

Original investigation

HIV-1 Proteins Influence Novelty-Seeking Behavior and Alter Region-Specific Transcriptional Responses to Chronic Nicotine Treatment in HIV-1Tg Rats

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Abstract

Introduction: Clinical studies suggest that HIV-1-infected patients are more likely to use or abuse addictive drugs than is the general population. We hypothesized that HIV-1 proteins impact novelty-seeking behavior and enhance the transcriptional response to nicotine in genes implicated in both novelty-seeking behavior and drug addiction.

Methods: We assessed the effects of HIV-1 proteins on novelty-seeking behavior by comparing baseline activity differences of HIV-1Tg and F344 control rats in the open-field test. One day after behavioral testing, all rats began daily subcutaneous injections of either nicotine (0.4 mg/kg, base) or saline (the same for each rat) for 27 days. At the end of treatment, the prefrontal cortex, nucleus accumbens, and ventral tegmental area were collected for RNA expression analysis of genes in the receptor families for dopamine, GABA, glutamate, and serotonin.

Results: Significant strain difference was detected in the distance moved in the center, such that HIV-1Tg rats traveled greater distance in the center of the arena than did F344 rats. Quantitative RT-PCR analysis showed that mRNA from *Drd3* and *Grm2* in the prefrontal cortex and *Drd5* and *Gabra6* in the ventral tegmental area was significantly upregulated, whereas that of *Drd5* in the nucleus accumbens was downregulated in HIV-1Tg rats compared with F344 rats. Further, more addiction-related genes were significantly modulated by nicotine in each brain region in the HIV-1Tg rats than in the control animals.

Conclusions: HIV-1 proteins may affect novelty-seeking behavior and modulate the expression of genes related to drug addiction and novelty-seeking behavior.

Implications: HIV-1 viral proteins and chronic nicotine treatment impact the expression of genes involved in novelty-seeking behavior and addiction in three brain regions of the HIV-1 transgenic rat. These findings implicate that HIV-1 proteins may be involved in novelty-seeking behavior and in modulating the expression of genes related to drug addiction and novelty seeking.

Introduction

According to the United Nations Program on HIV/AIDs, approximately 16 million persons utilize drugs of abuse; among them, about 3 million worldwide are living with HIV-1 infection.¹ It has been hypothesized that HIV-1 infection alters the structure and function of the central nervous system (CNS) reward pathways, as the infection is associated with drugs of abuse at both the cellular and molec-ular levels.^{[2](#page-6-1),[3](#page-6-2)} It is speculated that these alterations in the CNS make such individuals more vulnerable to the rewarding effects of drugs of abuse.[4–8](#page-7-0) Although this hypothesis has been supported by clinical evidence showing that the percentage of HIV-infected patients who use various addictive substances such as alcohol, nicotine, morphine/heroin, methamphetamine, and cocaine is greater than that in the general population,^{[6](#page-7-1)} the molecular mechanisms remain largely unknown.

HIV-1 infection involves the actions of viral proteins on targeted cells of the immune system, such as macrophages and T lymphocytes. $9-11$ Like the peripheral immune system, the CNS is highly vulnerable to HIV-1 infection. The viral proteins penetrate the CNS during the early phases of the infection and selectively target dopamine-rich regions, particularly the mesocortical and mesoaccumbens dopamine circuits.[12–14](#page-7-3) Further, the HIV-1 proteins Tat and gp120 are toxic to dopaminergic neurons, causing dopamine depletion in subcortical structures^{15,16} and contributing to the development of neurologic complications of HIV infection.[5,](#page-7-6)[17](#page-7-7)[,18](#page-7-8) These protein-induced alterations could contribute to a higher risk of substance abuse development because of the increased sensitivity of the brain to the pleasurable effects of psychostimulants[.17](#page-7-7) Thus, it is reasonable to assume that HIV-infected patients are more prone to escalate their drug-intake behavior as a result of viral protein-induced changes in dopamine-rich brain regions. The prefrontal cortex–ventral tegmental area–nucleus accumbens (PFC-VTA-NAc) neural circuits are critical for elicitation of reward perception, goal-directed behavior, and habit formation^{19,[20](#page-7-10)} and are targeted by HIV-1 proteins.^{[21](#page-7-11)[,22](#page-7-12)} However, it is less clear how viral protein-induced changes in the PFC–VTA–NAc circuitry of neurotransmitter systems mediate the neural and behavioral responses to drugs of abuse.

