenous RT assays showed that viruses containing the M184V substitution were highly resistant to 3TCTP, with an increase in the 50% inhibitory concentration of 250-fold in comparison with wild-type recombinant virus.**

Although mutations in the human immunodeficiency virus type 1 (HIV-1) reverse transcriptase (RT) are responsible for resistance to nucleoside analogs (10, 14, 18, 22), the molecular mechanisms involved are not well understood. For example, RT isolated from 3'-azido-3'-deoxythymidine (AZT)-resistant variants of HIV-1 displayed the same level of sensitivity to AZT triphosphate (AZTTP) in cell-free kinetics and endogenous enzyme assays as did wild-type (wt) material (17). In contrast, both recombinant L74V RT and K65R RT are less sensitive to each of ddITP, ddCTP, ddATP, and AZTTP than is wt RT in such assays (13, 20).

We have pursued this subject further by using fully endogenous RT reactions and enzyme kinetics to compare wt viruses and enzymes with those containing the M184V mutation. The fully endogenous RT reaction may be more physiologically relevant than enzyme kinetics assays, since virion RNA acts as a template and since viral proteins other than RT may influence the synthesis of viral DNA products (1, 7). M184V is located in the "palm" subdomain of RT, within a conserved YMDD motif thought to make up part of the polymerase active site (4, 16). Viruses that contain this mutation display high-level resistance (i.e., 250- to 1,000-fold) to  $(-)2'$ -dideoxy- $3'$ -thiacytidine ( $3TC$ ) (10, 14, 21, 22) and low-level resistance (i.e. 3- to 10-fold) to each of ddC and ddI in tissue culture (10, 14). We now report that the endogenous RT assay is a more sensitive means of determining levels of HIV resistance to dideoxynucleoside triphosphates (ddNTPs) than are enzyme kinetics measurements, at least in regard to 3TCTP.

**M184V and wt RTs have similar RDDP and DDDP activity.** Four homopolymeric templates, i.e.,  $poly(rA-dT_{12-18})$ , poly  $(rC-dG_{12-18})$ , poly  $(rI-dC_{12-18})$ , and poly  $(dC-dG_{12-18})$ , and one heteropolymeric RNA template were used to study RNAand DNA-dependent DNA polymerase activities. The AZTTP used for this work was a gift of Wellcome Laboratories, Research Triangle Park, N.C. 3TC and 3TCTP were gifts of Glaxo Group Research, Greenford, United Kingdom. Other ultrapure dNTPs, ddNTPs, and homopolymer template/primers (t/p), i.e., poly (rC)/oligo (dG)<sub>12–18</sub>, poly(dC)/oligo(dG)<sub>12–18</sub>, and poly(rA)/oligo(dT)<sub>12–18</sub> were obtained from Pharmacia Biotech, Inc., Montreal, Canada. Poly(rI)/oligo(dC)<sub>12-18</sub> was prepared from poly(rI) and oligo(dC)<sub>12–18</sub> (Pharmacia Bio-<br>tech). [<sup>3</sup>H]dNTPs (22 Ci/mmol) and [α-<sup>32</sup>P]dNTPs (3,000 Ci/ mmol) were purchased from ICN Biomedical Inc., Montreal, Canada, and Du Pont Inc., Mississauga, Canada. RNA template was prepared, using a MEGAscript transcription kit (Ambion, Austin, Tex.), from linearized plasmid pHIV-PBS, which consists of a 497-bp HIV-1 sequence spanning the R region of HIV-1 long terminal repeat and a portion of *gag* (2). Heteropolymeric RNA t/p was prepared as described previously (13) by annealing RNA template and a chemically synthesized 18-nucleotide DNA primer, complementary to the HIV-1 primer-binding sequence. Recombinant wt and mutated RTs and relevant plasmids were expressed and purified as described previously (24). RNA-dependent DNA polymerase (RDDP) and DNA-dependent DNA polymerase (DDDP) activities were measured as described previously (12, 13). The specific activities of the recombinant purified wt and M184V RT molecules used in this study were similar.

Table 1 shows that the two enzymes possessed similar *Km* and *V* values for each of the four dNTPs with all of the t/p used. This indicates that they had similar affinities for dNTPs. We considered that the location of codon 184 in the YMDD motif, thought to be part of the polymerase active site (16), might have affected the affinity of the enzyme for template. This possibility was examined by using both homopolymeric and heteropolymeric templates, yet no significant differences in *Km* values were observed between the two enzymes (data not shown).

