

The Expression of the Interleukin 6 Gene Is Induced by the Human Immunodeficiency Virus 1 TAT Protein

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Summary

Human immunodeficiency virus 1 (HIV1) infection is associated with severe psoriasis, B cell lymphoma, and Kaposi's sarcoma. A deregulated production of interleukin 6 (IL-6) has been implicated in the pathogenesis of these diseases. The molecular mechanisms underlying the abnormal IL-6 secretion of HIV1-infected cells may include transactivation of the IL-6 gene by HIV1. To test this hypothesis, we used the pIL6Pr-chloramphenicol acetyltransferase (CAT) plasmid, an IL-6 promoter-CAT construct, as a target of the transactivating function of the HIV1 TAT protein. By cotransfecting the pIL6Pr-CAT and the *tat*-expressing pSVT8 plasmid in MC3 B-lymphoblastoid or in HeLa epithelial cells, we observed that TAT transactivates the human IL-6 promoter. These results were confirmed when pIL6Pr-CAT was transfected in MC3 or HeLa cells that constitutively expressed the *tat* gene in a sense (pSVT8 cells) or antisense (pSVT10 cells) orientation. 5' deletion plasmids of pIL6Pr-CAT, in which regions at -658, -287, and -172 were inserted 5' to the *cat* gene, were transiently transfected in pSVT10 and pSVT8 cells and showed that TAT-induced activation of the IL-6 promoter required a minimal region located between -287 and -54 bp. Moreover, experiments with plasmids carrying the -658, -287, and -172 bp regions of the IL-6 promoter inserted downstream to a TAR-deleted HIV1-LTR identified the sequence of -172 to -54 as the minimal region of the IL-6 promoter required for TAT to transactivate the TAR-deleted HIV1-LTR. By DNA-protein binding experiments, *tat*-transfected cells expressed a consistent increase in κ B and nuclear factor (NF)-IL-6 binding activity. Accordingly, the pDRCAT and IL-1REK9CAT, carrying tandem repeats of NF- κ B or NF-IL6 binding motifs, respectively, were activated in TAT-expressing cells. The biological relevance of the TAT-induced IL-6 secretion was addressed by generating 7TD1 cells, an IL-6-dependent mouse cell line, stably expressing the *tat* gene. These *tat*-positive cells expressed the endogenous IL-6 gene, secreted high amounts of murine IL-6, and grew efficiently in the absence of exogenous IL-6. Moreover, the *tat*-positive 7TD1 cells sustained the growth of parental 7TD1 cells and showed a dramatic increase in their tumorigenic potency. These results suggest that TAT protein may play a role in the pathogenesis of some HIV1-associated diseases by modulating the expression of host cellular genes.

HIV1 infection causes various clinical and immunological abnormalities, including lymphadenopathy, activation of polyclonal B cells that manifests as hypergammaglobulinemia and auto-antibody production, Kaposi's sarcoma, and lymphoma of the B cell phenotype (1-3). In HIV1-infected subjects, there is a deregulated production of cytokines, that affects the growth and differentiation of lymphoid and mesenchymal cells, and that may contribute to the development of the clinical features of AIDS. IL-6 has a broad range of biological effects (4) and plays a major role in the

development of B cell malignancies and Kaposi's sarcoma (5-7). IL-6 gene transcription is induced in cells infected by HIV1, and increased levels of IL-6 have been reported in serum and cerebral spinal fluid of HIV1-infected patients (8-10). These findings suggest that HIV1 may transactivate the host cellular IL-6 gene.

The TAT protein of HIV1 is required for efficient viral gene expression (11). TAT increases the initiation of transcription from the HIV1-LTR (12) and affects RNA processing and utilization by interacting with a TAT-responsive element

(TAR) located between nucleotide +1 and +44 with respect to the initiation site (+1) of viral transcription (13). In addition, TAT interacts with upstream regulatory DNA sequences circumscribed within the nuclear factor (NF)¹-κB/Sp1 region of the HIV1 promoter (14) and with host cell proteins (15–19). TAT could affect the transcription of cell genes such as IL-6, whose regulatory sequences, like HIV1-LTR, possess NF-κB and NF-IL6 enhancer elements (20, 21). To test this hypothesis, we used the pIL6Pr-CAT, an IL-6 promoter-CAT construct, as a target of the transactivating function of the HIV1 TAT proteins. By cotransfecting the pIL6Pr-CAT and the *tat*-expressing pSVT8 plasmids (22) in MC3 B-lymphoblastoid or in HeLa epithelial cells, we observed that TAT transactivated the human IL-6 promoter. These results were confirmed when pIL6Pr-CAT was transfected in MC3 or HeLa cells that constitutively expressed the *tat* gene. The biological relevance of the *tat*-induced IL-6 secretion was addressed by generating 7TD1 cells, an IL-6-dependent mouse cell line (23), stably expressing the *tat* gene. These *tat*-positive cells expressed the endogenous IL-6 gene, grew efficiently in the absence of exogenous IL-6, and showed a dramatic increase in their tumorigenic potency.

Materials and Methods

Animals and Reagents. 4-wk-old female nude mice were purchased from Charles River Breeding Laboratories (Como, Italy). LPS *Escherichia coli* 055:B55 was obtained from Difco Laboratories (Detroit, MI). G418 (Geneticin, 658 μg of G418 base per mg) was purchased from Sigma Chemicals (Milan, Italy). Recombinant TAT protein and the rabbit anti-TAT antibody were obtained from the National Institutes of Health-AIDS Research and Reference Reagent Program. Rat monoclonal and goat polyclonal antibodies to mouse IL-6 were purchased from Boehringer Mannheim (Milan, Italy) and from Genzyme (Cambridge, MA), respectively. Elisa assays specific for human or mouse IL-6 were purchased from Genzyme. HPLC-purified human rIL-6 (sp act, 2.7 × 10⁹ U/mg) was provided by G. Ciliberto (Istituto di Ricerche di Biologia Molecolare, Pomezia, Italy).