In addition to molecular changes, persons with long-standing HIV-1 infection demonstrate a greater change in risk-taking and safety assessment behaviors.²³ It has been proposed that an alteration in this equilibrium renders HIV-infected patients more vulnerable to drugs of abuse. Nicotine, in the form of cigarette smoking, is the preferred drug of abuse among HIV-infected patients.²⁴ Clinical studies also show that smokers with HIV-1 infection engage in other risky behaviors such as unprotected sex and have higher rates of abuse of alcohol and club drugs^{[25,](#page-7-15)[26](#page-7-16)} despite their knowledge of the long-term negative consequences of these actions.

Studies of humans and of animal models of nicotine addiction reveal that novelty seeking as a component of risk behavior can predict the likelihood of compulsive use of cigarettes. Individuals who display high novelty-seeking behavior consume more cigarettes than do low novelty seekers.²⁷⁻²⁹ Previous experiments with mice have shown that novelty-seeking behavior correlates with consistent nicotine use.[30](#page-7-18)[,31](#page-7-19) Both nicotine effects and novelty-induced reward feedback can be localized to the mesolimbic circuitry with a prominent effect on dopamine.[32](#page-7-20)[,33](#page-7-21) Molecular mechanistic studies indicate that both novel stimuli and psychostimulant drugs, such as nicotine, affect behavior by producing a strong neuromodulator response through activating the cholinergic,³⁴⁻³⁷ GABA, glutamatergic, and serotogenergic systems.³⁸ However, there is limited knowledge of the effects of HIV-1 proteins on the molecular mechanisms underlying long-term changes in novelty-seeking behavior and response to nicotine.

In this study, we used both behavioral and molecular approaches to determine whether long-term HIV infection modifies high-risk behaviors and alters the molecular response to nicotine. The HIV-1Tg rat model used carries a *gag-pol*-deleted HIV-1 genome under the control of the HIV-1 viral promoter and expresses seven of the nine HIV-1 genes[.39](#page-7-24) The HIV-1Tg rat displays symptoms similar to those of human HIV patients receiving highly active antiretroviral therapy (HAART) and shows greater sensitivity to many psychostimulants, including morphine, alcohol, and nicotine.⁴⁰⁻⁴⁴ For our behavioral analysis, we measured novelty-seeking behavior using the open-field arena test⁴⁵⁻⁴⁷ to determine the modifying effects of viral proteins on behavior and vulnerability to drug abuse. Subsequently, we examined the expression of genes involved in dopaminergic, GABAergic, and glutamatergic neurotransmitter systems and related to nicotine- and novelty-induced reward behaviors. Such molecular studies were conducted in order to determine how gene expression is altered by viral proteins and chronic nicotine treatment.

Materials and Methods

Animals

Male HIV-1Tg rats and F344 genetic-background control rats (Harlan Industries, Cortland, NY) were used at 7–8 weeks of age. All rats were group-housed in standard plastic rat cages, maintained in a temperature $(20^{\circ}C - 22^{\circ}C)$ - and humidity $(45\% - 55\%)$ -controlled environment with a 12-hour light/dark cycle. Food and water were provided *ad libitum*. All experimental procedures were conducted during the light cycle and approved by the University of Virginia Animal Care and Use Committee.

