

Molecular Interactions of Human Immunodeficiency Virus Type 1 with Primary Human Oral Keratinocytes

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Infection of the oral mucosa of human immunodeficiency virus type 1 (HIV-1)-infected individuals remains an under-evaluated and somewhat enigmatic process. Nonetheless, it is of profound importance in the ongoing AIDS pandemic, based on its potential as a site of person-to-person transmission of the virus as well as a location of HIV-1 pathogenesis and potential reservoir of disease in the setting of virally suppressive highly active antiretroviral therapy. We utilized molecular and virological techniques to analyze HIV-1 infection of primary human mucosal cells and also evaluated the proapoptotic potential of selected HIV-1 proteins in primary isolated human oral keratinocytes. Primary isolated human oral keratinocytes were plated on 0.4 μ M polyethyleneterephthalate cell culture inserts to form an in vitro oral mucosal layer. The strength of this layer in forming a barrier was determined by measuring *trans*-epithelial electrical current passage across the monolayer. The oral keratinocyte monolayers had *trans*-epithelial electrical resistance of approximately 176 to 208 Ω . For viral infectivity assays, the macrophage-tropic (R5) HIV-1 strains, YU-2 and ADA, and T-cell-line-tropic (X4), NL4-3 virions, incubated with or without deoxynucleoside triphosphates (dNTPs) and/or the polyamines spermine and spermidine, were used to infect oral keratinocytes. Of importance, polyamines and dNTPs have been shown to enhance natural endogenous reverse transcription (NERT), a step essential for early lentiviral infection, and are abundantly present in human semen. The infectivities of HIV-1 strains YU-2, ADA, and NL4-3 for these primary keratinocytes were dramatically increased by the addition of physiological concentrations of dNTPs, spermine, and spermidine. Binding and viral internalization assay studies showed no differences in these oral mucosal cells, with or without NERT-altering agents. It was also observed that the recombinant, cell-free HIV-1 proteins Nef, Tat, and gp120 (R5) induced apoptosis in primary oral keratinocytes compared with the results seen with nontreated cells or cells treated with glutathione *S*-transferase protein as a control under similar conditions. Microarray analyses suggested that HIV-1 gp120 and Tat induce apoptosis in primary human oral keratinocytes via the Fas/FasL apoptotic pathway, whereas induction of apoptosis by Nef occurs through both Fas/FasL and mitochondrial apoptotic pathways. Thus, these findings suggest molecular mechanisms by which semen in particular, as well as other bodily fluids such as cervicovaginal secretions, could increase oral transmission of HIV-1 via increasing infectivity in confluent and low-replicating oral keratinocytes. As well, the induction of apoptosis in human oral keratinocytes with relevant HIV-1-specific proteins suggests another potential complementary mechanism by which the oral mucosa barrier may be disrupted during HIV-1 infection in vivo.

Infection of the oral mucosa by human immunodeficiency virus type I (HIV-1) in vivo is a critical area of HIV-1 pathogenesis which has been investigated previously, but an extensive number of important questions remain unanswered (5, 7, 20, 31, 57). This field of research is key, as the oral mucosa is increasingly found to be associated with horizontal (and vertical) transmission of this human lentivirus, as well as being potentially involved with a variety of pathogenic processes in the oral cavities of HIV-1-infected individuals (10, 26, 84, 106).

HIV-1 had been thought previously to rarely be transmitted between individuals via oral-genital and oral-oral contact of mucosa or through infected saliva (56, 60, 109). Nevertheless, several recent studies have suggested that a receptive oral sexual contact may lead to significant HIV-1 transmission in certain cohorts, especially in male homosexuals and crack cocaine and intravenous drug users (34, 97, 99, 102). A number of studies have demonstrated that HIV-1 can be detected in the saliva of infected individuals by culture, reverse transcriptase PCR, and DNA-PCR. Although the levels are usually lower than in peripheral blood or semen and cervical/vaginal secretions of infected individuals, certain patients have been shown to be “hyper-secretors” with respect to saliva (62, 92, 107). A recent study has demonstrated by heteroduplex mapping that the viral quasispecies in peripheral blood is quite similar to viral strains in the saliva of HIV-1-infected individuals (40). As well, both in vitro and in vivo studies have suggested that epithelial cells may be infected with HIV-1 (23, 61,

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71). Our laboratories have previously shown, utilizing in situ PCR, that HIV-1 could be detected in a significant majority of infected individuals within their oral mucosal epithelial cells (92). As well, detection of HIV-1 in oral mucosal cells was demonstrated by electron microscopy (91). Recent studies have demonstrated that oral keratinocytes can be infected by HIV-1 using galactosylceramide and CXCR4 as receptors (61), and ethanol may increase infectivity (23). As well, other studies have suggested that HIV-1 infection, both cell free and cell associated, of salivary gland cell lines via CXCR4 and CCR5 with a CD4-independent mechanism can be demonstrated in vitro (70). In addition, secretory immunoglobulin A from parotid glands may inhibit CCR5- but not CXCR4-tropic viruses. Data also suggest that oral shedding of HIV-1 RNA correlates with antiretroviral therapy, CD4⁺ T-cell counts, and tonsillectomy status (134).

It has been well described for the past decade or more that a variety of salivary inhibitors decrease HIV-1 infectivity in the oral cavity (108, 109). Nonetheless, ongoing studies have suggested that oral transmission of the virus remains quite common in a number of important settings in vivo (6, 7, 51). These enigmatic processes require significant evaluations to dissect, in a prioritized manner, the full virus:host cell interactions which occur in vivo and lead to oral HIV-1 transmission and pathogenesis. As such, in this study, we have utilized selected virological and molecular techniques to analyze HIV-1 infection and interactions with primary human oral keratinocytes. As well, we evaluated the potential for selected recombinant HIV-1 proteins to induce apoptosis in these primary human cells.

MATERIALS AND METHODS

Isolation of primary human oral mucosal cells. Primary oral mucosal cells were obtained from the gingiva of HIV-1-seronegative dental patients and processed as described, with some modifications (71). Briefly, the epithelial layer from the oral tissue was digested for 60 min at 37°C with collagenase type II (Millipore Corp., New Bedford, MA) (1.0 mg/ml) and dispase grade II (Boehringer Mannheim, Indianapolis, IN) (2.4 mg/ml) to separate the epithelium from the underlying submucosa. Epithelial sheets were then dispersed into individual cells by using trypsin. The cells were washed twice with Dulbecco's phosphate-buffered saline (PBS) containing amphotericin B (fungizone) plus penicillin/streptomycin and resuspended in keratinocyte growth medium (Clonetics Corp., San Diego, CA). Aliquots of 10⁴ cells were plated in fibronectin-coated six-well plastic tissue culture dishes (BD Biosciences Discovery Labware). Purity of keratinocytes was assessed by immunocytochemical analyses with the epithelial cell markers keratin and CD99. Cell isolates showing ~98% purity were used in subsequent studies. All studies were approved by our institutional review board.

Measurements of TEER. To measure the *trans*-epithelial electrical resistance (TEER) of confluent oral mucosal cells, the cells were cultured on 0.40 µM polyethyleneterephthalate track-etched membranes (BD Biosciences, Bedford, MA) to form an in vitro oral mucosa barrier layer. The integrity was determined by measuring *trans*-epithelial electrical resistance across the monolayer with the aid of a Millicell-ERS system (Millipore Corp., Bedford, MA), as described previously (72).

HIV-1 infection of primary oral mucosal cells. Viral infectivity assays were performed to determine the effects of natural endogenous reverse transcription (NERT) on HIV-1 infection of primary human oral mucosal cells. Aliquots (100 ng of p24 antigen equivalents) of the CXCR4-tropic HIV-1 strain, NL4-3, and the CCR5-tropic HIV-1 strains, YU-2 and ADA, were treated with either medium or NERT-stimulating agents, the polyamines spermine (3.0 mM) and spermidine (0.1 mM), with or without deoxynucleoside triphosphates (dNTPs) (100 nM), which are the physiological concentrations of these molecules in human semen (129). After incubation for 20 h at 37°C, aliquots were used to infect primary oral mucosal cells grown to confluence in fibronectin-coated, six-well tissue culture plates. After incubation for 4 h, the cells were washed 3 times with medium and viral growth was monitored for 1 week. Aliquots of the

supernatants were collected at specified time points for quantitation of the HIV-1 p24 antigen levels by enzyme-linked immunosorbent assays (ELISA) (Dupont).