Plasmids. To generate the pIL6Pr-CAT plasmid, the 1.2 Kbp BamHI-XhoI 5' upstream sequences of the IL-6 gene were excised from the pGEM672A plasmid (24) and inserted into compatible sites (BamHI-SalI) of the pEMBL-CAT plasmid using standard techniques (25). The 5' deletion mutants of pIL6Pr-CAT were generated by digesting the pIL6Pr-CAT with BalI, NheI, or HaellI, followed by fill-in with Klenow polymerase. The resulting constructs contained regions of -658, -287, and -172 relative to the first nucleotide of the ATG codon of the IL-6 genomic sequence (24). To generate the HIV1-LTR-IL-6 promoter fusion plasmids, the TAR-deleted EcoRI-BglII fragment of pILC-CAT (26) was isolated, filled-in, and inserted at the SstI site (filled) of the various IL-6 promoter mutants. The correct 5'-3' orientation was analyzed by multiple restriction digestions and by sequencing using the Sanger's method (25). The pSVT8 and pSVT10 plasmids, expressing the *tat* gene in a sense or antisense orientation, respectively (22), were obtained from A. Caputo (University of Ferrara, Ferrara, Italy). The β-galactosidase expressing pnlacZ plasmid was obtained from A. Weisz (University of Naples, Naples, Italy). The pDRCAT plasmid, carrying the HIV1-κB enhancer upstream to a thymidine

kinase (tk) basal promoter was provided by W. Greene (University of California at San Francisco, San Francisco, CA). The IL-1REK9CAT plasmid, carrying three copies of the NF-IL6 binding motif upstream to the IL-6 basal promoter, was donated by T. Kishimoto and T. Hirano (University of Osaka, Osaka, Japan). The pILC-CAT plasmid, carrying the HIV1-LTR and TAR region, and the pSV12 plasmid, harboring the *tat* gene, were a kind gift of A. Rabson (Center for Advanced Biotechnology and Medicine, Piscataway, NJ).

Cells, Transfection Procedures, and Analysis of the Transfected Cells. IM9, a human myeloma cell line (American Type Culture Collection, Rockville, MD), EBV-positive MC3 (6), and HeLa epithelial cells were cultured in DMEM supplemented with 10% vol/vol heat-inactivated FCS (Flow Laboratories, Milan, Italy), 3 mM glutamine, and 10 mM Hepes buffer, pH 7.2 (Gibco Laboratories, Milan, Italy). The IL-6-dependent 7TD1 cells (23) were cultured as previously reported (6). MC3 and HeLa cells were transfected by electroporation as previously described (6, 27). Briefly, cells were washed and resuspended in 0.8 ml of Ca²⁺- and Mg²⁺-free PBS (D-PBS) at a concentration of 1.5 × 10⁷/ml in the presence of 10 μg of pSV2neo control plasmid, pSVT8 or pSVT10 plasmids linearized by ScaI digestion. Cells were subjected to a single electrical pulse (0.2 kV, 960 μFD) using an apparatus (Bio-Rad Laboratories, Milan, Italy), recovered, and cultured for 48 h in DMEM supplemented with 10% FCS before selection in 3 mg/ml G-418 (Geneticin). Cells were routinely cultured in G-418-supplemented medium. The integration of the transfected plasmids was analyzed as described (27). High molecular weight DNA was isolated, digested with SalI or BamHI enzymes, electrophoresed in 0.6% agarose gel, blotted on filters (Gene Screen Plus; Du Pont, Milan, Italy), and analyzed by Southern blotting using either a 0.3-Kbp *tat* probe excised from the pSV12 plasmid by BglII digestion, or a 1.1-kbp EcoRI *neo* fragment isolated from the pSV2-neo plasmid. Specific *tat*- and human *β-actin* mRNAs from pSVT10- or pSVT8-transfected cells were analyzed by reverse transcription of total RNAs, followed by PCR amplification using the *tat* primers 5'-TTACTCCACACACCACACCAAGAA-3' and 5'-TCTCTC-TCTCTCTCCACCTTCTTCTCTAT-3', and with the *β-actin* primers 5'-ATGGATGATGATATCGCCGCG-3' and 5'-CTAGAA-GCATTTGCGGTGGACGATGGAGGGGCC-3'. Specific *tat* RNAs were visualized by electrophoresis in 1.5% agarose gel. The presence of transactivating TAT proteins was tested by transiently expressing the HIV1-LTR containing pILC-CAT plasmid (25) in mock-transfected cells and in cells expressing the *tat* gene in a sense or antisense orientation. The expression of the pIL6Pr-CAT plasmid in *tat*- or antit_{at}-positive MC3 and HeLa cells was analyzed by transiently transfecting cells with different amounts of pIL6Pr-CAT as detailed in the legend to the figures. 7TD1 cells were cultured in IL-6-supplemented medium as reported (6, 23), and transfected with the pSV2neo, pSVT8, or the pSVT10 plasmid. 48 h after transfection, cells were cultured in the presence of 2 mg/ml G-418 (Geneticin). 2 wk later, cells were tested for *tat* gene integration and for the presence of *tat* transcripts and TAT transactivating proteins as detailed above. Cells were then cultured without exogenous IL-6. The expression of the endogenous mouse IL-6 gene was analyzed by RNA reverse transcription and PCR amplification using the primers: 5'-ATGAAGTTCTCTCTGCAAGGACT-3', and 5'-CACTAGGTTTGCCGAGTAGATCTC-3'. Amplified PCR products were electrophoresed in 1% agarose gel, transferred onto nylon membranes (Gene Screen Plus) and hybridized with a 0.6-kbp EcoRI-HindIII fragment of mouse IL-6 cDNA probe (28). 7TD1-pSV2neo control cells were generated by transfection of 7TD1 cells with pSV2neo followed by G418 selection in the presence of exogenous IL-6. To evaluate the capacity of the 7TD1-pSVT8 cell

¹ Abbreviations used in this paper: CAT, chloramphenicol acetyltransferase; NF, nuclear factor; nt, nucleotide.

to sustain the growth of the IL-6-dependent 7TD1-pSV2neo cells, 7TD1-pSVneo cells were cultured (4×10^5 cells/well in 0.2 ml cultures) with 25% vol/vol of conditioned medium from 7TD1-pSVT8 cells in the absence of exogenous IL-6. Cell growth was evaluated after 72 h by cell counting and [^3H]thymidine incorporation as previously reported (6). In other experiments, 7TD1-pSV2neo cells were cocultured at 2×10^5 cells/ml with 7TD1-pSVT8 cells (2×10^5 cells/ml) in transwell plates, where 7TD1 cells (upper level) were separated by a 0.45- μm membrane from the *tat*-transfected 7TD1-pSVT8 cells (lower level). In these experiments, 7TD1 cells were counted daily. In other experiments, 7TD1 cells were cultured in the presence of 50 ng/ml of human IL-6 or with amounts of recombinant TAT protein ranging from 0.3 to 10 $\mu\text{g}/\text{ml}$. In parallel experiments, *tat*-positive 7TD1-pSVT8 cells were cultured in the presence of a rabbit polyclonal antibody to TAT at concentrations ranging from 1:100 to 1:1,000 final dilution. The proliferative rate was analyzed at 72 h as described (6).