**M184V RT displays resistance to 3TCTP.** The  $K_i$  values for different ddNTPs, including AZTTP and 3TCTP, are given in Table 2. The *K<sub>i</sub>* value of 3TCTP for M184V RT, with RNA heteropolymeric  $t/p$ , was 41.4  $\mu$ M, an increase of about 35-fold over that for wt RT  $(1.15 \mu M)$  (Table 2). The  $K_i$  value of

<sup>\*</sup> Corresponding author. Mailing address: Lady Davis Institute— Jewish General Hospital, 3755 Chemin Côte-Ste-Catherine, Montreal, Quebec, Canada H3T 1E2. Phone: (514) 340-8260. Fax: (514) 340- 7537.

$t/p^a$	Substrate	$K_m$ ( $\mu$ M) <sup>b</sup>		$L^{b,c}$	
		wt	M184V	wt	M184V
$Poly(rA-dT_{12-18})$	<b>dTTP</b>	$2.47 \pm 0.09$	$2.54 \pm 0.14$	$5.47 \pm 0.19$	$5.28 \pm 0.61$
Poly(rC-dG <sub>12-18</sub> )	dGTP	$1.99 \pm 0.24$	$3.71 \pm 0.09$	$4.02 \pm 0.35$	$4.47 \pm 0.32$
$Poly(dC-dG12-18)$	dGTP	$1.37 \pm 0.19$	$1.49 \pm 0.14$	$2.75 \pm 0.22$	$3.01 \pm 0.43$
Poly (rI-d $C_{12-18}$ )	dCTP	$28.65 \pm 1.44$	$26.35 \pm 0.31$	$7.30 \pm 0.47$	$7.48 \pm 0.52$
Heteropolymeric RNA	<b>dTTP</b>	$1.22 \pm 0.28$	$1.42 \pm 0.19$	$2.15 \pm 0.68$	$2.01 \pm 0.34$
	dCTP	$3.58 \pm 0.21$	$3.54 \pm 0.32$	$1.85 \pm 0.45$	$1.78 \pm 0.30$
	dGTP	$0.23 \pm 0.01$	$0.27 \pm 0.06$	$0.90 \pm 0.05$	$0.97 \pm 0.04$
	dATP	$3.36 \pm 0.33$	$4.39 \pm 0.30$	$1.54 \pm 0.16$	$1.67 \pm 0.24$

TABLE 1.  $K_m$  and *V* values of dNTPs for wild-type and M184V RTs

 $a$  In experiments with homopolymeric t/p, the latter were used at a concentration of 0.1 U/ml. For heteropolymeric RNA, a fixed concentration of 18 nM t/p was used and dNTP, for which the  $K_m$  value was to be determined, was added at a variety of concentrations. For heteropolymeric t/p, three dNTPs, other than the one to be the system of  $\pi$  and  $\pi$  and  $\pi$  and  $\pi$  and  $\pi$  and

<sup>b</sup> Values are means  $\pm$  standard deviations for at least three separate experiments.<br>
<sup>c</sup>V is expressed as nanomoles of dNMP incorporated per 30 min per microgram of RT.

ddATP, the active intracellular form of ddI, for M184V RT with heteropolymeric templates was threefold higher than that for wt RT. The  $K_i$  for ddCTP was increased only twofold. For AZTTP, ddGTP, and ddTTP, a twofold increase in  $K_i$  was also observed for M184V RT. No differences in the  $K_i$  or  $K_i/K_m$ values of ddITP were detected for wt and M184V RTs.

**M184V-containing viruses are resistant to 3TCTP in endogenous RT reactions.** MT-2 cells were used to generate viruses containing either the M184V or K65R substitution in RT as described previously (14). For endogenous RT reactions, viruses were harvested from the culture fluids of  $2 \times 10^8$  infected MT-2 cells as follows: low-speed centrifugation of cells at  $1,000 \times g$  for 30 min, and ultracentrifugation of supernatant fluids at 88,000  $\times g$  for 1 h at 4<sup>o</sup>C. Pelleted viruses were resuspended in 0.2 ml of TN buffer (10 mM Tris [pH 7.8], 100 mM NaCl), further purified as described previously (23) with 2-ml columns of Sephacel S-1000 (Pharmacia, Montreal, Canada) that had been prebalanced with TN buffer, and eluted with TN buffer. Purified viruses were quantified and kept at  $4^{\circ}$ C until use.

