Amino Acid Deletion at Codon 67 and Thr-to-Gly Change at Codon 69 of Human Immunodeficiency Virus Type 1 Reverse Transcriptase Confer Novel Drug Resistance Profiles

TOMOZUMI IMAMICHI,¹* MICHAEL A. MURPHY,¹ HIROMI IMAMICHI,¹ and H. CLIFFORD LANE²

*Laboratory of Molecular Retrovirology, Clinical Services Program, Science Applications International Corporation— Frederick, National Cancer Institute—Frederick, Frederick,*¹ *and Laboratory of Immunoregulation, National Institute of Allergy and Infectious Diseases, Bethesda,*² *Maryland*

Received 26 October 2000/Accepted 19 January 2001

The potential roles of an amino acid deletion at codon $67 (\Delta 67)$ and a Thr-to-Gly change at codon $69 (T69G)$ **in the reverse transcriptase of human immunodeficiency virus (HIV) type 1 in drug sensitivity and relative** replication fitness were studied. Our results suggest that the $\Delta 67$ and T69G changes can be categorized as **mutations associated with multidrug resistance. The combination of both mutations with an L74I change (**D**67**1**T69G/L74I) leads to a novel 3*****-azido-3*****-deoxythymidine resistance motif and compensates for impaired HIV replication.**

Current analyses of genotypic drug resistance patterns have revealed the presence of stable rearrangements, including amino acid insertions and deletions in the β 3- β 4 hairpin loop of the finger domain of the reverse transcriptase (RT) of human immunodeficiency virus type 1 (HIV-1) (3, 18a, 19). The emergence of these stable rearrangements is associated with failures of combination multiple antiretroviral therapy (3, 7, 11, 13, 16, 18, 18a, 19). A two-amino-acid insertion between codons 69 and 70 in the RT has been shown to confer resistance to multiple nucleoside RT inhibitors (NRTIs) (11, 13, 18). We have recently identified an amino acid deletion at codon 67 (Δ 67) and a Thr-to-Gly change at codon 69 (T69G) in the RT of HIV-1 from a patient who had failed combination therapy with $3'$ -azido-3'-deoxythymidine (AZT) and $2',3'$ dideoxyinosine (ddI) (7). The emergence of this same combination of mutations in the RT has also recently been reported under the selection pressure of 2',3'-didehydro-3'-deoxythymidine $(D4T)$ – $(-)$ β -L-2', $3'$ -dideoxy-3'-thiacytidine (3TC)–indinavir (14). Therefore, it appears that the emergence of the $\Delta 67$ deletion and the T69G change is not the result of any single treatment regimen but rather may be due to an unusual set of host and/or viral characteristics.

We have demonstrated that high-level resistance to AZT (up to 1,810-fold) was seen in the setting of the $\Delta 67$ and T69G changes in association with the AZT resistance mutations K70R, T215F, and K219Q and the non-NRTI (NNRTI) resistance mutations L74I and K103N (7). Further study has revealed that the emergence of the T69G mutation confers drug resistance at the expense of fitness (8). Subsequently, the development of the $\Delta 67$ deletion led to a virus with improved replication and high-level AZT resistance (8). The purpose of the present study was to determine the overall profiles of

resistance to licensed NRTIs and NNRTIs in viruses containing the $\Delta 67$ deletion and the T69G mutation.

A series of HIV-1 variants were created in a cloned proviral DNA pNL4.3 (1) backbone using a Quickchange site mutagenesis kit (Stratagene, La Jolla, Calif.) (8). Drug resistance assays were carried out using MT-2 cells (4, 5) following previously described methods (8). Sensitivities to each drug were reported as the concentrations of the drugs that inhibited p24 production by 50% (IC₅₀s) in tissue culture systems (8, 20). Significant differences between HIV-1 variants in drug sensitivity were calculated by using the unpaired *t* test of the StatView program (Abacus Concepts, Berkeley, Calif.).