For transient expression experiments, cells were transfected by electroporation at 5×10^6 cells in 0.8 ml of D-PBS as detailed above and cultured for 48 h in complete DMEM medium. The amounts of transfected DNA were equalized either with the pZipSVneo control plasmid or with pUC18 plasmid DNA. Moreover, in preliminary experiments, we found that cell transfections generated CAT activity that was linear in the range of 5 to 70 $\mu\text{g}/\text{ml}$ of pSV2-CAT, after which it declined. Under these conditions, CAT activity in cellular extracts from duplicate transfections varied by <15%. Transfection efficiency was monitored by cotransfecting the cells with 5 μg of pnl-lacZ plasmid. β -Gal activity was assayed using 80–100 μg of protein cell extract as described (25). A random 25% variation of the β -Gal levels was observed throughout this study. The transient expression experiments were performed at least five times with different plasmid preparations. Primer extension was carried out as described (25). 20 μg of total RNA was annealed to the oligonucleotides 5'-CAACGG-TGGTATATCCAGTG-3' (for *cat* RNA), and 5'-CAGATACTACTCTTG-3' (for U2 RNA). RNA was elongated with reverse transcriptase, digested with RNase A and separated over a 6% denaturing 7 M urea-acrylamide gel.

CAT Assays. 48 h after transfection, cells were harvested and washed once with PBS. Cell extracts were prepared by three cycles of freeze-thawing in 0.25 M Tris, pH 7.8, and CAT assays were performed as previously described (27, 29). Proteins were measured in each cell extract with an assay kit (Bio-Rad) and equal amounts of proteins were analyzed for each sample. Each assay contained 50 μg of cell extract, 20 μl of 4 mM acetyl-coenzyme A (Boehringer Mannheim), 1 μl (0.5 μCi) of D-threo-(1,2- ^{14}C)-chloramphenicol (NEN, Boston, MA) in a final volume of 150 μl of 0.25 M Tris, pH 7.8. Reactions were incubated for 3 h at 37°C, extracted with ethyl acetate, dried, and spotted on silica gel plates (Polygram Sil G; Macherey-Nagel, Duren, Germany). Plates were run in a tank containing a mixture of chloroform/methanol (95:5). After a 16-h autoradiography, the plates were cut and samples were counted in a scintillation counter (LS5000TD; Beckman Instruments, Inc., Palo Alto, CA).

Electrophoretic Mobility Shift Assays (EMSA). Nuclear extracts and gel shift assays were performed as described elsewhere (27, 29). After incubation, cells were harvested, washed once in cold PBS, and transferred to 1.7-ml microfuge tubes for a second wash in cold PBS. The supernatant was removed and the cell pellet was resuspended in lysing buffer (10 mM Hepes, pH 7.9, 1 mM EDTA, 60 mM KCl, 1 mM dithiothreitol (DTT), 1 mM PMSF, and 0.2% vol/vol NP-40 for 5 min. Nuclei were collected by centrifugation (500 g for 5 min), rinsed with NP-40-free lysing buffer, and re-

suspended in 150 μl of buffer containing 250 mM Tris HCl, pH 7.8, 20% glycerol, 0.42 M NaCl, 60 mM KCl, 1 mM DTT, and 1 mM PMSF. Nuclei were then subjected to three cycles of freezing and thawing. The suspension was cleared by centrifugation (7,000 g for 15 min), and aliquots were immediately tested in a gel retardation assay or stored in liquid phase N_2 until use. Oligonucleotide probes used included 5'-GATCGGACGTCACATTGCACAATC-TTAATAAT-3' (NF-IL-6), 5'-GGACGTCACACTACAACTCT-TAATAA-3' (mutant NF-IL6), 5'-TGGGATTTCCCA-3' (κB), and 5'-TAATATTTTCCCA-3' (mutant κB). Each oligonucleotide was annealed to its complementary strand and end-labeled with γ - ^{32}P -ATP (Amersham Corp., Arlington Heights, IL) using polynucleotide kinase (New England Biolabs, Beverly, MA). Equal amounts of cell extracts were incubated in a reaction mixture consisting of 20 μl buffer containing 20 mM Hepes, pH 7.9, 20% glycerol, 100 mM KCl, 0.2 mM EDTA, 0.5 mM DTT, 0.5 mM PMSF, 2 μg poly (d[I-C]) (Boehringer Mannheim), and 5 μg of extract for 5 min on ice. 1 μl of γ - ^{32}P -labeled double-stranded probe (0.2 ng, $4\text{--}6 \times 10^4$ cpm) was then added with or without a 100-fold excess of competitor wild type or mutant oligonucleotide. The reactions were incubated at room temperature for 30 min and run on a 5% acrylamide/bis-acrylamide (30:1) gel in 22.5 mM Tris-borate, 0.5 mM EDTA. Gels were dried and autoradiographed.

Results and Discussion

The HIV1 TAT Protein Transactivates the Promoter of the IL-6 Gene. The pIL6Pr-CAT was cotransfected with either the pSVT8 or the pSVT10 plasmid (22) that carry the *tat* gene in sense and antisense orientation, respectively, and allow G-418 selection by expressing the *neo* gene. As shown in Fig. 1, *a* and *b*, the *tat* gene product efficiently increased the IL-6 promoter-dependent CAT activity in both B lymphoblastoid MC3 and epithelial HeLa cells. In these experiments LPS, a likely contaminant of plasmid preparations and a possible inducer of IL-6 promoter activity (30), exerted no effect. These data suggested that the *tat* gene, transiently expressed in both B lymphoid and epithelial cells, transactivated the promoter of IL-6.

In further experiments, MC3 and HeLa cells were stably transfected with either the pSVT8 or the pSVT10 plasmid. After selection in G-418, we obtained stable bulk cultures of MC3-pSVT8, MC3-pSVT10, HeLa-pSVT8, and HeLa-pSVT10 cells. The integration of the *tat* gene was tested by Southern blots of genomic DNA, and the *tat* transcription was analyzed by PCR amplification of *tat*-specific transcripts. To ascertain the presence of TAT proteins, we exploited the transactivating effects that TAT exerts on HIV1-LTR (11), and conducted transient expression experiments in which pSVT8- or pSVT10-positive cells were transfected with the pILC-CAT, an HIV1-LTR-carrying plasmid (26). These experiments showed that the transfected MC3 and HeLa cells (*a*) harbored the *tat* gene in a random genomic integration with an average of one copy per cell genome (data not shown); (*b*) efficiently expressed *tat*-mRNAs (data not shown); and (*c*) had a high constitutive activation of the HIV1-LTR-CAT plasmid, which indicated the endogenous production of functional TAT proteins (Fig. 2 *a*). We used transient expression experiments to test whether pSVT8- or pSVT10-transfected