Virion binding, uptake, and internalization studies. Aliquots of the CXCR4-tropic HIV-1 strain, NL4-3, and the CCR5-tropic HIV-1 strain, YU-2 (100 ng of HIV-1 p24 antigen), were incubated with NERT-stimulating agents, the polyamines spermine (3 mM) and spermidine (0.1 mM), with or without 100 nM dNTPs at 37°C for 4 h, according to a previously described methodology (29, 129-131). The ensuing cocktail was used to infect primary human oral mucosal cells. As described previously (29), controls included incubated but untreated virions. The unbound virions are then washed off with phosphate-buffered saline (PBS) (twice), and subsequently the uninternalized virions were removed by incubation with 0.25% trypsin (Sigma) for 5 min at 37°C. Three further washings, including one wash with a serum-containing medium, were then performed. The levels of HIV-1 p24 antigen in the cell lysates were quantitated by ELISA.

HIV-1 protein induction of programmed cell death in primary oral keratinocytes. To investigate the potential ability of selected HIV-1 proteins to induce apoptotic cell death in primary oral keratinocytes, three specific HIV-1 proteins, gp120, Tat, and Nef, which we and other groups have shown to induce apoptosis in various human cells (1, 82, 83), were evaluated. Nef and gp120 were expressed in our laboratory in baculovirus and vaccinia virus systems, respectively, whereas HIV-1 Tat was obtained from the National Institutes of Health AIDS Reagent and Repository Program. Low-passage, primary human oral keratinocytes were seeded onto fibronectin-coated, four-well chamber slides (BD Biosciences Discovery Labware). The cells were allowed to attach for 48 h before being exposed to 10 ng/ml of recombinant HIV-1 Nef, gp120 BaL (R5), Tat, and glutathione *S*-transferase (GST) (recombinant control protein). Selection of this particular initial concentration was based on our previous studies of primary human brain microvascular endothelial cells (1). Following treatment with viral proteins, cells were incubated for 48 h at 37°C and 5% CO₂ in a humidified environment. As additional negative controls, cells that had not been treated with any protein were used. Terminal deoxynucleotidyltransferase-mediated dUTP-biotin nick end labeling (TUNEL) assays were performed on treated and untreated cells, using an in situ cell death detection system, TMR Red (Roche Diagnostics, Mannheim, Germany), following the manufacturer's instructions. Briefly, cells were washed twice with 1× PBS and fixed for 10 min with acetone at room temperature. The fixed cells were washed three times with 1× PBS, and 50 µl of TUNEL reaction mixture was added to each well. Cells treated with the TUNEL mixture were incubated at 37°C in a humidified environment for 1 h. On completion of incubation, the treated cells were washed four times with PBS and evaluated by fluorescence microscopy. As positive controls, primary oral keratinocytes were treated with DNase (Life Technologies, Gaithersburg, MD) at a concentration of 1.0 µg/ml and incubated for 10 min at room temperature to induce DNA strand breaks. Untreated cells were also used as negative apoptosis controls, in addition to the GST-treated cells.

Semiquantitative apoptosis determinations. For semiquantitative measurements of apoptosis, images generated with a charge-coupled device array camera (RT Color; Diagnostic Instruments, Inc., Sterling Heights, MI) were subjected to fluorescence brightness value determinations on a monochromatic scale in red, blue, and green (RGB) values (0 to 255 ASCII numbers). Apoptosis was defined based on a red monochromatic scale in the range of 0 to 255, as described previously (1). Blank values were subtracted from the average of seven random values from different cells. Each image was further analyzed on a visual scale ranging from 1 to 10. This number was multiplied with ASCII numbers to generate final semiquantitative values for each image.

Human apoptosis gene microarrays. The CLONTECH Atlas cDNA expression human apoptosis array was used in this study. Total cellular RNA was extracted from third-passage human oral keratinocytes with an ULTRASPEC RNA isolation kit (Biotex Laboratories, Inc., Houston, TX), following the manufacturer's instructions. After RNA extraction, cDNA probe mixtures were synthesized by reverse transcribing respective RNA using cDNA synthesis primer mix provided by the manufacturer and [α -³²P]dATP (Perkin Elmer Sciences, Inc., Boston, MA). Each radioactively labeled probe mix was then hybridized to separate Atlas arrays overnight at 68°C. After high-stringency washes, as suggested by the manufacturer, the hybridization patterns were analyzed by autoradiography and quantified with a phosphorimager (Molecular Dynamics). The relative expression levels of a given cDNA from RNA obtained from oral keratinocytes treated with recombinant HIV-1 Nef, Tat, and gp120 protein were assessed by comparing the signal obtained with a probe of RNA from treated keratinocytes to that obtained with a probe of control RNA.

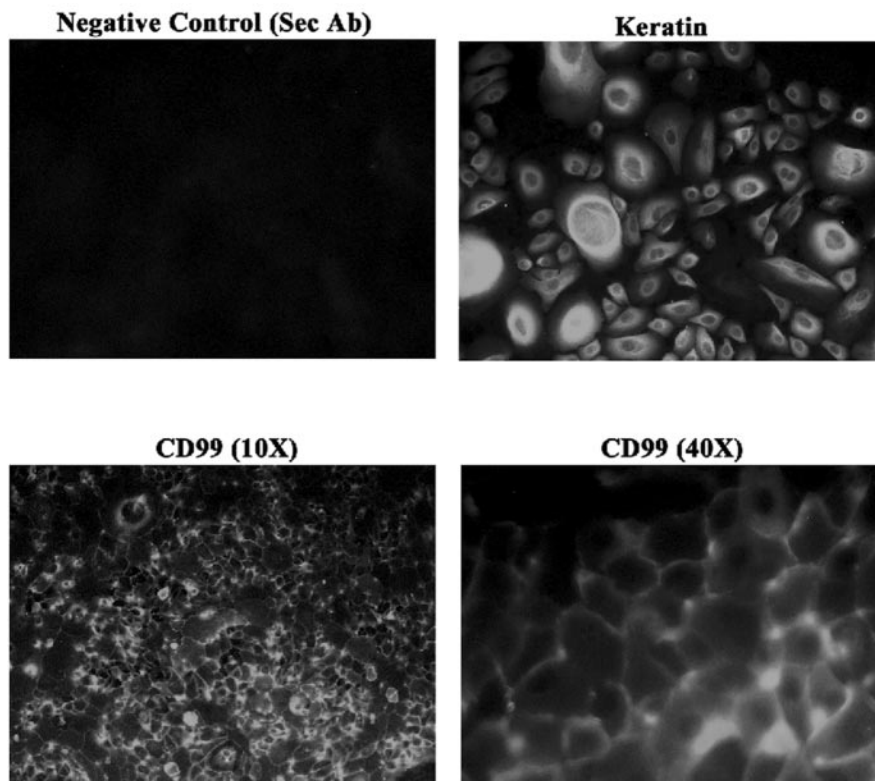


FIG. 1. Illustration of immunofluorescence staining for keratin, a specific characterization marker for keratinocytes, and CD99. Aliquots of 10⁴ cells were plated in fibronectin-coated six-well plastic tissue culture dishes (BD Biosciences Discovery Labware) and incubated for 72 h. Purity of keratinocytes was then assessed by immunocytochemical analyses with the epithelial cell markers keratin and CD99. Sec Ab, secondary antibody only.