Constructs containing either the $\Delta 67$ or the T69G change demonstrated increases in the IC_{50} s of 3'-dideoxycytidine (ddCi Sigma, St. Louis, Mo.) ($P < 0.01$), ddI (Sigma) ($P <$ 0.01), D4T ($P < 0.05$ for $\Delta 67$, $P < 0.01$ for T69G) (Sigma), 3TC ($P < 0.01$), and abacavir ($P < 0.01$) (Table 1). They were sensitive to AZT (Sigma), nevirapine, and delavirdine. Interestingly, the $\Delta 67$ or T69G change increased sensitivity to efavirenz ($P < 0.01$). A mutant containing both the $\Delta 67$ and T69G changes $(\Delta 67 + T69)$ was sensitive to AZT, ddI, nevirapine, delavirdine, and efavirenz. The $\Delta 67$ and T69G changes increased the ICs₅₀ of ddC (*P* < 0.05), 3TC (*P* < 0.01), D4T (*P* < 0.05), and abacavir $(P < 0.05)$ (Table 1). Ross et al. recently reported that the $\Delta 67$ mutation in the HXB2 backbone did not lead to high-level resistance to NRTIs (14). Differences in assay methodology or subtle difference in the HIV genotypic backbone may explain this difference between our results and theirs.

Since the $\Delta 67$ or T69G change increased sensitivity to efavirenz, further phenotypic studies were performed to assess the impact of these mutations on NNRTI-resistant variants of HIV-1. As described by other groups, the variant containing the K103N change showed resistance to nevirapine, delavirdine, and efavirenz (15). Addition of either the $\Delta 67$ or T69G mutation to the K103N mutant diminished resistance to efavirenz. In contrast, the presence of both the $\Delta 67$ and T69G

Corresponding author. Mailing address: Laboratory of Molecular Retrovirology, Clinical Services Program, SAIC—Frederick, NCI— Frederick, Building 550, Room 126, P.O. Box B, Frederick, MD 21702- 1201. Phone: (301) 846-5450. Fax: (301) 846-6762. E-mail: timamichi @nih.gov.

FIG. 1. Impact of the $\Delta 67+T69G/L74I$ motif in combination with known NNRTI resistance mutations on AZT sensitivity. Recombinant mutants were constructed by site-directed mutagenesis. The $\Delta 67 + T69G/L74I$ motif in the RT gene of NL4.3 was induced in each variant containing a single NNRTI resistance mutation. Recombinant viruses were assessed for AZT susceptibility with the drug resistance assay (7, 20). Open bars show the nanomolar IC₅₀s of AZT for mutants without the $\Delta 67 + T69G/L74I$ mutations, and closed bars show the IC₅₀s of AZT for mutants with the $\Delta 67+T69G/L74I$ mutations. The data shown are means \pm standard errors. Each construct was independently tested at least three times.

changes in the K103N mutant partially decreased resistance to all NNRTIs (Table 1).

We have previously reported that the combination of an L74I change, an NNRTI resistance mutation with the $\Delta 67$ deletion, and a T69G change $(\Delta 67 + T69G/L74I)$ in the RT resulted in AZT resistance. The IC_{50} of AZT for the Δ 67+T69G/L74I mutant was fivefold higher than that for the wild type (7). The further addition of a K103N change to the Δ 67+T69G/L74I motif led to a further increase (16-fold) in the IC_{50} of AZT (7).