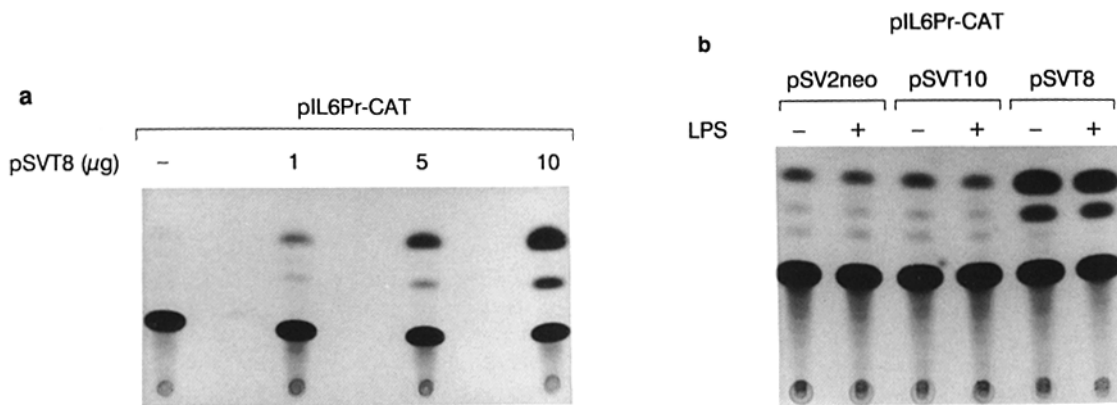


Figure 1. The *tat* gene product transactivates the IL6 promoter. (a) HeLa cells were transiently transfected with 10 μg of pIL6Pr-CAT plasmid together with the indicated amounts of the *tat*-expressing pSVT8 plasmid. The percent specific acetylations, calculated by directly counting the acetylated spots, were: 0.3 (baseline), 2.2 (1 μg), 6.8 (5 μg), and 16.8 (10 μg). In parallel experiments, cells were transfected with 10 μg of pIL6Pr-CAT plasmid and with similar amounts of pSVT10 (anti-*tat*) plasmid with no significant variations over the baseline pIL6Pr-CAT expression (data not shown). The data are representative of several independent experiments. Similar results were obtained by transfecting MC3 cells (data not shown). (b) MC3 B-lymphoblastoid cells were transfected with 10 μg of pIL6Pr-CAT and with 10 μg of the indicated plasmids. At 24 h after transfection, cell aliquots were stimulated with *Escherichia coli* LPS (1 μg/ml) for additional 24 h. Percent acetylations were: pSV2neo, 0.80 (LPS-), 0.75 (LPS+); pSVT10, 1.10 (LPS-), 1.15 (LPS+); pSVT8, 10.5 (LPS-), 10.8 (LPS+). The data are representative of several independent experiments. Similar results were obtained by transfecting HeLa cells (data not shown).

cells transactivated the pIL6Pr-CAT plasmid. Both MC3-pSVT8 and HeLa-pSVT8 produced a higher IL-6 promoter-driven CAT activity than did the control pSVT10-transfected cells (Fig. 2 b).

When the *tat*-transfected MC3 and HeLa cells were tested for the secretion of IL-6 by specific ELISA assay, pSVT8-transfected cells secreted a higher amount of IL-6 than the pSV2neo- or pSVT10-transfected cells (Table 1). These data indicate that stable expression of the HIV1 *tat* gene results in the activation of the endogenous IL-6 gene.

Identification of the Region of the IL-6 Promoter Responsive to TAT. To gain further insights into the molecular mechanisms of the TAT-induced activation of the IL-6 gene, we performed a primer extension analysis of *cat* mRNA in pSVT10- and pSVT8-cells transfected with the pIL6pr-CAT

plasmid. As shown in Fig. 3, HeLa-pSVT8 cells expressed higher amount of *cat* mRNA than the anti-*tat*-transfected pSVT10-cells. The elongated product extended for 98 bp, locating the transcription start site at -63 bp, corresponding to the major start site of the IL-6 gene (24). Similar results were obtained by primer extension analysis of the IL-6 mRNA transcribed from the endogenous IL-6 gene in *tat*- or anti-*tat*-transfected MC3 and HeLa cells (data not shown).

To address the question of whether TAT could interact with IL-6 promoter sequences located upstream to the transcription start site, we constructed 5' deletion mutants of pIL6Pr-CAT in which regions at -658, -287, and -172 were inserted 5' to the *cat* gene (Fig. 4 a). These plasmids were transiently transfected in pSVT10 and pSVT8 cells and showed that the pIL6Pr(-287)-CAT plasmid was efficiently transac-

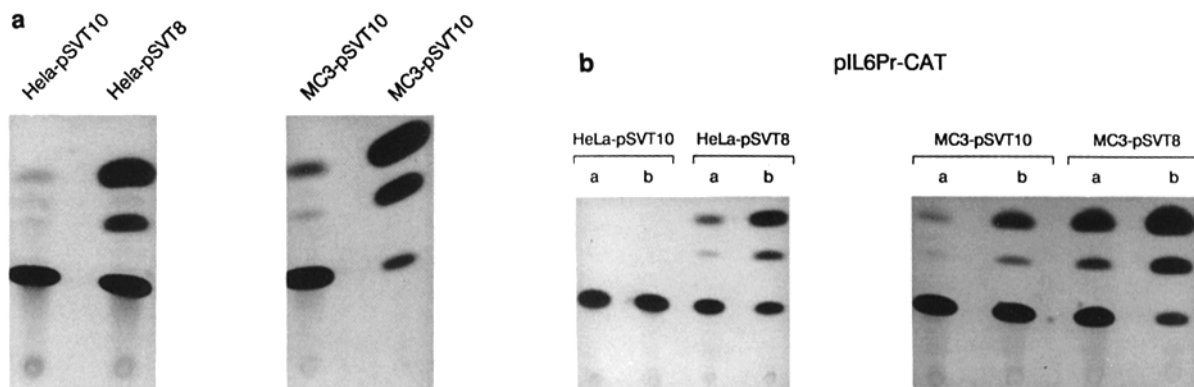


Figure 2. Constitutive activation of the IL-6 promoter in HeLa and MC3 cells stably expressing the *tat* gene. (a) Constitutive expression of pIL6Pr-CAT in the *tat*-transfected cells. *Tat* (pSVT8)- or anti-*tat* (pSVT10)-expressing cells were transiently transfected with 10 μg of the HIV-1-LTR-carrying pIL6Pr-CAT plasmid. Percent acetylations were: HeLa-pSVT10, 0.6; HeLa-pSVT8, 38.0; MC3-pSVT10, 1.8; MC3-pSVT8, 92.0. (b) Expression of pIL6Pr-CAT plasmid in *tat*-transfected cells. Cells were transfected with 5 μg (a) or 10 μg (b) of pIL6Pr-CAT plasmid. Percent acetylations, calculated 48 h after transfection, were: HeLa-pSVT10: (a) 0.10; (b) 0.15; HeLa-pSVT8: (a) 5.3; (b) 14.0; MC3-pSVT10: (a) 1.0; (b) 2.4; MC3-pSVT8: (a) 12.5; (b) 23.0.