RESULTS

Isolation and characterization of primary human oral keratinocytes. To analyze the ability of HIV-1 to infect primary oral mucosa cells, we first isolated and cultured pure keratin-positive, human oral mucosal epithelial cells. The purity of the cells was determined by immunocytochemical staining with monoclonal anti-pan cyokeratin antibodies from Sigma Aldrich Corp. (St. Louis, MO). The results are illustrated in Fig. 1. Keratin staining was robustly demonstrated using the CY2 chromophore. In these analyses, using immunofluorescence microscopy, few if any cells were shown to be fibroblastic by morphology or negative staining. In addition, we have begun to characterize the intercellular proteins that may be important in further studies of HIV-1-infected cell migration across an oral mucosal barrier. As demonstrated by initial studies illustrated in Fig. 1, CD99 was shown to have reproducible and significant expression in these primary oral mucosal epithelial explants. This is an extremely important molecule that has been demonstrated recently to play a major role in the migration of monocytes across endothelial junctions (101). As such, this may be a key molecule in understanding transmigration of HIV-1-infected monocytes and potentially CD4⁺ T lymphocytes across an oral mucosal barrier.

In addition, we sought to ascertain whether or not cultured human oral keratinocytes were able to generate an oral mucosal barrier system in vitro, analogous to our studies of in

vitro blood-brain barrier models (72). We utilized near-100%-pure primary oral mucosal cells cultured on 0.4 μM polyethyleneterephthalate membranes. These cultures, in multiple separate and independent samples, demonstrated a transepithelial electrical resistance (TEER) of approximately 176 to 208 Ω (Table 1). This is somewhat lower than that obtained for human brain microvascular endothelial cells but is still significantly elevated compared to baseline (e.g., brain microvascular endothelial cells frequently lead to approximately 250 Ω in models of this type in our hands). Whether these TEER values obtained for oral mucosal keratinocytes barriers reveal properties similar to other epithelial barriers requires further elucidation. Of note, we were unable to detect the expression of zonula occludens proteins, mainly involved in the creation of

TABLE 1. TEER measurements of human oral keratinocytes

Sample	TEER (ohms)
Control	131, 130
1.....	177
2.....	181
3.....	187
4.....	176
5.....	178
6.....	208
7.....	181
8.....	186

barrier properties of certain epithelial layers (not illustrated). These initial studies have suggested, however, not only that human oral keratinocytes can be grown in an in vitro explant system but also that they do have barrier functions, as described previously (25, 94).

Virological and microenvironmental factors which alter HIV-1 infection of oral keratinocytes. To evaluate critical virological and environmental parameters which alter HIV-1 infection of the oral keratinocytes, we first infected the cells in the presence of the NERT-stimulating agents: deoxynucleoside triphosphates (dNTPs) and polyamines. Of note, NERT represents stimulated reverse transcription within intact virions, which augments viral infectivity in many human cell types (29, 129). Following infection of oral keratinocytes with HIV-1 NL4-3, YU-2, and ADA in the presence or absence of these agents, we assayed HIV-1 *gag* p24 production in cell culture supernatants by ELISA. As shown in Fig. 2, in all three strains of HIV-1, the CXCR4-tropic NL4-3 virus as well as the CCR5-tropic viruses YU2 and ADA, viral infectivity of these primary oral explant mucosal cells could be dramatically increased with the use of agents which stimulate NERT, including polyamines and dNTPs, at physiological concentrations demonstrated in human semen (129). Although the CXCR4-tropic NL4-3 virus as well as the CCR5-tropic YU2 and ADA viruses all had their infectivities increased by these NERT-stimulating agents, it is interesting that YU-2 demonstrated a more dramatic effect from NERT stimulation. Control viral binding and internalization studies showed no differences in these oral cells with or without NERT-altering agents, demonstrating the specificity of these molecular effects (data not illustrated).

Induction of apoptosis in human primary keratinocytes by HIV-1 proteins. HIV-1 infection of cells can release specific viral products (123), which may induce apoptosis in bystander cells. It has been shown that HIV-1 infection can also lead to the production and release of proinflammatory cytokines (90), which are toxic to cells. Therefore, in addition to determining the effects of NERT on HIV-1 replication in primary human oral keratinocytes, we wished to determine whether HIV-1 viral proteins released during infection induce apoptosis in these cells. The ability of HIV-1 proteins to induce apoptotic cell death in primary keratinocytes, by exposing the cells to three selected HIV-1 specific proteins, gp120, Tat, and Nef, was investigated.

Primary oral keratinocytes cells were exposed to recombinant viral proteins Nef, Tat, gp120 BaL, and GST at a concentration of 10 ng/ml. At 48 h postexposure, apoptosis was evaluated utilizing the TUNEL assay; the results are depicted in Fig. 3A. It was observed that the use of nontreated cells and a control protein, GST, led to little or no programmed cell death in the primary oral keratinocytes. Individually each of the viral proteins showed apoptosis in oral keratinocytes, but with quite different intensities compared with control results. Strict quantitation is difficult in these assay systems, but in evaluations of multiple independent assays, visual examination of the photomicrographs suggested that the effect of Tat was the most pronounced, followed by gp120 BaL and Nef. However, comparison of all treatments using double-tailed Student's *t* tests indicated that the differences between Tat and gp120 did not rise to significance ($P > 0.08$). Similarly, there was no significant difference between gp120 and Nef ($P > 0.8$), but there was

a significant difference between Tat and Nef ($P < 0.05$). Of importance, though, there were various degrees of quite significant differences between all three HIV-1 proteins and the GST-treated, control oral keratinocytes, with all *P* values less than 0.001.

Human apoptosis gene expression. The mechanisms involved with induction of apoptosis in primary human oral keratinocytes are not known. Therefore, to elucidate the molecular mechanisms involved in HIV-1 protein-induced apoptosis, we utilized targeted gene microarrays to investigate the up-regulation of apoptosis-related genes in primary human oral keratinocytes exposed to HIV-1 Nef, gp120, and Tat proteins. Cells exposed to GST protein were used as controls.

Table 2, Table 3, and Table 4 depict apoptosis-related genes up-regulated in response to gp120, Nef, and Tat, respectively. The genes up-regulated when cells were exposed to gp120 included tumor necrosis factor (TNF) receptor super family member 1, and caspases 3, 6, and 8, as well as tumor proteins p53, Bcl-2, NF- κ B, cyclin B1, cyclin-dependent kinase 1 (cdk1), and other apoptosis-related genes. As Table 3 indicates, Nef-treated oral keratinocytes showed up-regulation in the expression of p53, DAXX, mitogen-activated protein kinase (MAPK) 7, and caspases 3, 6, 8, 9, and 10, as well as TNF receptor super family members 1 and 2 and Bcl-xL plus other apoptosis-related genes, compared with the GST-treated control. The results from the Tat-treated primary human oral keratinocytes are depicted in Table 4. As indicated in this table, modest up-regulation in the expression of MAPK 7, cyclin B, caspases 9 and 10, TNF receptor super family 12, and Ras homolog gene family member A were observed in primary oral keratinocytes exposed to Tat. PICTAIRE protein kinase 3 and E2F transcription factor 5 were modestly down-modulated. As can be observed in Tables 2, 3, and 4, it is clear that most of the up-regulated genes have been implicated in the Fas/FasL apoptotic pathways. Therefore, increases in their mRNA expression levels suggest possible involvement of the Fas/FasL apoptotic pathway in HIV-1 gp120-, Nef-, and Tat-induced apoptosis in oral keratinocytes. We also observed up-regulation in mRNA expression levels of genes involved in the mitochondrial apoptotic pathway, notably caspase 9, in cells exposed to HIV-1 Nef and Tat. It could therefore be strongly suggested that HIV-1 Nef and Tat may utilize more than one pathway in inducing apoptosis in primary human oral keratinocytes.

DISCUSSION

These studies evaluated virological and microenvironmental factors which potentially may increase HIV-1 infection of oral mucosal epithelial cells. First, natural endogenous reverse transcription (NERT) (i.e., intravirion reverse transcription), which has been described previously (29, 129) as a new phase of the viral life cycle, was analyzed for its effects on augmenting infection of human oral keratinocytes in vitro.