Drug, Treatment and Behavioral Testing

(−)-Nicotine hydrogen tartrate (Sigma, St Louis, MO) was dissolved in 0.9% saline, and its concentration was calculated as nicotine free base. Rats from each strain were divided randomly into two groups: saline-treated control and nicotine-treated experimental, designated as follows: F344_Saline (*n* = 11); F344_Nicotine (*n* = 12); HIV-1Tg_ Saline (*n* = 9); and HIV-1Tg_Nicotine (*n* = 11). To determine whether HIV-1 proteins or chronic nicotine treatment modified the function of the central reward system and related behaviors, baseline noveltyseeking behavior was investigated at 8 weeks of age. The open-field test, a widely used method to measure novelty-seeking behavior, was adopted.[45](#page-7-26) Briefly, novelty-seeking behavior was assessed by recording a unit of activity each time a rat moved from the center of the box to the sides or from the sides to the center by using the Any-Maze video tracking system for 10 minutes (Stoelting Co., Wood Dale, IL). The data on time mobile (total amount of time that the animal was mobile during testing), total distance traveled, and distance traveled in the center and peripheral zones of the open-field arena were automatically tracked by Any-Maze software and analyzed as dependent variables. The test apparatus consisted of a clear plastic box (50 \times 50 \times 50 cm) raised 28 cm above the ground.⁴⁵⁻⁴⁷ Each rat was placed at the center of the box and allowed to move freely through the apparatus without reinforcements or inhibitions.

One day after the open-field test, rats were injected subcutaneously with either saline or nicotine once per day (0.4 mg/kg/day) for 27 days before brain tissues were collected. The duration of nicotine exposure was chosen according to our recently reported findings^{[48](#page-7-27)[,44](#page-7-28)} demonstrating that this amount of time can alter mRNA concentrations in the brains of HIV-1Tg rats. This choice was shown to be appropriate, given that our results demonstrated both behavioral and molecular alterations among nicotine-treated in comparison with saline-treated rats. The concentration of nicotine was selected on the basis of our recently reported results.[44](#page-7-28) Because of the congenital cataracts in HIV-1Tg rats,⁴⁹ all behavioral experiments were conducted under dimmed red light in order to minimize visual differences between strains.

On day 28, the animals were sacrificed, and different brain regions were collected. Using a rat brain matrix (Kent Scientific, Torrington, CT), 1-mm slices were taken from each brain. The slices containing the prefrontal cortex (PFC; second frontal area), ventral tegmental area (VTA), and nucleus accumbens (NAc) were identified according to the rat brain atlas.⁵⁰ These regions were chosen because of their implications for the reward system, which can be activated by both nicotine and novel stimuli. Tissue from the regions of interest was collected using a 1.5-mm brain punch (Stoelting, Wood Dale, IL) and stored at −80°C until use. We did not pool brain tissues from multiple animals.

RNA Extraction and Primer Design

Total RNA was isolated using the Trizol reagent (Invitrogen, Carlsbad, CA) according to the manufacturer's protocol. The purity and quantity of total RNA were measured at optical densities of 260 nm and 280 nm with the NanoDrop 2000c (Thermo Scientific, Waltham, MA). The primers for all 17 genes examined were designed using Primer Express (v. 3.0) software (Applied Biosystems). To avoid amplifying genomic DNA, two primers of each gene of interest were designed to span at least one intron. Each pair of primers and their amplicon sequences were tested using the Basic Local Alignment Search Tool (BLAST; [http://blast.ncbi.nlm.nih.gov/Blast.](http://blast.ncbi.nlm.nih.gov/Blast.cgi) [cgi](http://blast.ncbi.nlm.nih.gov/Blast.cgi)) to ensure the specificity of the primers for the targeted genes. Dissociation curves were generated to check the specificity of the primers before including them in the qRT-PCR array. The primer sequences used are shown in Supplementary Table 1.

Quantitative RT-PCR (qRT-PCR) Array

A custom-designed RT-PCR array was used to measure gene expression as described previously. 44,51–53 Briefly, 2 μg of total RNA was reverse transcribed into first-strand cDNA using Superscript II Reverse Transcriptase. The cDNA mixture was incubated at 25°C for 10 minutes, 42°C for 1.5 hour, and 70°C for 15 minutes and then amplified in a volume of 10 μL containing 5.0 μL of $2 \times$ Power SYBR Green PCR Master Mix (Applied Biosystems) and combined sense and antisense primers (2.5 μL). The final concentration was 20 nM in a 384-well plate using the 7900HT Sequence Detection System (Applied Biosystems). The PCR conditions were as follows: 95°C for 10 minutes, followed by 40 cycles of 95°C for 15 seconds and 60°C for 1 minute. After the last cycle, a dissociation curve was generated to check for non-specific products.