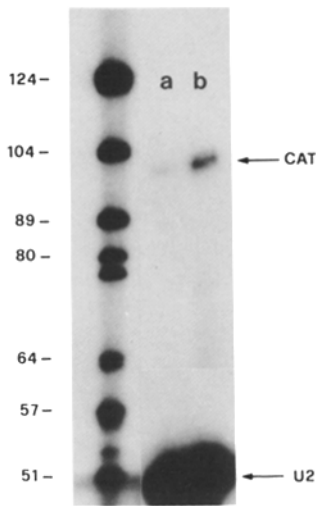


Figure 3. Primer extension analysis of pIL6Pr-CAT mRNA in pSVT10- or pSVT8-transfected cells. Anti-*tat* (pSVT10)- or *tat* (pSVT8)-expressing HeLa cells were transfected with 10 μ g of pIL6Pr-CAT plasmid. 36 h later, total RNA was isolated and analyzed by primer extension. Protected bands of *cat* and U2 mRNAs are indicated. A 24 h exposure of U2 band is shown, while *cat* mRNA band is a 4-d exposure. (a) HeLa-pSVT10; (b) HeLa-pSVT8.

tivated by TAT, whereas the pIL6Pr(-172)-CAT construct was unresponsive to TAT (Fig. 4 *b*). This suggests that TAT-induced activation of the IL-6 promoter required a minimal region located between -287 and -54 bp (at the XhoI site). Next, we generated plasmids where the -658, -287, and -172 bp regions of the IL-6 promoter were inserted downstream to a TAR-derived HIV1-LTR sequence. The resulting p Δ ILC-IL6Pr-CAT plasmids (Fig. 4 *a*) were transiently expressed in *tat* or anti-*tat*-transfected cells. In these experiments, the sequence of -172 to -54 of the IL-6 promoter was unresponsive to TAT (pIL6Pr(-172)-CAT, Fig. 4 *b*). This sequence, however, conferred TAT responsiveness to the TAR-deleted HIV1-LTR promoter (compare p Δ ILC-CAT and p Δ ILC-IL6Pr(-172)-CAT plasmids in Fig. 4 *b*).

A primer extension analysis of *cat* mRNA transcribed by the p Δ ILC-IL6Pr(-172)-CAT identified a major protected

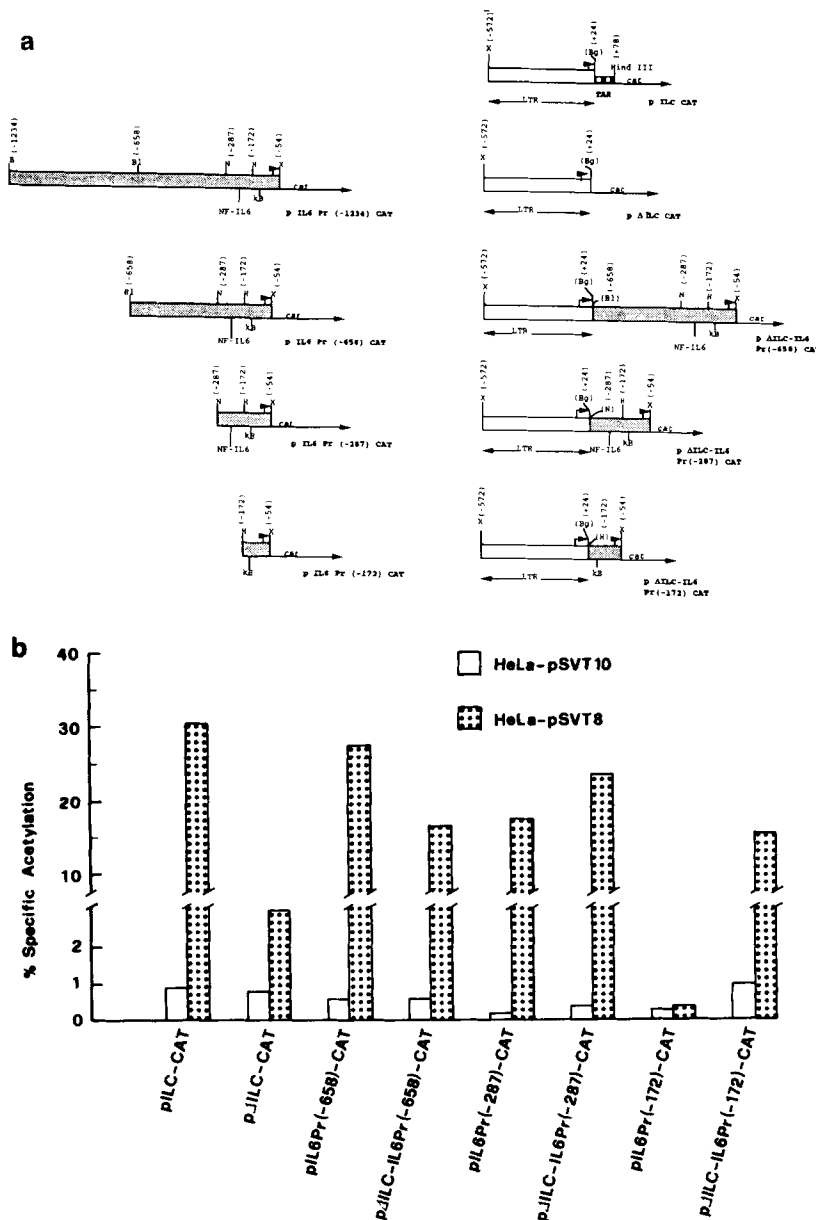


Figure 4. (a) Schematic representation of pIL6Pr-CAT and HIV1-LTR-IL6 promoter fusion plasmids. Restriction sites are indicated as: (B) BamHI; (B)l BglII; (N) NheI; (H) HaeIII; (X) XhoI. (b) Induction of mutant pIL6Pr-CAT and of HIV1-LTR-pIL6Pr-CAT fusion plasmids by TAT. 10 μ g of the indicated plasmids, carrying 5' deletions of pIL6Pr-CAT or discrete regions of the IL-6 promoter fused to a TAR-deleted HIV1-LTR sequence, were transiently transfected in pSVT10- or pSVT8-HeLa cells. CAT activity was assayed at 48 h as detailed in Materials and Methods.

Table 1. Secretion of IL-6 Molecules by MC3, HeLa, and 7TD1 Cells

Cells	IL-6 secretion (pg/ml)
MC3-pSV2neo	14.0
MC3-pSVT10	16.6
MC3-pSVT8	318
HeLa-pSV2neo	336
HeLa-pSVT10	384
HeLa-pSVT8	1,913
7TD1-pSV2neo	<3
7TD1-pSVT8	216
M ϕ + LPS	2,350

pSV2neo-transfected control cells and *tat*- or anti-*tat*-transfected cells were cultured for 48 h. Supernatants were then collected and tested for IL-6 content by ELISA specific for human or murine IL-6. Data express the amounts of IL-6 secreted by 10^6 cells over 48-h culture time. Human monocytes were isolated by centrifugation over Ficoll-Hypaque followed by 46% Percoll gradients and stimulated with LPS at 1 μ g/ml, as described elsewhere (45).