Our laboratories were involved in the initial discovery of NERT in lentivirions. These studies demonstrated that reverse transcription could be driven in replication-competent virions by microenvironments containing physiological concentrations of polyamines and dNTPs. This process increases viral infectivity, being especially prominent in resting cells, and may

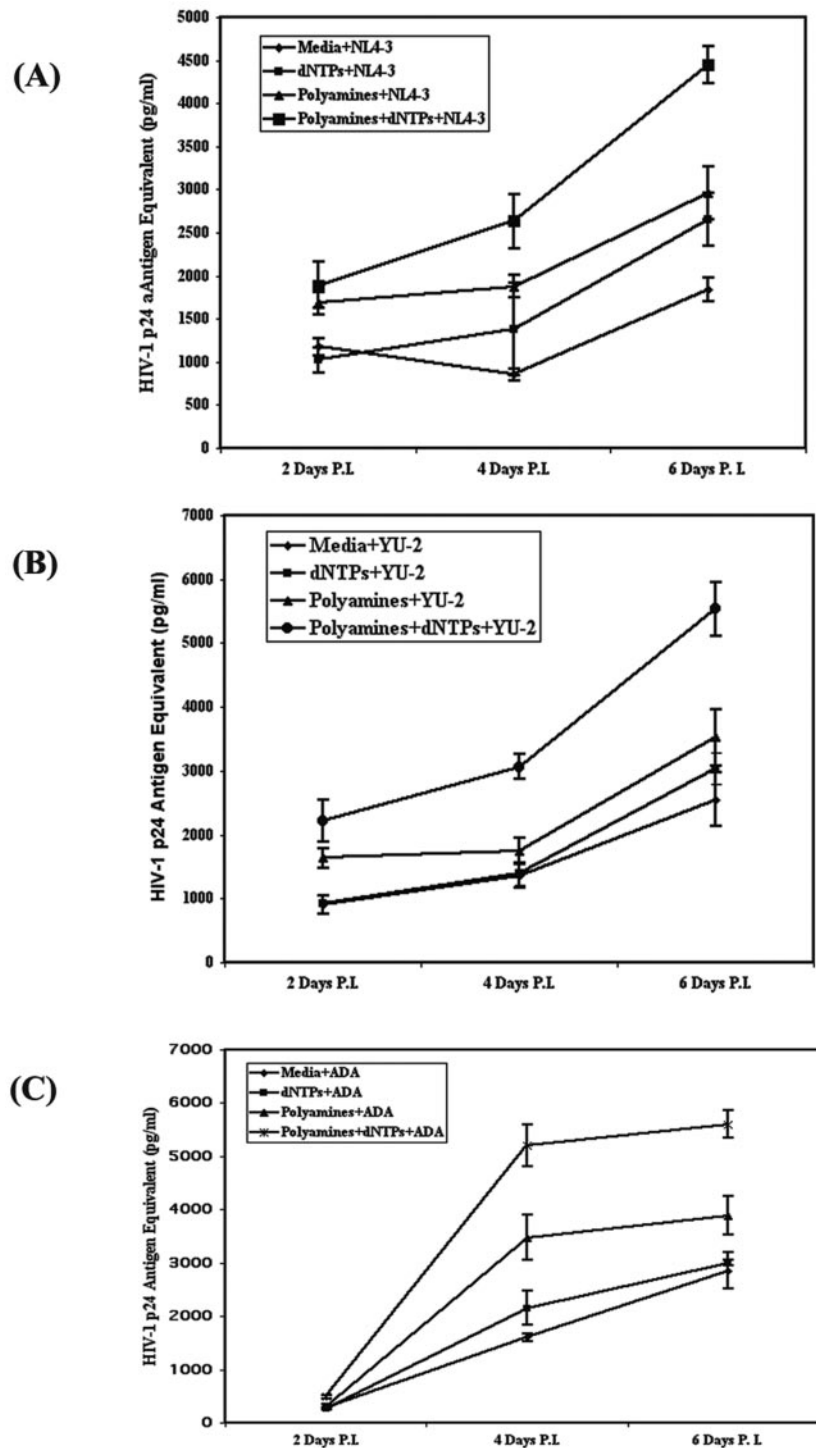


FIG. 2. Natural endogenous reverse transcription (NERT) stimulation of HIV-1. Virions increase infectivity on human oral keratinocytes. Primary human keratinocytes were grown to confluence, and then 100 ng of p24 antigen equivalents of the CXCR4-tropic HIV-1 strain, NL4-3 (A), and the CCR5-tropic HIV-1 strains YU-2 (B) and ADA (C) were treated with either medium or NERT-stimulating agents (the polyamines spermine [3 mM] and spermidine [0.1 mM] and/or dNTPs [100 nM], which are the physiological concentrations in human semen), prior to infection of 1×10^6 primary human oral keratinocytes. Certain viral inputs were treated with dNTPs alone and others with polyamines alone, while still others were treated with both polyamine and dNTPs. After infection for 4 h, the cells were vigorously washed three times with medium and viral growth was monitored for 1 week. Viral outgrowth using ELISA for HIV-1 p24 antigen expression in the culture supernatants was analyzed. These data are representative of three independent experiments performed in triplicate. P.I., postinfection.