To define the combined effects of the $\Delta 67 + T69 \text{G/L}74 \text{I}$ motif and other NNRTI resistance mutations on AZT resistance, a series of constructs containing the well-described NNRTI resistance mutations were created by site-directed mutagenesis (8). Addition of an A98G change to the $\Delta 67 + T69G/L74I$ motif also led to an increase in the IC_{50} (520 \pm 70 nM; $P < 0.01$) of AZT. No significant changes in AZT resistance were seen upon addition of V106A (92 \pm 20 nM), Y108A (100 \pm 20 nM), Y188C (164 \pm 48 nM), or G190S (205 \pm 53 nM; *P* < 0.1). The Y181C and L100I changes can lead to suppression of AZT resistance (2, 10). Addition of the Y181C or L100I change diminished the IC₅₀ of AZT for the $\Delta 67 + T69$ G/L74I motifcontaining mutant to 10 ± 1.7 or 12 ± 8 nM, respectively (Fig. 1). An M184V change in the RT confers resistance to 3TC and suppresses AZT resistance (15). Addition of an M184V mutation also downregulated AZT resistance in the $\Delta 67 + T69$ G/ L74I construct (24 \pm 4.5 nM).

Mutations M41L, D67N, K70R, T215F/Y, and K219Q/E are associated with well-described AZT resistance (15). Codons 41, 67, and 70 are present in the β 3- β 4 loop region of the finger domain of the RT and separated by approximately 25 Å from codons 215 and 219 in the β 11a- β 11b loop. Meanwhile, codons 98 and 103 are present in the β 5- β 6 loop of the NNRTI binding site and separated by approximately 40 \AA from the finger domain (17). Despite this distance, the combination of the $\Delta 67$ deletion and the T69G and L74I changes with A98G or K103N led to AZT resistance (Fig. 1). The crystal structure of a covalently trapped catalytic complex of HIV-1 RT has been reported (6). Binding of the template-primer and a deoxynucleoside triphosphate to the DNA-binding cleft between the finger and palm domains of the RT has been shown to induce the outer part of the finger domain, the fingertip, to bend inward toward the palm domain (6). Recently, Winters et al. have reported that the $\Delta 67$ deletion in the $\beta 3$ - $\beta 4$ loop and the T69G mutation in HIV-1 are associated with nucleoside analogs and have proposed a model to explain their findings (18a). In this model, these mutations lead to changes in the three-dimensional structure of the β 3- β 4 loop and alpha helices C and E in both RT-DNA open (17) and RT-DNA-dTTP closed structures (6). These changes lead to the loss of hydrogen bonds between the RT and dTTP (18a). Therefore, the changes in these regions may lead to an inefficient RT and then allow a novel interaction between the RT and the templateprimer-deoxynucleoside triphosphate, subsequently leading to a novel drug resistance.

In the absence of L74I, AZT resistance was not seen in the Δ 67+T69G/K103N construct (data not shown). Therefore, the L74I mutation or the combination of $\Delta 67 + T69G/L74I$ changes in the finger domain might facilitate long-distance "communication" between the finger and palm domains through the template-primer, as first reported by Kleim et al. (9).

We have previously reported that the T69G mutation led to ddI resistance in an AZT-resistant backbone and was associated with impaired HIV replication (8) . Emergence of the $\Delta 67$ deletion and the L74I change in the RT compensated for this decrease in HIV replication, even though the $\Delta 67$ deletion alone led to impaired HIV replication (8). To characterize the effect of the $\Delta 67 + T69$ G/L74I motif on the replication capa-

 (B)

FIG. 2. Replication properties of recombinant HIV-1 variants. Recombinant HIV-1 mutants were constructed by site-directed mutagenesis. MT-2 cells were infected with 1,250 50% tissue culture-infective loses/3 \times 10⁶ cells and cultured for 7 days. To determine virus growth properties, levels of p24 antigen in day 7 culture supernatant were measured with a p24 antigen capture kit (8). The $\Delta 67+T69G/L74I$ motif in the RT gene of NL4.3 was induced in each variant containing a single NNRTI resistance mutation. Viruses were cultured in the absence (A) or presence (B) of 1 μ M AZT. Closed bars show results for variants without the Δ 67+T69G/L74I mutations, and hatched bars show results for variants with the Δ 67+T69G/L74I mutations. Results are expressed as the mean percentage of growth \pm the standard error of the mean compared to WT growth in the absence of AZT from three independent experiments. In this experiment, the WT p24 concentration was 713 \pm 53 ng/ml.