band of 248 nucleotides (nt), corresponding to the start site (+1) of HIV1-LTR (Fig. 5, lanes a and c). Moreover, consistent with the data shown in Fig. 3, the amount of *cat* mRNA in *tat*-expressing cells was significantly higher than the *cat* mRNA transcribed by anti-*tat*-transfected cells. In fact, both the 98-nucleotide *cat* band generated by transfecting pIL6Pr(-287)-CAT (Fig. 5, lanes b and d), and the 248 nt *cat* band generated by the p Δ ILC-IL6Pr(-172)-CAT (Fig. 5, lanes a and c) were highly activated in TAT-positive cells. A longer exposure of the primer extension experiments revealed the presence of a 98-nt additional *cat* band in cells transfected with the p Δ ILC-IL6Pr(-172)-CAT, indicating that the start sites of the IL-6 promoter and of the HIV1-LTR were both functional. A densitometer analysis of the *cat* bands revealed that the HIV1-LTR start site was preferentially utilized, with a minimal transcription from the IL-6 promoter start site (data not shown). This suggests that both the pIL6Pr(-287)-CAT and the p Δ ILC-IL6Pr(-172)-CAT plasmids were responsive to TAT. Results shown in Fig. 4 b identified the sequence of -172 to -54 (at the XhoI site) as the minimal region of the IL-6 promoter required for TAT to transactivate the Δ TAR HIV1-LTR. This suggests that the -172/-54 bp region could function as a TAT-responsive sequence, possibly allowing TAT to be directed close to the TATA box of the IL-6 promoter.

The Expression of the HIV1 tat Gene Induces an Increase in NF- κ B and NF-IL6 Binding Factors. The data shown in Fig. 4 b, where TAT activated the pIL6Pr(-287)-CAT plasmid while the pIL6Pr(-172)-CAT was unresponsive to TAT, suggested that TAT-mediated activation of the IL-6 gene required transcription factors binding sequences located within -287 and -54 bp of the IL-6 promoter. In this region, functional

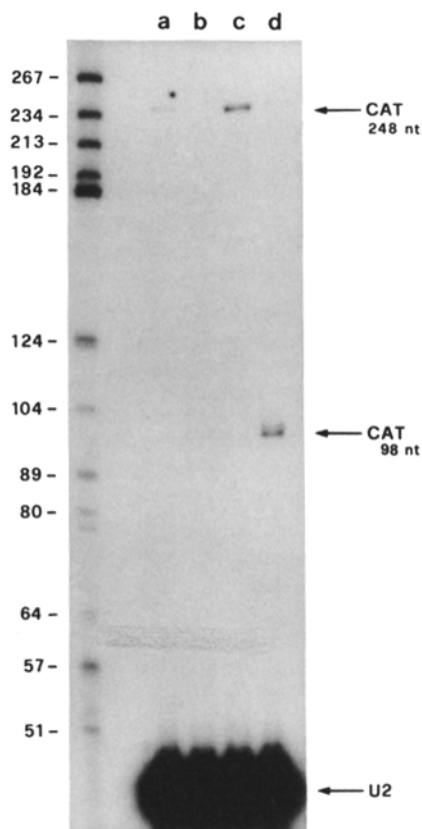


Figure 5. Primer extension analysis of *cat* mRNA in pSVT10- or pSVT8-cells transfected with pIL6Pr(-287)-CAT and of p Δ ILC-IL6Pr(-172)-CAT plasmids. Anti-*tat* (pSVT10)- or *tat* (pSVT8)-expressing HeLa cells were transfected with 10 μ g of pIL6Pr(-287)-CAT or p Δ ILC-IL6Pr(-172)-CAT. 36 h later, total RNA was isolated and analyzed by primer extension. The 98-nt and the 248-nt *cat* messages were generated by transfecting the pIL6Pr(-287)-CAT or the p Δ ILC-IL6Pr(-172)-CAT, respectively. HeLa pSVT10 cells transfected with p Δ ILC-IL6Pr(-172)-CAT (a) or with pIL6Pr(-287)-CAT (b); HeLa-pSVT8 cells transfected with p Δ ILC-IL6Pr(-172)-CAT (c) or with pIL6Pr(-287)-CAT (d). A 24-h exposure is shown. Similar results were obtained in the case of transfected MC3 lymphoblastoid cells.

NF- κ B and NF-IL6 enhancers have been identified (20, 21). Therefore, we tested whether TAT might induce nuclear factors binding to the κ B or NF-IL6 sequences of the IL-6 promoter. As shown in Fig. 6, *tat*-transfected cells expressed a consistent increase in κ B and NF-IL6 binding activity. Accordingly, the IL-1REK9CAT and pDRCAT, carrying tandem repeats of NF-IL6 or NF- κ B binding motifs, were consistently activated in TAT-expressing cells (Table 2). In other experiments, we also found that suboptimal amounts of the *tat*-expressing pSVT8 plasmid synergized with the pCMV-NFIL6 expression vector (21) in inducing a full activation of the IL-6 promoter (data not shown).

The Expression of the HIV1 tat Gene in the IL-6-dependent 7TD1 cells Confers Growth Independency from IL-6 and Enhanced Tumorigenicity. Deregulated expression of the IL-6 gene has been associated with the abnormal growth and tumorigenic phenotype of a variety of lymphoid and mesenchymal cells,

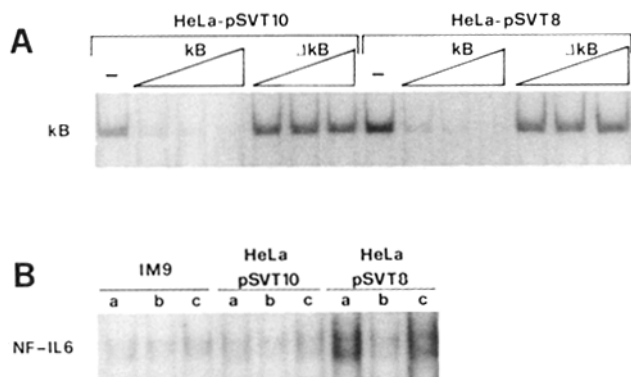


Figure 6. Induction of NF- κ B and NF-IL6 binding factors by TAT. 10 μ g of nuclear proteins were tested for binding to the κ B or NF-IL6 sequences of the IL-6 promoter. IM9 is a myeloma cell line constitutively expressing NF-IL6. Only retarded bands are shown. Competitions were tested with 25-, 50-, or 100-fold molar concentrations of unlabeled κ B or mutant κ B (Δ - κ B). In the NF-IL6 band-shift experiments, nuclear extracts were competed with 50-fold molar concentrations of unlabeled NF-IL6 (b) or mutant NF-IL6 (c). Similar results were obtained with MC3 cells.

including EBV-positive B cell lymphoma and Kaposi's sarcoma (4–7) which are common malignancies among HIV1-infected patients. In vitro, IL-6 molecules enhance the survival and the growth potential of neoplastic B cells and Kaposi's sarcoma cells and they induce a tumorigenic phenotype in immortalized EBV-positive cells (5, 7). To gain further insight into the biological relevance of the TAT-mediated induction of IL-6, we transfected the IL-6-dependent 7TD1 cells (23) with pSVT8, pSVT10, or pSV2neo control plasmids. These cells, which are strictly dependent on exogenous IL-6 for their growth, were cultured in the presence of optimal amounts (20 ng/ml) of human rIL-6 and G-418 for 15 d.