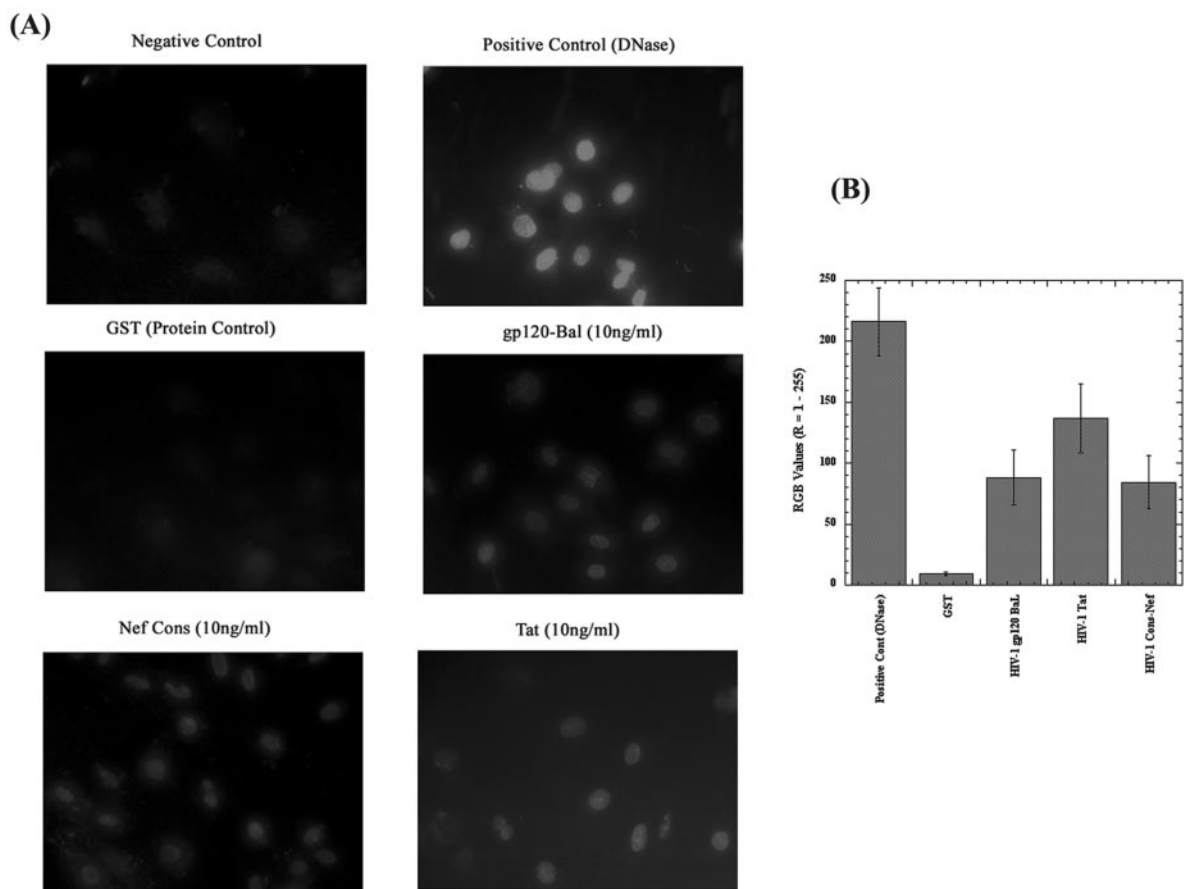


FIG. 3. (A) TUNEL assays of human oral keratinocytes treated with selected HIV-1-specific proteins. Representative TUNEL staining of HIV-1 Nef-, gp120-, and Tat-treated primary human oral keratinocytes is shown. Cells were exposed to HIV-1 Nef, gp120, and Tat for 72 h, after which cells were subjected to TUNEL staining. The concentration of protein was 10 ng/ml. As a positive control, keratinocytes were exposed to DNase at a concentration of 10 μ g/ml for 10 min at room temperature. TUNEL assays were performed using an in situ cell death detection kit, TMR Red (Roche Diagnostics). The cells were analyzed with fluorescence microscopy (Olympus System microscope, model BX60, with fluorescence attachment BX-FLA). The top panels illustrate a negative control and a positive control with apoptosis-induced nuclei of primary oral keratinocytes after treatment with nuclease (DNase; Life Technologies). These results are representative of experiments performed in triplicate and repeated at least twice. The other photomicrographs of the figure illustrate apoptosis-induced nuclei of cells treated with selected HIV-1-specific proteins. Nef Cons, Nef consensus sequence protein. (B) Semiquantitative analysis of apoptosis in primary human oral keratinocytes, using red, blue, and green values ($R = 1$ to 255). The details of this methodology for evaluating TUNEL-positive cell numbers are presented in Materials and Methods.

increase transmissibility of the virus in vivo. In a study by Zhang et al. (129), strong-stop negative-strand DNA was detected in HIV-1 virions obtained from a cell culture system. Most importantly, it was demonstrated that without detergent, addition of dNTPs to isolated HIV-1 virions could stimulate RT activity in these extracellular virions, leading to higher levels of virions carrying strong-stop negative-strand moieties and more-complete negative-strand intermediates. This was demonstrated at nanomolar concentrations of dNTPs. As well, several human physiological fluids, including seminal and blood plasma, could drive intravirion HIV-1 DNA synthesis. This was hypothesized to be due to dNTPs and polyamines within these fluids (129). These data suggested that in physiological fluids, HIV-1 virions might synthesize significant quantities of partial, and possibly full-length, reverse transcripts. This hypothesis was then supported by our initial data in which HIV-1 virions, isolated from the peripheral blood of infected individuals, were demonstrated to harbor HIV-1 DNA (for a

TABLE 2. cDNA microarray analyses of human oral mucosa (keratinocytes) exposed to extracellular HIV-1 gp120 and GST protein as a control

Gene	Severalfold increase in gene expression
TNF receptor superfamily member 1.....	2.00
Retinoblastoma binding protein 4	2.30
Tumor protein p53.....	1.80
NF- κ B	2.40
Cyclin dependent kinase 1 (Cdk 1)	2.50
Proliferating cell nuclear antigen.....	-2.00 ^a
Caspase 3.....	1.7
Caspase 6.....	1.71
Caspase 8.....	1.83
Bcl-2.....	-2.50
Growth arrest-specific gene 1 (GAS 1).....	2.15
Cyclin B1	1.80
MAPK/ERK.....	2.00

^a Negative (-) signs indicate genes that were moderately down-modulated.

TABLE 3. cDNA microarray analyses of human oral mucosa (keratinocytes) exposed to extracellular HIV-1 Nef and GST protein as a control

Gene	Severalfold increase in gene expression
Tumor protein p53	2.00
DAXX	2.00
Mitogen-activated protein kinase (MAPK) 7	2.50
Caspase 3	1.5
Caspase 6	1.8
Caspase 8	1.71
Caspase 9	1.83
Caspase 10	2.45
Proline dehydrogenase (oxygenase)	2.15
TNF receptor superfamily member 1	1.85
TNF receptor superfamily member 12	1.50
Ras homolog gene family member A	1.75
Metalloproteinase-disintegrin TNF- α convertase (TACE)....	2.00
Bcl-xL	-1.75 ^a

^a Negative (-) signs indicate genes that were down-modulated.

review, see references 130 and 131). Based on the studies in the present report, it could be hypothesized that these molecular processes at the oral mucosa may also alter interhost vertical and horizontal transmission of HIV-1.

To test the ability of NERT-stimulating agents in augmenting HIV-1 infection of primary human oral keratinocytes in vitro, we exposed the cells to HIV-1 NL4-3, ADA, and YU2. Our results show that oral keratinocytes are productively infected by both R5- and X4-tropic HIV-1 isolates and that R5 isolates may do so most efficiently. These findings are fully consistent with the results of other studies (70, 71, 132). However, Liu et al. (61) have reported productive infection of oral keratinocytes with X4- and dual-tropic isolates (X4/R5) and minimal infection with R5 isolates. In contrast, Quinones-Mateu et al. (91) could not demonstrate infection of oral keratinocytes with either X4 or R5 isolates. The apparent discrepancy

TABLE 4. cDNA microarray analyses of human oral mucosa (keratinocytes) exposed to extracellular HIV-1 Tat and GST protein as a control

Gene	% Increase in gene expression
PCTAIRE protein kinase 3 (apoptosis related).....	-1.3 ^a
Cyclin C	18.5
Mitogen-activated protein kinase (MAPK) 7	2.4
E2F transcription factor 5 (transcription activators and repressors)	-62.0
Cyclin-dependent kinase inhibitor.....	28.0
Transcription factor Dp-2.....	17.8
Caspase 9 (apoptosis related).....	30.0
Caspase 10 (apoptosis related).....	25.0
Proline dehydrogenase (oxygenase)	31.0
TNF receptor superfamily member 12	17.6
Insulin-like growth factor 1	27.0
Insulin-like growth factor binding protein 6.....	9.0
Retinoid X receptor beta (nuclear receptor, death receptor)	34.0
Ras homolog gene family member A.....	22.0

^a Negative (-) signs indicate genes that were down-modulated.

between the results of our study and those of the other studies may be explained by the differing experimental conditions which were used. For example, the amount of virus used by Liu et al. to infect their cells was about 30-fold higher than the level we used in our study. We also did not add Polybrene in our viral infections, as we have found this to be toxic to these cells and also because it is not representative of the physiological environment. In addition, Liu et al. used 2-week-old cultures of oral keratinocytes, whereas ours were approximately 3 days old. Liu et al. (61) also reported minimal expression of CCR5 in comparison with the expression of CXCR4. It is not clear whether the age of their cultures adversely affected the expression of CCR5 in oral keratinocytes, since other studies have reported the expression of this receptor at levels comparable to that of CXCR4 (50, 70, 71).

In addition to our observations of the infection of oral keratinocytes by both HIV-1 R5 and X4 isolates, we also demonstrated that stimulation by NERT increases infectibility of these cell-free virions on oral mucosal epithelium, as well as virus newly produced from HIV-1-infected cell types. Importantly, these factors model oral sexual contact and/or oral inflammation. Higher concentrations of these agents may not only increase HIV-1 replication but also increase HIV-1-specific protein-induced apoptosis (programmed cell death) in target cells (see below).