Statistical Analysis

Graphpad prism (v. 6.01) was used for the statistical analyses, and data are shown as mean ± S.E.M. The unpaired two-tailed *t* test was employed to determine group differences in the measurements of novelty-seeking behavior. Expression of each gene of interest was first normalized to the expression of the housekeeping gene glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and then analyzed using a comparative C_t method.^{[54](#page-8-2)} The relative expression of each gene was compared between the groups of HIV-1Tg control versus

F344 control (to determine viral effects), HIV-1Tg nicotine versus HIV-1Tg saline (to determine the pharmacological effects of nicotine in the presence of viral proteins), and F344 nicotine versus F344 saline (to determine the pharmacological effects of nicotine in the absence of viral proteins) rats using the Student *t* test. Significant alteration in mRNA expression was defined as a fold change > 20% with a *p* value < .05 ($N = 4-6$ per group). According to the literature, a difference of less than 20% between two groups is less likely to be biologically significant.^{55,[56](#page-8-4)}

Results

HIV-1Tg Rats Showed Greater Novelty-Seeking Behavior than F344 Control Rats

To investigate the role of HIV-1 proteins in novelty-seeking behavior, HIV-1Tg and F344 rats were tested in the open-field arena. [Figure 1](#page-3-0) shows the strain differences in the measurements of novelty-seeking behavior between the two groups. There was a significant strain difference in the baseline measurements. The HIV-1Tg rats spent more time in the center of the arena (t_{37} = 3.924; p = .0004) and made more entries into the center $(t_{37} = 3.907; p = .0004)$ than did the F344 rats [\(Figure 1A](#page-3-0) and [B](#page-3-0)). No significant differences were observed in the mobility scores between HIV-1Tg and F344 rats ([Figure 1C](#page-3-0)). Ambulation was also measured as total distance traveled in the openfield arena. Results showed that there was a significant effect of strain on distance traveled in the center zone of the arena, such that HIV-1Tg rats traveled greater distance in the center than did F344 rats $(t_{37} = 4.187; p = .0002)$ ([Figure 1E\)](#page-3-0). Total distance traveled and distance moved in the peripheral zone during the 10-minutes of testing in the open-field was not significantly different between HIV-1Tg and F344 rats [\(Figure 1D](#page-3-0) and [F\)](#page-3-0). Taken together, these data demonstrate an increase in novelty-seeking behavior in HIV-1Tg rats compared with control animals, suggesting that viral proteins affect the central reward circuit of the brain, which controls novelty-seeking behavior.

Effect of Nicotine or HIV-1 Proteins on Expression of Genes in the NAc that Are Implicated in Drug Addiction and Novelty-Seeking Behavior

As shown in [Table 1](#page-4-0), our qRT-PCR array analysis revealed that 6 of 17 genes implicated in drug addiction were significantly changed, at $p \le 0.05$, by nicotine or HIV-1 proteins in the NAc of HIV-1Tg or F344 rats. In the F344 rats, only the GABA, receptor α 5 subunit gene (*Gabra5*) showed significant upregulation, 135%, by nicotine $(p = .0001)$. In the HIV-1Tg rats, nicotine significantly upregulated the expression of *Drd5* (132.5%; *p* = .002), *Gabra1* (96%; *p* = .007), *Gabra6* (97.6%; *p* = .007), and metabotropic glutamate receptor 1 (*Grm1*) (38%; $p = .011$) but significantly downregulated the expression of *Gabra2* (−18.7%; *p* = .012) and *Grm5* (−30.1%; *p* = .012). Compared with the effects of nicotine, only the expression of *Drd5* was downregulated by HIV-1 proteins (-55.9%; *p* = .025). Thus, more genes were significantly modulated by nicotine in HIV-1Tg rats than in F344 rats, suggesting that the presence of viral proteins could alter the susceptibility of animals to nicotine addiction.

Effect of Nicotine and HIV-1 Proteins on Expression in the PFC of Genes that Are Implicated in Drug Addiction and Novelty-Seeking Behavior

Of the 17 genes examined in the PFC, we found that the expression of *Drd3* (143%; $p = .011$) and metabotropic glutamate receptor

Figure 1. Difference in baseline measures of novelty-seeking behavior in HIV-1Tg and F344 rats in the open-field arena test (*N* = 9–12 per group). The HIV-1Tg rats spent more time in, exhibited higher numbers of entries into, and traveled greater distance in the center zone of the arena compared with F344 rats. Difference from F344 rats ********p* ≤ .001.