Isogenic 7TD1-pSVT98 (TAT positive) and 7TD1-pSVT10 (TAT negative) cells were then cultured without exogenous IL-6. Only the *tat*-expressing 7TD1-pSVT8 cells survived and grew efficiently without exogenous IL-6 (shown in Fig. 7 a). pSVT8- and pSVT10-transfected cells expressed *tat*-specific mRNA (Fig. 7 b). Moreover, *tat*-transfected cells showed a constitutive activation of pILC-CAT plasmid higher than the anti-*tat*-transfected cells, indicating that functional TAT proteins were produced by the *tat*-expressing cells (Fig. 7 c). These *tat*-positive cells expressed IL-6 mRNAs, whereas control 7TD1 cells did not express any detectable IL-6 mRNAs (Fig. 7 d). Moreover, 7TD1-pSVT8 cells constitutively secreted consistent amounts of murine IL-6 (Table 1).

The IL-6-dependent 7TD1-pSV2neo and 7TD1-pSVT10 cells could grow both in the presence of conditioned medium of 7TD1-pSVT8 cells and when they were cocultured with TAT-positive 7TD1-pSVT8 cells in transwell culture plates where cell contacts were prevented (data not shown). These effects were probably caused by the IL-6 secreted by *tat*-transfected cells, rather than by TAT proteins released by pSVT8-transfected cells. In fact, a polyclonal antibody to TAT protein did not inhibit the IL-6-independent growth of TAT-positive 7TD1-pSVT8 cells, whereas antibodies to mouse IL-6 consistently decreased the growth of 7TD1-pSVT8 cells (data not shown). Furthermore, recombinant TAT was unable to substitute for IL-6 in the growth of 7TD1 parental cells (data not shown). These data argue against a direct role of TAT as growth factor.

The expression of the *tat* gene and the subsequent IL-6-independent growth of 7TD1-pSVT8 cells would be expected to induce a modulation of the tumorigenic phenotype of the TAT-positive 7TD1 cells. This possibility was tested by subcutaneously injecting either the control 7TD1-pSV2neo or the TAT-positive 7TD1-pSVT8 cells in athymic nude mice.

Table 2. Constitutive Expression of pDRCAT and pIL-1REK9CAT in *tat*- or Anti-*tat*-transfected HeLa Cells

Cells	Transfected plasmid			CAT activity [‡]	
	pDRCAT	pIL-1REK9CAT	pILC-CAT	Acetylation	Fold induction [§]
					%
HeLa pSVT10*	+	–	–	3.2	
HeLa pSVT8	+	–	–	55.4	17.3
HeLa pSVT10	–	+		0.3	
HeLa pSVT8	–	+		5.7	19.0
HeLa pSVT10	–	–	+	2.8	
HeLa pSVT8	–	–	+	62.0	22.1

The data are representative of five independent experiments. Similar results were obtained in the case of MC3-pSVT10 and MC3-pSVT8 cells. * HeLa-pSVT10 (anti-*tat*-transfected) or HeLa-pSVT8 (*tat*-transfected) were transiently transfected with 10 μ g of the pDR CAT (carrying two κ B motifs) or with 10 μ g of the pIL-1REK9-CAT (carrying three NF-IL6 binding motifs). Cells were also transfected with 10 μ g of pILC-CAT, a HIV1-LTR-CAT construct.

[‡] Determined at 48 h after transfection by using 50 μ g of cell extract.

[§] Expressed as the ratio of percentages acetylated.

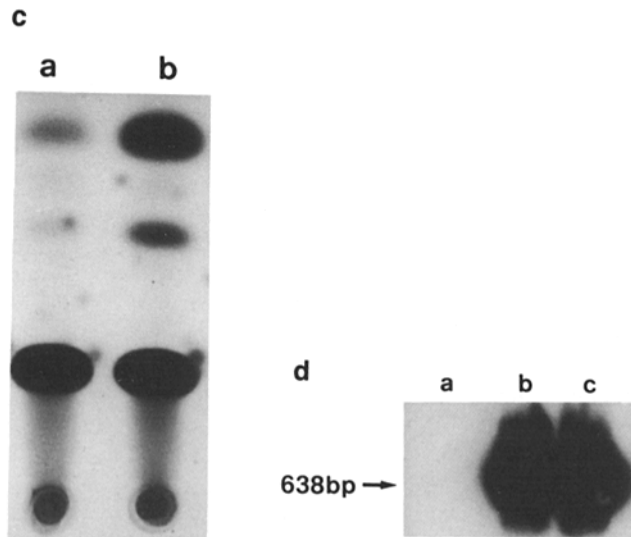
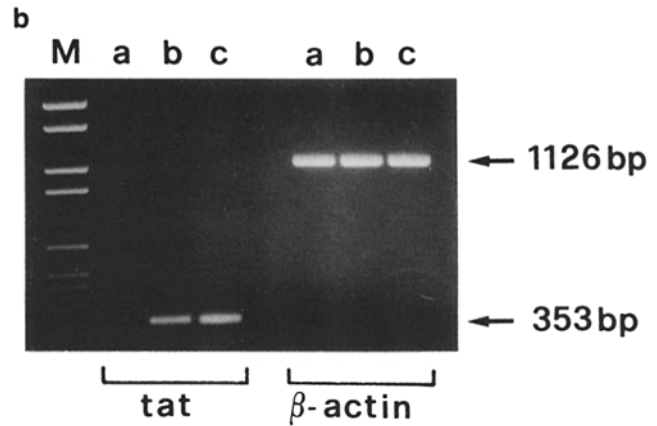
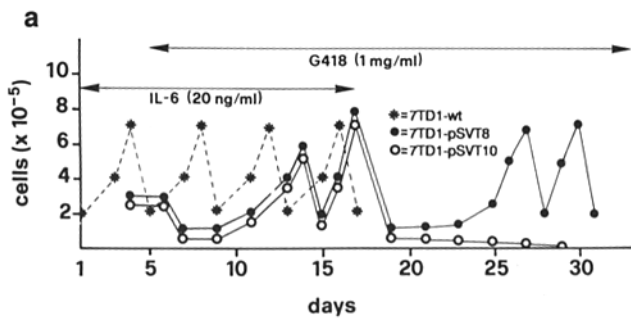