HIV-1 proteins and apoptosis in oral keratinocytes. One of the hallmarks of HIV-1 infection is programmed cell death of infected cells (9, 12). Of note, HIV-1 proteins individually have also been shown to induce and/or augment apoptosis in various primary isolated human cells, as well as in established cell lines in vitro (9, 12, 82, 83). Thus, in addition to studying the effects of NERT in augmenting HIV-1 infection of oral keratinocytes, we also sought to discern the role of HIV-1 proteins on oral mucosa cells. To validate the experiments in this study, two complementary controls were utilized, a non-HIV-1 recombinant protein expressed in our laboratories, GST, and non-treated cells. In the present study, we observed that several selected HIV-1 proteins mediated apoptosis of human oral keratinocytes. Previous studies have suggested that apoptosis (or programmed cell death) of CD8⁺ and/or CD4⁺ T lymphocytes may play a major role in AIDS pathogenesis (81). Increased apoptosis in neurons, astrocytes, and microvascular endothelial cells in the central nervous system of macaques infected with a neurovirulent strain of simian immunodeficiency virus has been described (2). Recent studies have also shown in vitro apoptosis induced by HIV-1 gp120 and/or SDF-1 α/β , binding to the chemokine coreceptor, CXCR4, on certain T-lymphocyte as well as neuronal cell populations (3, 45, 59, 74, 103), and HIV-1 envelope-specific apoptosis was demonstrated in primary brain mixed-cell cultures (78). The present study adds to these data and shows that HIV-1 Tat, gp120, and Nef induced substantial apoptosis in primary human oral keratinocytes.

In addition to HIV-1 gp120, which has been well studied in previous apoptosis experiments, other lentiviral proteins were evaluated for their effects on oral keratinocytes. Tat is an HIV-1 regulatory protein and the major transcriptional transactivator of this lentivirus (58). Tat is expressed in the early stage of infection from multiply spliced HIV-1 mRNA and, importantly, is also excreted into the extracellular milieu from

infected cells (33). Tat can be taken up by noninfected cells at levels which, when internalized, can alter expression from the HIV-1 long terminal repeat and the promoters of certain cellular genes, both directly and by increasing nuclear factor-kappa B (NF- κ B) levels (33). HIV-1 Tat may also induce T-lymphocyte cell death by apoptosis (98). Thus, HIV-1 Tat has been shown to have deleterious effects on certain cells *in vitro*. This was clearly demonstrated in the present study for human oral keratinocytes.

The role of Nef in inducing apoptosis in T lymphocytes is quite controversial (28, 79, 93, 110). In a recent study, we observed that baculovirally expressed consensus HIV-1 recombinant Nef induced apoptosis in primary human brain microvascular endothelial cells at concentrations of 10 to 100 ng/ml. These apoptotic effects of Nef were augmented with the addition of 0.3% ethanol (1). In the present study, we observed Nef-mediated apoptosis in primary human oral keratinocytes. It therefore appears that contrary to the results of certain studies of other cell types, HIV-1 Nef induces apoptosis, *in vitro*, in human oral mucosal cells.

Mechanisms of apoptosis in oral mucosal cells induced by HIV-1 proteins. The molecular mechanisms involved in apoptosis of primary human oral keratinocytes are mostly unknown. Therefore, to begin to elucidate the molecular mechanisms involved, we employed targeted gene microarrays to analyze the regulation of apoptotic genes in primary human oral keratinocytes exposed to soluble HIV-1 gp120, Nef, and Tat proteins. The gene microarray analyses demonstrated that Tat-treated primary human oral keratinocytes induced increased expression of TNFR12, whereas both gp120- and Nef-treated oral keratinocytes induced up-regulation in the expression of TNFR 1 and caspases 6 and 8. In addition, Nef up-regulated the expression of DAXX, a protein that has also been implicated in apoptosis.

Apoptosis is comprised of two major pathways, the mitochondrial apoptotic pathway and the Fas/Fas ligand plus the TNF/TNFR-1 apoptotic pathways (14, 53, 105). In all cases, the downstream target of these apoptotic pathways is the activation of caspase 3. Caspase 8 is the downstream indicator of activation of the Fas-Fas ligand pathway, whereas caspase 9 is associated with the mitochondrial apoptotic pathway (53, 105). Receptors in the TNF receptor super family are involved with the induction of apoptosis as well as inflammatory signaling. In response to Fas ligation, Fas-associated death domain protein binds and recruits caspase 8 to form the death-induced signaling complex, thereby activating Fas apoptotic signaling (13, 66, 76). It has also been shown that upon Fas engagement, molecules such as DAXX may also bind to Fas and mediate Fas-associated death domain protein-independent apoptotic signaling (54). Thus, the fact that in the present studies, genes involved in the TNF/TNF-R and Fas/Fas ligand pathways were found up-regulated suggests the involvement of the TNF/TNF-R and Fas/Fas ligand pathways in HIV-1 gp120-, Nef-, and Tat-induced apoptosis in primary human oral keratinocytes. These results are consistent with previous studies which suggest that Nef (41-44), gp120 (125), and Tat (126, 127) up-regulate the expression of genes in the TNF/TNF-R and Fas/Fas ligand pathways, resulting in apoptosis. For instance, the HIV-1 proteins gp120 (96, 115), Tat (32, 120), and Nef (128) have been reported to increase CD95 and FasL levels, which

enhance cellular susceptibility to Fas-mediated apoptosis. Nef, on the other hand, is thought to induce FasL expression by interacting specifically with the zeta chain of the TCR complex (124), whereas Tat has been shown to up-regulate the expression of the initiator, caspase 8 (11).

In addition to the induction of the TNF/TNF-R and Fas/Fas ligand pathways, oral keratinocytes exposed to Nef showed up-regulation in the expression of caspases 3, 6, 8, 9, and 10, whereas Tat-induced expression of caspase 9 and 10 and gp120 induced the expression of caspases 3, 6, and 8. The caspase 9 activation pathway involves the release of cytochrome *c* from mitochondria, which associates with pro-caspase 9 and Apaf-1 in the presence of dATP to form a proapoptotic complex (53, 105). The pro-caspase 9-Apaf-1-cytochrome *c* complex so formed induces autoactivation of caspase 9 by aggregation (133). Once activated, caspase 9 initiates a cascade involving caspase 3, 6, and 7, which act by themselves to cleave cellular targets such as poly(ADP-ribose) polymerase, leading to cell death (133). In this study, we also observed increases in the expression of various caspases when mucosal cells were exposed to HIV-1 gp120, Nef, and Tat. This increase in caspase 9 expression in the cells exposed to HIV-1 Nef and Tat suggests that induction of apoptosis by these specific proteins may also involve the mitochondrial apoptotic pathway. Furthermore, this observation supports the hypothesis that synergy between the upstream caspase pathways may have an important role in HIV-1 proteins-induced oral keratinocyte apoptosis. Together, the data presented herein support an important function for caspases in HIV-1-induced injury and apoptosis in this cell type.

In addition to the TNF/TNF-R and Fas/Fas ligand pathways and the caspase cascades initiated by exposure of oral keratinocytes to gp120, Nef, and Tat, cells exposed to Nef exhibited down-regulation in the expression of Bcl-xL. Similarly, gp120 induced down-modulation in the expression of Bcl-2. Bcl-2 is a human proto-oncogene located on chromosome 18. Its product is an integral membrane protein (entitled Bcl-2) located in the membranes of the endoplasmic reticulum and nuclear envelope and in the outer membranes of the mitochondria. At least 16 members of the Bcl-2 family of proteins have been sequenced in recent years, both in mammals and in viruses (19, 95, 112). The Bcl-2 family of proteins regulates the mitochondrial potential and early apoptotic changes in cells (75). Some of these, such as Bcl-2, Bcl-xL, Bcl-w, or Mcl-1, protect cells against Fas-induced programmed cell death by decreasing the permeability of the mitochondrial membrane, while others, such as Bad, Bak, Bax, Bid, Bim, or Hrk, induce apoptosis (96). The antiapoptotic activity of these proteins is related to their ability to form heterodimers (96). For instance, in murine FL5.12 cells, Oltvai et al. (80) showed that overexpression of Bax after withdrawal of interleukin 3 led to the formation of homodimers, resulting in programmed cell death. Overexpression of antiapoptotic proteins such as Bcl-2 and Bcl-xL resulted in the formation of Bcl-2/Bax or Bcl-xL/Bax heterodimers, thereby inhibiting the lethal functions of Bax.