Endogenous RT assays were modified from previously described procedures (3, 8) and were performed with a total volume of 30  $\mu$ l containing 50 mM Tris (pH 7.8), 5 mM MgCl<sub>2</sub>, 60 mM KCl, 10 mM dithiothreitol, 10 mM NaCl, 1 mM ethylene glycol-bis( $\beta$ -aminoethyl ether)-*N*,*N*,*N'*,*N'*-tetraacetic acid (EGTA), 0.1% Nonidet P-40, 0.4 mM each dATP, dGTP and dTTP, 10  $\mu$ M dCTP, 10  $\mu$ Ci of [ $\alpha$ -<sup>32</sup>P]dCTP, 20  $\mu$ l of purified HIV normalized on the basis of p24 content (i.e., 280 ng of p24, representing approximately  $2 \times 10^7$  cpm of RT activity per ml per reaction), and variable concentrations of inhibitors. After 6 h at  $39^{\circ}$ C, reactions were terminated by addition of an equal volume of stop buffer (1% sodium dodecyl sulfate, 50 mM EDTA, 0.2 M NaCl). The reaction mixture was digested with 20  $\mu$ g of protease at 56 $\degree$ C for 30 min, extracted with phenol-chloroform, and precipitated with ethanol. Products were boiled and separated on a 1% denaturing agarose gel (20 mM NaOH, 1 mM EDTA), visualized by autoradiography (Fig. 1), and analyzed by molecular imaging (Bio-Rad, Mississauga, Canada) (Fig. 2).

We found that the 50% inhibitory concentration  $(IC_{50})$  of 3TCTP was 1,970  $\mu$ M for M184V versus 7.8  $\mu$ M for wt virus, an increase of about 250-fold (Fig. 2). The recombinant K65R virus displayed less resistance than the M184V virus to 3TCTP in endogenous RT reactions (Fig. 1 and 2), i.e., a 13-fold difference in comparison with wt virus (Fig. 2). Similar results were obtained on at least three separate occasions.

Mutated RTs that contain a YVDD instead of a YMDD motif almost fully retained DNA polymerase activity, consistent with a recent report that also documented a slight decrease in the processivity of mutated M184V RT (5). Viruses





*a* The concentration of dNTPs was 7  $\mu$ M, and that of t/p was 0.1 U/ml. Inhibitor was added at variable concentrations. *b* Values are means determined on the basis of at least three separate experiments.



FIG. 1. Analysis of endogenous RT reactions. Concentrations of 3TCTP from the left for wt viruses were 0, 0.6, 4.0, 20, 60, and 120  $\mu$ M. For M184V and K65R viruses, concentrations of 3TCTP were 0, 6, 40, 200, 600, and 2,400  $\mu$ M.

that contain the M184V substitution replicate almost as well as wt viruses do (19, 24). However, this mutation confers highlevel resistance to 3TC and 3TCTP in tissue culture (10, 14, 21, 22) and cell-free assays, respectively.

The products of our endogenous RT reaction must contain both minus- and plus-strand DNA, since substantial amounts of this material migrated between 9.4 and 23 kb on a neutral agarose gel (results not shown), a pattern similar to that previously reported for murine leukemia virus (11). In contrast (8, 17), most DNA products migrated faster than 4.3 kb on denaturing gels (Fig. 1), a result attributable to the fact that DNA products were partially double-stranded and/or because circular intermediate complexes formed during reverse transcription (11). Upon denaturation, relatively short single-stranded DNA molecules, including plus strands initiated at multiple sites (6), would be separated from minus strands and would move faster than intermediate complexes.

The similarity in  $V$  and  $V/K_m$  values between wt and M184V RTs indicates that the two enzymes possess comparable catalytic efficiencies. Furthermore, the M184V mutation may not affect the ability of the enzyme to bind to template, because the  $K<sub>m</sub>$  of M184V RT for different t/p was similar to that of wt (results not shown). Recently, another "helix clamp" structure was proposed to be responsible for template binding (15).



FIG. 2. Inhibition of endogenous RT reactions by 3TCTP. RT activities are expressed as percentages of those of controls performed in the absence of drug. Data are expressed as mean values of three separate experiments.