Figure 7. Generation and characterization of 7TD1 cells stably expressing the *tat* gene. (a) 7TD1 cells were transfected with pSVT8 or pSVT10 plasmid as previously reported (6, 27). G-418 selection (2 mg/ml, Geneticin) was applied 48 h after transfection. 2 wk later, cells were cultured without exogenous IL-6 whereas G-418 selection was continued. Cells were monitored daily for growth. Aliquots of 7TD1 cells were transfected with pSV2neo plasmid and cultured in IL6-supplemented medium (data not shown). (b) Expression of the transfected *tat* gene. Total RNA from pSV2neo, pSVT8-, or pSVT10-transfected 7TD1 cells was amplified with reverse PCR using *tat*- or β -actin-specific primers and visualized by electrophoresis in 1.5% agarose. (M) BglI-HinI digested pBR328; (a) RNA from 7TD1-pSV2neo cells; (b and c) RNA from pSVT8- or pSVT10-transfected cells. (c) Constitutive expression of the pILC-CAT plasmid in 7TD1 and 7TD1-pSVT8 cells. 7TD1-pSV2neo (a) and 7TD1-pSVT8 (*tat*-positive cells, b) were transfected with 10 μ g of the HIV1-LTR-carrying pILC-CAT plasmid. Percent acetylations 48 h after transfection were: (a) 1.5; (b) 30.4. (d) Expression of the endogenous IL-6 gene in *tat*-transfected 7TD1 cells. RNA from 7TD1-pSV2neo (a), 7TD1-pSVT8 (b), and mouse spleen cells stimulated for 24 h with 1 μ g/ml of LPS (c) was amplified by reverse PCR, electrophoresed in agarose gel, transferred onto nylon membrane, and hybridized with a mouse IL-6 cDNA probe.

Tat-transfected 7TD1-pSVT8 cells caused a dramatic increase in both the number of mice with tumors and in the number of tumors per animal (Table 3). These data suggest that TAT proteins could play a role in the generation of a tumorigenic phenotype by sustaining the autocrine growth of susceptible cells.

In this paper we show that the HIV1 *tat* gene, either transiently expressed or stably transduced in MC3 lymphoblastoid cells and in HeLa epithelial cells, transactivates the human IL-6 promoter. Moreover, the stable expression of the HIV1 *tat* gene resulted in the activation of the endogenous IL-6 gene and in the secretion of consistent amounts of IL-6 molecules. This finding may be of relevance because IL-6 molecules are secreted by HIV1-infected cells and can increase HIV1 gene expression (8, 31), and because a variety of cell types, including B and epithelial cells, can be infected by HIV1 (32, 33). The foregoing data suggest that there is a potent autocrine or paracrine mechanism of stimulation of HIV1 gene expression in which HIV1 infection results in a TAT-mediated secretion of IL-6 molecules, which in turn enhance HIV1 gene expression. Interestingly, IL-6 gene expression can also be triggered by the HIV1 envelope glycoproteins gp120 and

gp160 (34, 35), suggesting that a deregulated production of IL-6 could take place in the early phase of HIV1 infection, and could result in the stimulation of different bystander cell types. A recent paper (36) reported that HIV1 infection resulted

Table 3. *In Vivo* Tumorigenicity of *tat*-transfected 7TD1 Cells

Cells	Tumorigenicity in vivo	
	No. of mice with tumors/ No. of inoculated mice	Average of tumors per positive mouse
7TD1-pSV2neo	1/6	1
7TD1-pSVT8	6/6	3.0

Tumorigenicity in nude mice was assayed by a single subcutaneous injection of 2×10^6 cells in 0.2 ml of PBS (6). Typically, tumors >5 mm in diameter developed after 10–15 d of latency at the challenge site and in the regional LN and showed a histological picture of infiltrating blast cells (data not shown). Data are the numbers of tumors recorded through 6 wk of observation.

in the activation of TNF- β , whose gene product is a potent inducer of IL-6 gene (4), likely through the TAT protein. In the same study, the IL-6 production was not detected, suggesting that TAT may selectively activate specific promoters in different cell phenotypes.

The molecular mechanisms whereby TAT activates the IL-6 gene are at present unknown. The TAT protein of HIV1 is required for efficient viral gene expression (12). TAT increases the initiation of transcription from the HIV1 LTR (13) and affects RNA processing and utilization by interacting with a TAT-responsive element (TAR) located between nucleotide +1 and +44 with respect to the start site (+1) of viral transcription (11). TAT could activate IL-6 transcription by interacting with RNA stem-loop structures in the 5' untranslated region of the IL-6 promoter. At least three functional transcription start sites have been identified in the IL-6 promoter at -176, -86, and -63 from the first nucleotide of the ATG codon (24). A sequence analysis of these regions, carried out using the energy minimizing algorithm of Zuker (37), defines a RNA stem-loop structure at the 5' untranslated region of IL-6 mRNA potentially able to bind to TAT. Here we report that the region of the IL-6 promoter located at -172 to -54 can substitute for the TAR sequence, allowing TAT to transactivate the TAR-deleted HIV1-LTR. Our data suggest that TAT may be directed at this region of the IL-6 promoter by binding to a RNA structure, and could induce the transcription of the IL-6 gene by possibly cooperating with transcriptional factors such as NF- κ B and NF-IL6. This possibility is currently under investigation. Alternatively, TAT may bind to cell transcription factors, as recently suggested (15, 17), and ultimately activate IL-6 transcription. Another possibility is that TAT could increase the stability of a baseline constitutive transcription of IL-6 gene

in MC3 and HeLa cells. However, such posttranscriptional activity cannot apply to 7TD1 cells because they do not express any detectable IL-6 mRNA.

Recent reports suggest that TAT cooperates with upstream regulatory DNA sequences circumscribed within the NF- κ B/Sp1 region of the HIV1 promoter (14) and with host cell proteins (15-19). TAT could affect the transcription of cell genes such as IL-6, whose regulatory sequences, like HIV1-LTR, possess NF- κ B and NF-IL6 enhancer elements (20, 21). In support of this possibility, we found that expression of *tat* leads to an increase in κ B and NF-IL6 binding factors. In addition, since the pIL6Pr(-172)-CAT plasmid, which lacks a NF-IL6 binding motif but retains a κ B site, was not induced by TAT, our data suggest that in *tat*-transfected cells NF- κ B and NF-IL6 could function as a unique transcriptional complex, as recently suggested (38), in activating the IL-6 promoter.

A variety of clinical lesions are associated with HIV1 infection. This includes B lymphomas, psoriasis, and Kaposi's sarcomas, where HIV1 transcripts have been detected (1-3, 39). There is evidence that abnormal secretion of IL-6 plays a major role in the pathogenesis of these diseases by cooperating with such other cytokines as oncostatin M and IL-1, which in turn are potent inducers of IL-6 secretion (4, 40-42). In this paper we showed that TAT is a major stimulant of IL-6 secretion and could induce autocrine growth and enhanced tumorigenicity of the IL-6-dependent 7TD1 cells. These data are consistent with and support the recent observations that TAT can activate the transcription of genes from heterologous viral and cellular promoters (36, 43, 44), and suggest that TAT proteins may directly participate in the pathogenesis of HIV1-associated diseases by modulating the expression of host cellular genes.

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