bility of HIV containing NNRTI resistance mutations, relative replication fitness assays were performed in the presence or absence of $1 \mu M$ AZT. Infected MT-2 cells were cultured at 0.1×10^6 /ml for 7 days, and p24 levels in day 7 supernatants were measured by a p24 antigen capture kit (8) (Fig. 2). In the absence of AZT (Fig. 2A), a K103N, V108I, Y181C, Y188C, or G190S change in the RT did not lead to a significant difference in growth compared with the wild-type (WT) virus $(P > 0.05)$. In contrast, a A98G, L100I, or V106A change, decreased virus replication to 28 \pm 7.0, 66 \pm 6.5, or 40 \pm 4.8% of that of the WT, respectively. The impaired HIV replication induced by the A98G or V106A change was compensated for by the addition of the $\Delta 67 + T69G/L74I$ motif to the mutant. Meanwhile, addition of the $\Delta 67 + T69$ G/L74I motif to the L100I, Y181C, or Y188C change significantly decreased HIV replication to 37 \pm 4.4, 46 \pm 21, or 10 \pm 2.0% of that of the WT, respectively (*P* < 0.01). In the presence of 1 μ M AZT (Fig. 2B), all variants without the $\Delta 67 + T69G/L74I$ motif were sensitive to AZT. Addition of the motif to the WT or A98G or K103N mutant resulted in partial restoration of replication to 28 \pm 11, 38 \pm 11, and 65 \pm 6.3%, respectively. Archer et al. demonstrated that NNRTI resistance mutations could alter the rate of RNase H cleavage, which is correlated with HIV replication (1a). Since addition of the $\Delta 67 + T69 \text{G/L}74$ motif to variants containing mutations associated with resistance to NNRTI affected their replication capability in this study (Fig. 2A), the motif might also influence RNase H cleavage.

To define the novel AZT resistance mechanism, a virionassociated RT inhibition assay was also attempted using serial twofold dilutions of AZT-TTP (DuPont NEN, Boston, Mass.) and a commercially available RT assay kit (Roche Molecular Biochemical, Indianapolis, Ind.). The virion-associated RT activity of the $\Delta 67 + T69 \text{G/L}74$ I motif did not show resistance to AZT-TTP (data not shown). Further biochemical study of this Δ 67+T69G/L74I motif is necessary to understand the precise mechanism(s) of the novel AZT resistance motif, as well as the compensated replication fitness. Recently, Meyer et al. demonstrated a novel mechanism of AZT resistance due to an increase in primer-unblocking activity (12). It is hoped that such knowledge will lead to the development of better therapies and salvage therapies.

We thank Hiroaki Mitsuya at the NCI for kindly providing 3TC and abacavir. Nevirapine, delavirdine, and efavirenz were provided by Boehringer Ingelheim (Ridgefield, Conn.), Pharmacia & Upjohn (Kalamazoo, Mich.), and DuPont Pharmaceutical Company (Wilmington, Del.), respectively. MT-2 cells and pNL4.3 were obtained through the AIDS Research and Reference Reagent Program, Division of AIDS, NIAID, NIH.

This study was supported by the National Institute of Allergy and Infectious Diseases under contract N01-CO-56000 with SAIC—Frederick.

REFERENCES

- 1. **Adachi, A., H. E. Gendelman, S. Koenig, T. Folks, R. Willey, A. Rabson, and M. A. Martin.** 1986. Production of acquired immunodeficiency syndromeassociated retrovirus in human and nonhuman cells transfected with an infectious molecular clone. J. Virol. **59:**284–291.
- 1a.**Archer, R. H., C. Dykes, P. Gerondelis, A. Lloyd, P. Fay, R. C. Reichman, R. A. Bambara, and L. M. Demeter.** 2000. Mutants of human immunodeficiency virus type 1 (HIV-1) reverse transcriptase resistant to nonnucleoside reverse transcriptase inhibitors demonstrate altered rates of RNase H cleavage that correlate with HIV-1 replication fitness in cell culture. J. Virol. **74:**8390–8401.