In contrast, significant differences in  $K_i$  values were shown to be associated with M184V with regard to 3TCTP (Table 2). M184V RT had a 35-fold-elevated  $K_i$  of 41.4  $\mu$ M compared with 1.15  $\mu$ M for wt. The  $K_i/K_m$  ratio, useful in determining the recognition of dNTP and ddNTP by an enzyme, was also increased by about 35-fold (11.69 for M184V versus 0.32 for wt RT), in agreement with results of earlier studies (9). As shown here, endogenous RT reactions showed even greater differences between M184V-containing and wt viruses, i.e., a 252 fold elevated  $IC_{50}$ . Thus, endogenous RT assays may, in some cases, represent a more sensitive means of determining the resistance to nucleoside analog triphosphates than kinetics assays performed with purified RTs. The former system may also be of greater physiological relevance, since viral RNA genome serves as the template and other viral proteins can interact with RT in the reaction.

This research was supported by grants to Mark A. Wainberg from the Medical Research Council of Canada and by Health and Welfare Canada. Mark A. Wainberg is a National AIDS Scientist of Health Canada.

## **REFERENCES**

- 1. **Allain, B., M. Lapadat-Tapolsky, C. Berlioz, and J. L. Darlix.** 1994. Transactivation of the minus-strand DNA transfer by nucleocapsid protein during reverse transcription of the retroviral genome. EMBO J. **13:**973–981.
- 2. **Arts, A. J., X. Li, Z. Gu, L. Kleiman, M. A. Parniak, and M. A. Wainberg.** 1994. Comparison of deoxyoligonucleotide and tRNA<sup>Lys3</sup> as primers in an endogenous human immunodeficiency virus-1 in vitro reverse transcription/ template-switching reaction. J. Biol. Chem. **269:**14672–14680.
- 3. **Borroto-Esoda, K., and L. R. Boone.** 1994. Development of a human immunodeficiency virus-1 in vitro DNA synthesis system to study reverse transcriptase inhibitors. Antiviral Res. **23:**235–249.
- 4. **Boyer, P. L., A. L. Ferris, and S. H. Hughes.** 1992. Cassette mutagenesis of the reverse transcriptase of human immunodeficiency virus type 1. J. Virol. **66:**1031–1039.
- 5. **Boyer, P. L., and S. H. Hughes.** 1995. Analysis of mutations at position 184 in reverse transcriptase of human immunodeficiency virus type 1. Antimicrob. Agents Chemother. **39:**1624–1628.
- 6. **Charneau, P., M. Alizon, and F. Clavel.** 1992. A second origin of plus-strand synthesis is required for optimal human immunodeficiency virus replication. J. Virol. **66:**2814–2820.
- 7. **Darlix, J. L., C. Gabus, M.-T. Nugeyre, F. Clavel, and F. Barre´-Sinoussi.** 1990. Cis elements and trans-acting factors involved in the tRNA dimeriza-tion of the human immunodeficiency virus HIV-1. J. Mol. Biol. **216:**689–699.
- 8. **Debyser, Z., A.-M. Vandamme, R. Pauwels, M. Baba, J. Desmyter, and E. De Clercq.** 1992. Kinetics of inhibition of endogenous human immunodeficiency virus type 1 reverse transcription by 2',3'-dideoxynucleoside 5'-triphosphate, tetrahydroimidazo-[4,5,1-jk][1,4]-benzodiazepin-(phenylthio)thymine derivatives. J. Biol. Chem. **267:**11769–11776.
- 9. **Faraj, A., L. A. Agrofoglio, J. K. Wakefield, S. Mcpherson, C. D. Morrow, G. Gosselin, C. Marthe, J.-L. Imbach, R. F. Schinazi, and J.-F. Sommadossi.** 1994. Inhibition of human immunodeficiency virus type 1 reverse transcriptase by the 5'-triphosphate  $\beta$  enantiomers of cytidine analogs. Antimicrob. Agents Chemother. **38:**2300–2305.
- 10. **Gao, Q., Z. Gu, M. A. Parniak, J. Cameron, N. Cammack, C. Boucher, and M. A. Wainberg.** 1993. The same mutation that encodes low-level human immunodeficiency virus type 1 resistance to  $2^{\prime},3^{\prime}$ -dideoxyinosine and  $2^{\prime},3^{\prime}$ dideoxycytidine confers high-level resistance to the  $(-)$  enantiomer of 2',3'dideoxy-3'-thiacytidine. Antimicrob. Agents Chemother. 37:1390-1392.
- 11. **Gilboa, E., S. Goff, A. Shields, F. Yoshimura, S. Mitra, and D. Baltimore.** 1979. In vitro synthesis of a 9 kbp terminally redundant DNA carrying the infectivity of Moloney murine leukemia virus. Cell **16:**863–874.
- 12. **Gu, Z., E. J. Arts, M. A. Parniak, and M. A. Wainberg.** 1995. Mutated K65R recombinant reverse transcriptase of human immunodeficiency virus type 1 shows diminished chain termination in the presence of  $2'3'$ -dideoxycytidine 59 triphosphate and other drugs. Proc. Natl. Acad. Sci. USA **92:**2760–2764.
- 13. **Gu, Z., R. S. Fletcher, E. J. Arts, M. A. Wainberg, and M. A. Parniak.** 1994. The K65R mutant reverse transcriptase of HIV-1 cross-resistant to  $2^{\prime},3^{\prime}$ dideoxycytidine, 2',3'-dideoxy-3'-thiacytidine and 2',3'-dideoxyinosine shows reduced sensitivity to specific dideoxynucleoside triphosphate inhibitors in vitro. J. Biol. Chem. **269:**28118–28122.
- 14. **Gu, Z., Q. Gao, X. Li, M. A. Parniak, and M. A. Wainberg.** 1992. Novel mutation in the human immunodeficiency virus type 1 reverse transcriptase gene that encodes cross-resistance to  $2^{\prime}, 3^{\prime}$ -dideoxyinosine and  $2^{\prime}, 3^{\prime}$ -dideoxycytidine. J. Virol. **66:**7128–7135.
- 15. **Hermann, T., T. Meier, M. Gotte, and H. Heumann.** 1994. The "helix clamp"