2 (126%; $p = .014$) was significantly upregulated by viral proteins in HIV1-1Tg rats (see [Table 2\)](#page-4-1). By comparing the saline-treated F344 with HIV-1Tg rats, we found that viral proteins significantly increased the expression of *Drd3* (143%; $p = .011$) and *Grm2* $(126.2\%; p=.014).$

In the F344 rats, nicotine significantly upregulated the expression of *Gabra 3* (34.2%; $p = .04$) and metabotropic glutamate receptor 2 ($Grm2$) (78.8%; $p = .0001$). In the HIV-1Tg rats, nicotine significantly upregulated the expression of *Grm2* (41.8%; $p = .026$) but significantly downregulated the expression of five genes, namely *Drd1a* (−28.3%; *p* = .01), *Drd2* (−55.1%; *p* = .032), *Drd3* (−39.1%; *p* = .038), *Drd5* (−48.9%; *p* = .007), and *Gabra5* (−47.4%; *p* = .037).

Effect of Nicotine and HIV-1 Proteins on Expression in the VTA of Genes that Are Implicated in Drug Addiction and Novelty-Seeking Behavior

With the same qRT-PCR approach, we examined expression of the same set of genes in the VTA region of the brains of HIV-1Tg and F344 rats treated with saline or nicotine ([Table 3\)](#page-5-0). Nicotine significantly upregulated the expression of *Gabra6* (334%; *p* = .022) but downregulated the expression of *Grm1* (−50%; *p* = .002) and *Grm2* (−61.5%; *p* = .029) in the F344 rats. In HIV-1Tg rats, nicotine significantly downregulated the expression of *Drd5, Gabra6, Grm1, Grm2*, and *Grm5*, by −28.8% to −55.5%, with a *p* value of .041 to .002. In addition, HIV-1 proteins significantly upregulated the expression of *Drd5* (110%; *p* = .002) and *Gabra6* (282.4%; *p* = .006) in the VTA.

Discussion

The findings of the current study extend our knowledge of HIV-1 infection such that the viral proteins were shown to affect noveltyseeking behavior in the open-field test by altering gene expression related to major neurotransmitter systems in the brain. HIV-1Tg

rats exhibited high novelty-seeking behavior with an abnormal baseline expression of D1- and D2-like dopamine and GABA receptors in the PFC–VTA–NAc neural circuits. We also found that nicotine differentially affects the transcription of excitatory and inhibitory neurotransmitter receptors in HIV-1Tg rats compared with F344 control rats. HIV-1Tg rats presented high levels of novelty-seeking behavior, suggesting that the viral proteins alter the central reward system by changing gene expression related to the response to nicotine.

Among humans, novelty seeking is one of the key personality traits associated with engaging in high-risk behaviors and with the risk of developing substance abuse. Recent studies of drug addiction in neuroAIDS have suggested that the use of addictive drugs is related to high-risk behaviors in persons infected with HIV-1. However, how the virus modifies this association during HIV-1 infection remains largely unclear. To explore this connection, we first investigated whether viral proteins cause any change in novelty-seeking behavior by using HIV-1Tg rats as an animal model that mimics the conditions present in human HIV-1 infection. HIV-1-infected patients receiving HAART show controlled viral replication with a persistent reservoir of viral proteins. These patients demonstrate mild to moderate cognitive deficits and progression of neurodegeneration. The HIV-1Tg rat mimics such patients, as these animals express seven of the nine viral proteins with controlled viral replication. Also, progression of cognitive and motor deficits occurs in such rats, similar to that seen in patients receiving HAART.⁵⁷⁻⁵⁹ Our model attempts to replicate the biology of HIV-1 infected individuals receiving HAART who are chronic smokers. We showed that viral proteins increase novelty-seeking behavior, as HIV-1Tg rats displayed increased time in, made more entries into, and traveled more distance in the center of the arena (see [Figure 1\)](#page-3-0) than did the control rats. These results are consistent with clinical findings showing that drug-dependent HIV-1-positive patients exhibit higher scores in novelty seeking and harm avoidance and lower scores in self-directedness relative to healthy controls according to the measurement of personality profile using