Although the mechanism of apoptosis regulation by Bcl-2 is not yet fully understood, whether or not a cell becomes susceptible to programmed cell death may depend on the balance between proapoptotic proteins, such as Bax, and antiapoptotic proteins, such as Bcl-2 or Bcl-xL. It has been reported that the

HIV-1-encoded proteins gp120, Tat, and Nef inhibit expression of Bcl-2, and Nef alone inhibits the expression of Bcl-xL (4, 100). In the present study, moderate down-modulation in the expression of Bcl-2 was observed in keratinocytes exposed to gp120. Similarly, oral keratinocytes exposed to Nef exhibited moderate decreases in the expression of Bcl-xL. The decrease in the relative amounts of these proteins in this study is likely important, as increased quantities of antiapoptotic proteins may be able to overwhelm even larger numbers of proapoptotic signals. The results also suggest that the reduced expression of antiapoptotic Bcl-2 family of proteins in gp120- and Nef-treated cells might have disturbed the balance between the pro- and antiapoptotic regulators, thereby promoting proapoptotic signaling in response to exposure to these HIV-1 proteins. Thus, these results further confirm the involvement of the Fas/FasL apoptotic pathway in both gp120- and Nef-induced apoptosis in oral keratinocytes.

Besides the expression of Bcl-2 proteins, our targeted microarray analyses also revealed the up-regulation in the expression of p53, NF- κ B, MAPK/extracellular signal-regulated kinase (ERK), growth arrest-specific gene 1 (GAS) 1, retinoblastoma binding protein 4, cyclin B1, and cyclin-dependent kinase 1 (Cdk 1) in primary human oral keratinocytes exposed to gp120. There was down-modulation in the expression of proliferating cell nuclear antigen (PCNA) in these same cells. Oral keratinocytes exposed to Nef, on the other hand, demonstrated up-regulation in the expression of the tumor protein p53, mitogen activated protein kinase (MAPK) 7, and Ras homolog gene family member A. Similarly, Tat induced increased expression of Ras homolog gene family member A, MAPK 7, and cyclin C plus other genes and down-regulated the expression of the apoptotic-related PICTAIRE and E2F transcription factor 5.

Ras is an essential component of signal transduction pathways that control cell proliferation, differentiation, and survival (73, 104). The gene and its expression products have been established as modulators of apoptosis and, depending on the cell type and the presence of other proapoptotic or antiapoptotic stimuli (21, 30), can either inhibit or promote apoptosis. Numerous studies have shown that Ras activity is manifested through ceramide signaling in the Fas/Fas ligand apoptotic pathway and that overexpression of *ras* accelerates the Fas-induced death process (21, 47). For example, in human or mouse lymphocytes, it has been shown that expression of Ras resulted in Fas-induced apoptosis (48), and this apoptotic process was blocked by the overexpression of *bcl-2* (21, 22). Similarly, it has been demonstrated in Jurkat T-cells that overexpression of *ras* resulted in sensitization of the cells to Fas engagement and low levels of expression of Bcl-2 (22, 35). In a study to examine the cellular responses to high-intensity Ras signaling, Joneson and Bar-Sagi (52) reported that the expression of increasing amounts of the oncogenic form of human HRas, HRasV12, resulted in a dose-dependent induction of apoptosis in both primary and immortalized cells (52). At low levels of Ras expression, this group reported that the antiapoptotic pathway is activated to an extent that is sufficient to counteract the apoptotic signals. At high levels of Ras signaling, this balance was tilted in favor of the proapoptotic signals presumably because the component(s) of the antiapoptotic pathway are limiting. The apoptotic effect of HRasV12 re-

quires the activation of both the ERK and Jun N-terminal protein kinase (JNK) mitogen-activated protein kinase cascade and is independent of the tumor protein, p53 (52). It is important to note that the p53-independent apoptotic effects may be exclusive to the HRasV12, since the expression of the molecule has been shown to play important roles in apoptosis (67, 68). For instance, wild-type p53 activates BH-3-containing proteins and mediates the down-modulation of Bcl-2 expression and up-regulation of Bax, thereby predisposing the cells to apoptosis (67, 68).

Our targeted microarray analyses clearly demonstrated that HIV-1 Nef and gp120 down-modulated the expression Bcl-xL and Bcl-2, respectively. Also, Nef and Tat induced increased expression of Ras mRNAs in primary oral keratinocytes. Taken together, these data suggest that HIV-1 Nef, gp120, and Tat-induced apoptosis in oral keratinocytes occurs potentially through the down-modulation of Bcl-2 and Bcl-xL and the up-regulation of Ras expression. It is possible that, in the present studies, the reduced expression of Bcl-2, in the case of gp120, and Bcl-xL, in the case of Nef, may have prevented the binding of these proteins to Ras, thereby enhancing the apoptotic signaling mediated by Ras, during the Fas-mediated apoptotic process. Furthermore, p53-dependent apoptotic pathways may have contributed to gp120- and Nef-elicited Bcl-2 expression.

The tumor protein, p53, is constitutively expressed in cells, and when bound to the ubiquitinase-like enzyme HDM2, is transported to the nucleus from the cytoplasm. The binding of p53 to the ubiquitinase-like enzyme HDM2 results in the degradation of p53 (39), and phosphorylation of p53 reduces its affinity for HDM2, thereby causing an increase in the levels of p53. Phosphorylated p53 induction mediates the down-modulation of Bcl-2 expression and up-regulation of Bax, thereby predisposing the cells to cells-to-cell cycle arrest and apoptosis (67, 68). Furthermore, Cande et al. (15) recently reported that activation of cyclin-dependent kinase-1 (Cdk1) resulted in the activation of mammalian target of rapamycin (mTOR), phosphorylation of p53 at S15, and apoptosis. They also showed that neutralization of p53 expression in peripheral blood mononuclear cells from HIV-1-infected patients suppressed apoptosis. These cells also exhibited an increase in cyclin B and mTOR expression, correlating with p53^{S15} phosphorylation and viral load. Thus, it appears that the sequential activation of cyclin B-Cdk1, mTOR, and p53 leads to induction of apoptosis. Unlike p53, MAPK and MAPK/ERK have not been directly linked to the mitochondrial apoptotic pathway. Also, the relationship between the activation of MAPK cascades and induction of apoptosis is still not clear and may differ among different cell types (118, 122).

The HIV-1 envelope protein gp120 elicits the transcriptional activation of p53 (18), resulting in the activation of the *bax* gene (69) and subsequent apoptosis. In a recent publication, Perfettini et al. (86) reported that p53 and NF- κ B are the dominant apoptosis-inducing transcription factors elicited by the HIV-1 envelope. Utilizing microarray technology, they identified several HIV-1 envelope-elicited, p53-dependent proapoptotic transcripts that include Puma, a proapoptotic "BH-3-only" protein from the Bcl-2 family, which is known to activate Bax/Bik. These authors further reported that p53 inhibition prevented the expression of several proapoptotic

genes and disabled the apoptotic machinery. Furthermore, Castedo et al. (18) have suggested that in gp120-elicited apoptosis, gp120 interacts with CD4 and CXCR4 to form a gp120/CD4/CXCR4 complex. The formation of this complex transiently results in the up-regulation in the expression of cyclin B at the protein level, which in turn induces cyclin B-dependent kinase 1-dependent (cyclin B/Cdk 1) activation of the mTOR, a serine/threonine kinase of the phosphatidyl inositol kinase family. Accumulation of the mTOR in the nucleus results in the direct phosphorylation of p53 at the Ser15 residue and the subsequent transcriptional activation of proapoptotic Bcl-2 protein family member Bax. This ultimately causes a reduction in the mitochondrial membrane potential and the release from the mitochondria of cytochrome *c* and other proapoptotic proteins, particularly apoptosis-inducing factor 1 (AIF), resulting in programmed cell death (17, 36, 37). Note also that retinoblastoma binding protein 4, which was demonstrated to be up-regulated in oral keratinocytes exposed to gp120, is a substrate for Cdk1 and has been found to be induced in apoptotic cells (46).