in HIV-1 reverse transcriptase: a new nucleic acid binding motif common in

- nucleic acid polymerases. Nucleic Acids Res. 22:4625–4633.<br>16. Kohlstaedt, L. A., J. Wang, J. M. Friedman, P. A. Rice, and T. A. Steitz. 1992.<br>Crystal structure at 3.5 Å resolution of HIV-1 reverse transcriptase complexed with an inhibitor. Science **256:**1783–1790.
- 17. **Lacey, S. F., J. E. Reardon, E. S. Furfine, T. A. Kunkel, K. Bebenek, K. A. Eckert, S. D. Kemp, and B. A. Larder.** 1992. Biochemical studies on the reverse transcriptase and RNase H activities from human immunodeficiency virus strains resistant to 3'-azido-3'-deoxythymidine. J. Biol. Chem. 267: 15789–15794.
- 18. **Larder, B. A., and S. D. Kemp.** 1989. Multiple mutations in HIV-1 reverse transcriptase confer high-level resistance to zidovudine (AZT). Science **246:** 1155–1158.
- 19. **Larder, B. A., S. D. Kemp, and P. R. Harrigan.** 1995. Potential mechanism for sustained antiretroviral efficacy of AZT-3TC combination therapy. Science **269:**696–699.
- 20. **Martin, J. L., J. E. Wilson, R. L. Haynes, and P. A. Furman.** 1993. Mecha-

nism of resistance of human immunodeficiency virus type 1 to 2',3'-dideoxyinosine. Proc. Natl. Acad. Sci. USA **90:**6135–6139.

- 21. **Schinazi, R. F., R. M. Lloyd, Jr., M.-H. Nguyen, D. L. Cannon, A. McMillan, N. Ilksoy, C. K. Chu, D. C. Liotta, H. Z. Bazmi, and J. W. Mellors.** 1993. Characterization of human immunodeficiency viruses resistant to oxathiolane-cytosine analogs. Antimicrob. Agents Chemother. **37:**875–881.
- 22. **Tisdale, M., S. D. Kemp, N. R. Parry, and B. A. Larder.** 1993. Rapid in vitro selection of human immunodeficiency virus type 1 resistant to 3'-thiacytidine inhibitors due to a mutation in the YMDD region of reverse transcriptase. Proc. Natl. Acad. Sci. USA **90:**5653–5656.
- 23. **Trono, D.** 1992. Partial reverse transcripts in virions from human immunodeficiency and murine leukemia viruses. J. Virol. **66:**4893–4900.
- 24. **Wakefield, J. K., S. A. Jablonski, and C. D. Morrow.** 1992. In vitro enzymatic activity of human immunodeficiency virus type 1 reverse transcriptase mutants in the highly conserved YMDD amino acid motif correlates with the infectious potential of the proviral genome. J. Virol. **66:**6806–6812.