2. **Byrnes, V. W., E. A. Emini, W. A. Schleif, J. H. Condra, C. L. Schneider,**

 \wedge 0.05. **W. J. Long, J. A. Wolfgang, D. J. Graham, L. Gotlib, A. J. Schlabach, B. S. Wolanski, O. M. Blahy, J. C. Quintero, A. Rhodes, E. Roth, D. L. Titus, and V. V. Sardana.** 1994. Susceptibilities of human immunodeficiency virus type 1 enzyme and viral variants expressing multiple resistance-engendering amino acid substitutions to reverse transcriptase inhibitors. Antimicrob. Agents Chemother. **38:**1404–1407.

- 3. **De Jong, J. J., J. Goudsmit, V. V. Lukashov, M. E. Hillebrand, E. Baan, R. Huismans, S. A. Danner, J. H. ten Veen, F. de Wolf, and S. Jurriaans.** 1999 Insertion of two amino acids combined with changes in reverse transcriptase containing tyrosine-215 of HIV-1 resistant to multiple nucleoside analogs. AIDS **13:** 75–80.
- 4. **Haertle, T., C. J. Carrera, D. B. Wasson, L. C. Sowers, D. D. Richman, and D. A. Carson.** 1988. Metabolism and anti-human immunodeficiency virus-1 activity of 2-halo-29,39-dideoxyadenosine derivatives. J. Biol. Chem. **263:** 5870–5875.
- 5. **Harada, S., Y. Koyanagi, and N. Yamamoto.** 1985. Infection of HTLV-III/ LAV in HTLV-carrying cells MT-2 and MT-4 and application in a plaque assay. Science **229:**563–566.
- 6. **Huang, H., R. Chopra, G. L. Verdine, and S. C. Harrison.** 1998. Structure of a covalently trapped catalytic complex of HIV-1 reverse transcriptase: implication for drug resistance. Science **282:**1669–1675.
- 7. Imamichi, T., T. Sinha, H. Imamichi, Y. M. Zhang, J. A. Metcalf, J. Falloon, and H. C. Lane. 2000. High-level resistance to 3'-azido-3'-deoxythimidine due to a deletion in the reverse transcriptase gene of human immunodeficiency virus type 1. J. Virol. **74:**1023–1028.
- 8. **Imamichi, T., S. C. Berg, H. Imamichi, J. C. Lopez, J. A. Metcalf, J. Falloon,** and H. C. Lane. 2000. Relative replication fitness of a high-level 3'-azido-3'-deoxythimidine-resistant variant of human Immunodeficiency virus type 1 possessing an amino acid deletion at codon 67 and a novel substitution $(Thr\rightarrow Gly)$ at codon 69. J. Virol. **74:**10958–10964.
- 9. **Kleim, J. P., M. Rosner, I. Winkler, A. Paessens, R. Kirsch, V. Hsiou, E. Arnold, and G. Riess.** 1996. Selective pressure of a quinoxaline non-nucleoside inhibitor of human immunodeficiency virus type 1 (HIV-1) reverse transcriptase (RT) on HIV-1 replication results in the emergence of nucleoside RT-inhibitor-specific (RT Leu-743Val or Ile and Val-753Leu or Ile) HIV-1 mutants. Proc. Natl. Acad. Sci. USA **93:**34–38.
- 10. Larder, B. A. 1992. 3'-Azido-3'-deoxythymidine resistance suppressed by a mutation conferring human immunodeficiency virus type 1 resistance to nonnucleoside reverse transcriptase inhibitors. Antimicrob. Agents Chemother. **36:**2664–2669.
- 11. **Larder, B. A., S. Bloor, S. D. Kemp, K. Hertogs, R. L. Desmet, V. Miller, M. Sturmer, S. Staszewski, J. Ren, D. K. Stammers, D. I. Stuart, and R. Pauwels.** 1999. A family of insertion mutations between codons 67 and 70 of human immunodeficiency virus type 1 reverse transcriptase confer multinucleoside analog resistance. Antimicrob. Agents Chemother. **43:**1961–1967.