Cyclin B1 is a key regulator of apoptosis in some cell types (88). Recently, it was reported that the cyclin B1 protein is important for the induction of radiation-induced apoptosis in hematopoietic cells (89). It has also been shown that accumulation of the protein is necessary for the induction of apoptosis caused by T-cell-receptor activation (38). Recently, it was demonstrated that induced cell death required the expression of cyclin B1 in T cells undergoing activation and that overexpression of the protein enhances the expression of *FasL*, which may ultimately lead to cell death (116). This effect of cyclin B1 is mediated by the binding of the protein to Cdk1 to form the cyclin B1/Cdk1 complex. The regulation of the transcription of *FasL* by the cyclin B1/Cdk1 complex is achieved through the control of NF- κ B activation. Overexpression of the NF- κ B subunit, Rel A, enhanced the activation and induction of *FasL* promoter activity and subsequent cell death (116).

Consistent with the HIV-1 gp120-induced apoptotic pathway described above, our microarray analyses demonstrated concomitant up-regulation in the expression of p53, NF- κ B, MAPK/ERK, growth-arrest-specific gene 1 (GAS 1), retinoblastoma binding protein 4, cyclin B1, and cyclin-dependent kinase 1 (Cdk 1) in primary human oral keratinocytes exposed to gp120. The formation of cyclin B1/Cdk1 complex might have contributed to gp120-induced *FasL* expression and subsequent programmed cell death of primary oral keratinocytes. The data also strongly suggest that in addition to the *Fas/FasL* and TNF/TNFR apoptotic pathways, gp120 might induce apoptosis in primary human oral keratinocytes via the mitochondrial apoptotic pathway.

The proliferating cell nuclear antigen (PCNA), whose expression was up-regulated by gp120, is a protein involved with the major DNA replication and repair machinery of the cell (55, 64). Depending on the physiological conditions of the cell, the protein may perform several functions in mammalian cells. For instance, in the absence of p53, the presence of large quantities of PCNA results in DNA replication in cells. The presence of p53 leads to DNA repair. However, low quantities or a complete absence of p53 may drive the cells into apoptosis (85). Consistent with this idea, there was a decrease in the expression of PCNA in keratinocytes exposed to gp120, and

this may have led to apoptosis. The increase in the expression of growth-arrest-specific gene 1 (GAS 1) may have also contributed to overall apoptotic mechanism, since the gene is known to inhibit DNA synthesis in vivo (24) and also to act as a mediator of apoptosis (49).

Nef possesses the capacity to regulate numerous transcriptional activities by targeting and controlling various *src* family kinases and MAPK and p53 activities. However, most reports regarding Nef-mediated transcriptional activation of p53 and MAPK pathways have been less than conclusive. Some of these studies suggest that Nef inhibits while others suggest that Nef enhances p53 and MAPK activities (63, 111). Nevertheless, it appears that the regulation of the transcriptional activities of various kinases and p53 by Nef is very complicated, and whether or not Nef inhibits or enhances these activities may depend on the cell type and possibly on the nature of the Nef protein involved. In the present study, Nef was found to up-regulate the expression of p53, and oral keratinocytes exposed to Nef and Tat were found to undergo MAPK cascade activation. In view of the fact that these proteins have been shown in some studies to contribute to the induction of apoptosis and that Nef and Tat were also found in the present study to up-regulate the expression of other proapoptotic genes, it could be argued these HIV-1 proteins might modulate the sensitivity of oral keratinocytes to proapoptotic molecules, thus contributing to the induction of programmed cell death. This is consistent with other studies that have shown that Tat and Nef induce certain cells to undergo MAPK cascade activation and apoptosis (27, 117). Thus, the complexity of the differential gene expression regulated by p53, as identified by our targeted DNA microarray in primary human oral keratinocytes exposed to all three HIV-1 proteins, suggests that a diverse array of genes may potentially play synergistic or additive roles in p53-mediated mitochondrial apoptosis in these cells in vitro.

In addition to the increased expression of p53-regulated genes by keratinocytes exposed to gp120, Nef, and Tat, it was also observed that both Nef and Tat induced increased expression of proline dehydrogenase. Nef alone induced the expression of metalloproteinase-disintegrin TNF- α convertase (TACE or ADAM17), and Tat induced the expression of PCTAIRE protein kinase, insulin-like growth factor 1 (IGF-1), insulin-like growth factor binding protein 6 (IGFBP-6), and retinoid X receptor beta.

Proline dehydrogenase, a mitochondrial enzyme, is involved in the transfer of redox potential across the mitochondrial membrane through the proline/delta-P5C (proline-P5C) pathway (87). In vitro, it has been shown that the proline dehydrogenase gene is induced early by the tumor suppressor protein p53 and that it inhibits cell proliferation plus localizes to the mitochondria and triggers programmed cell death. Similarly, TACE and the insulin-like growth factor binding protein 6 (IGFBP-6) are inducers of apoptosis whereas IGF-1 inhibits apoptosis. TACE, a key enzyme (shedase) that releases TNF- α from its inactive cell-bound precursor (119), has been shown to induce apoptosis in U937 cells via the release of TNF- α and soluble *FasL* (114).

IGFBP-6 is a family of six high-affinity secreted proteins that modulate the availability and hence the antiapoptotic actions of the IGFs. Thus, IGF-1 and IGFBP-6 exert opposing effects on cell growth and apoptosis (77, 113). Unlike proline dehy-

drogenase, TACE, and IGF-1 is the most prevalent and potent survival factor for many cells. Studies have shown that IGF-1 inhibits apoptosis, promotes matrix formation, and can decrease TNF- α -induced proteoglycan degradation and apoptosis in numerous cells (8).

Given their proapoptotic role, the production of proline dehydrogenase, TACE, and IGF-1 in keratinocytes exposed to gp120, Nef, or Tat might have impacted the induction of apoptosis favorably. The increase in the expression of IGF-1 by keratinocytes exposed to Tat may have been an attempt by these cells to counteract the effects of IGF-1 and apoptosis. Indeed, some studies suggest that Tat triggers IGF-1 gene expression, resulting in IGF-1-induced resistance to apoptosis in several cell types (27, 121).

The role of HIV-1 Tat in inducing apoptosis in cells is not quite clear. Tat has been described as pro- and antiapoptotic depending on the cell type, microenvironment, and culture conditions. For example, it has been shown in vitro that Tat induces apoptosis of uninfected T cells (65) through a Fas/FasL apoptotic pathway (120), whereas it exerts the opposite effect in HIV-1-infected T cells (65). Similarly, this protein has been shown to be antiapoptotic in different tumor cell lines of neuronal, epithelial, and lymphoid origin (16). Tat is able to trigger the expression of several genes responsible for cytokines and growth factors, and the differential expression of these may determine the sensitivity of cells to apoptosis (27). In the present study, Tat elicited the expression of numerous proapoptotic and growth factors. Collectively, the increase in the expression these genes may have resulted in the release of proapoptotic proteins, cytokines, and growth factors released in the cell microenvironment, leading to the proapoptotic effects of Tat.

In summary, the present studies strongly suggest, for the first time, that infection of human oral keratinocytes can be augmented by microenvironments containing physiological concentrations of polyamines and dNTPs and that cell-free lentiviral proteins may lead to programmed cell death in these cells. The data regarding NERT augmentation of HIV-1 infection of oral mucosa cells suggest that seminal fluids, and possibly cervical-vaginal fluids, in which NERT-stimulating molecules are at relatively high concentrations may antagonize the HIV-1 salivary inhibitors. In addition, cell-free HIV-1-specific proteins (i.e., Tat, Nef, and gp120) may lead to the disruption of the oral mucosal barrier in vivo. As such, these represent two potential molecular mechanisms by which HIV-1 infects humans after orogenital contact with infected individuals and by which infants of HIV-1-infected women may acquire HIV-1 infection perinatally, during vaginal delivery.

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