- 12. **Meyer, P. R., S. E. Matsuura, A. M. Mian, A. G. So, and W. A. Scott.** 1999. A mechanism of AZT resistance: an increase in nucleotide-dependent primer unblocking by mutant HIV-1 reverse transcriptase. Mol. Cell **4:**35–43.
- 13. **Ross, L., M. Johnson, N. Graham, M. Shaefer, and M. St. Clair.** 1999. The reverse transcriptase codon 69 insertion is observed in nucleoside reverse transcriptase inhibitor-experienced HIV-1-infected individuals, including those without prior or concurrent zidovudine therapy. J. Hum. Virol. **2:**290– 295.
- 14. **Ross, L., M. Johnson, R. G. Ferris, S. A. Short, L. R. Boone, T. E. Melby, R. Lanier, M. Shaefer, and M. St. Clair.** 2000. Deletions in the beta3-beta4 hairpin loop of HIV-1 reverse transcriptase are observed in HIV-1 isolated from subjects during long-term antiretroviral therapy. J. Hum. Virol. **3:**144– 149.
- 15. **Schinazi, R. F., B. A. Larder, and J. W. Mellors.** 2000. Mutations in retroviral genes associated with drug resistance: 1999–2000 update. Int. Antiviral News **8:**65–91.
- 16. **Tamalet, C., J. Izopet, N. Koch, J. Fantini, and N. Yahi.** 1998. Stable rearrangements of the β 3- β 4 hairpin loop of HIV-1 reverse transcriptase in plasma viruses from patients receiving combination therapy. AIDS **12:**F161– F166.
- 17. **Tantillo, C., J. Ding, A. Jacobo-Molina, R. G. Nanni, P. L. Boyer, S. H. Hughes, R. Pauwels, K. Andries, P. A. J. Janssen, and E. Arnold.** 1994. Locations of anti-AIDS drug binding sites and resistance mutations in the three dimensional structure of HIV-1 RT. J. Mol. Biol. **243:**369–387.
- 18. **Winters, M. A., K. L. Coolley, Y. A. Girard, D. J. Levee, H. Hamdan, R. W. Shafer, D. A. Katzenstein, and T. C. Merigan.** 1998. A 6-basepair insert in the reverse transcriptase gene of human immunodeficiency virus type 1 confers resistance to multiple nucleoside inhibitors. J. Clin. Investig. **102:** 1769–1775.
- 18a.**Winters, M. A., K. L. Coolley, P. Cheng, Y. A. Girard, H. Hamdan, L. C. Kovari, and T. C. Merigan.** 2000. Genotypic, phenotypic, and modeling studies of a deletion in the beta3-beta4 region of the human immunodeficiency virus type 1 reverse transcriptase gene that is associated with resistance to nucleoside reverse transcriptase inhibitors. J. Virol. **74:**10707–10711.
- 19. **Yahi, N., C. Tamalet, C. Tourres, N. Tivoli, F. Ariasi, F. Volt, J. A. Gastaut, H. Gallais, H. J. Moreau, and J. Fantini.** 1999. Mutation pattern of the reverse transcriptase and protease genes in human immunodeficiency virus type 1-infected patients undergoing combination therapy: survey of 787 sequences. J. Clin. Microbiol. **37:**4099–4106.
- 20. **Zhang, Y.-M., H. Imamichi, T. Imamichi, H. C. Lane, J. Falloon, M. B. Vasudevachari, and N. P. Salzman.** 1997. Drug resistance during indinavir therapy is caused by mutations in the protease gene and in its Gag substrate cleavage sites. J. Virol. **71:**6662–6670.