Table 3. Effect of Nicotine on Expression of Genes Related to Drug Addiction and Novelty-Seeking Behavior in the Ventral Tegmental Area of F344 and HIV-1Tg Rats

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the Temperament and Character Inventory.⁶⁰ However, our results show differences from the preclinical results, in that HIV-1Tg rats displayed a low locomotor response to placement in a novel environ ment.⁶¹ Other authors used adult female ovariectomized HIV-1Tg rats and tested novelty-seeking behavior in the activity chamber at 6, 7, and 11 months of age. These HIV-1Tg rats constantly exhibited a weaker response across monthly spaced testing of novelty-seeking behavior compared with control animals.^{[61](#page-8-7)} The discrepancy between our findings and those of Moran et al.⁶¹ may indicate that there are age- and sex-related differences in sensitivity to HIV-1 proteins in the CNS that could result in divergent neurobehavioral adaptations dur ing HIV infection. Similarly, a previous study involving age-depend ent motor function in HIV-1Tg rats found that ethanol-treated adult rats demonstrated a significant decrease in locomotor activity in the open-field test compared with water-treated controls.[43](#page-7-29) However, no difference was observed between ethanol- and water-treated adolescent HIV-1Tg rats in that study[.43](#page-7-29) Further research is needed to examine whether there are sex- and age-specific differences in the pathogenesis of neurobehavioral diseases in response to HIV-1 infec tion. Data from total distance traveled in the arena and distance traveled in the peripheral zone of the arena indicated no significant differences between strains; such results may be understood in the context of such extrapolated data on locomotor activity. Namely, HIV-1Tg rats traveled more distance in the center of the arena but may have traveled less distance in the peripheral zone, contributing to the result that both strains traveled similar total distances. The center zone of the arena, a novel environment, may have stimulated the rats' locomotor activity. However, we would like to note that locomotion has been found to "weakly predict" psychostimulant reward[.45](#page-7-26) Future studies aimed at more directly measuring rewardor drug-seeking behavior, such as conditioned place preference or drug self-administration, respectively, should be conducted with nicotine.

Next, we extended our behavioral findings by examining poten tial molecular mechanisms involved in the synergistic effects of HIV-1 proteins and nicotine in the neural circuits of the PFC–VTA– NAc (see [Table 4](#page-6-3)). For this objective, we analyzed the expression patterns of genes implicated in drug addiction and novelty-seeking behavior in HIV-1Tg and F344 rats after repeated nicotine or saline injections. In the saline-treated groups, HIV-1 proteins significantly increased the baseline mRNA concentrations of D2-like dopamine receptor *Drd3* in the PFC and D1-like dopamine receptor *Drd5* in the VTA, whereas they decreased the baseline expression of *Drd5* in the NAc of HIV-1Tg rats compared with the control rats. These results indicate that HIV-1 proteins lead to regionally dis tinct alterations in the mesocorticolimbic dopaminergic system. The HIV-1 proteins selectively target dopamine-rich regions in the brain and impair baseline dopaminergic synaptic transmission.^{[12](#page-7-3)[,14](#page-7-30)} The prefrontal dopaminergic circuits of the mesocorticolimbic system are the main targets of HIV-1 proteins and undergo dis ease-related alterations in synaptic connections that could lead to neurocognitive disorders.^{[15](#page-7-4),62-64} Our results provide further evidence for the idea that HIV-1 proteins have a selective effect on PFC dopaminergic activity by enhancing the amounts of D2-like dopamine receptors that may regulate the baseline dopamine quan tities in the NAc. Such differences in the mesocortical and mesoac cumbens dopaminergic systems might contribute to the enhanced striatal response to novel stimuli and addictive drugs, such as nico tine, in HIV-1Tg rats.

Table 4. Summary of Genes Significantly Modulated by Nicotine and HIV-1 Proteins in NAc, PFC, and VTA of HIV-1Tg and F344 Control **Rats**

Nac = nucleus accumbens; PFC = prefrontal cortex; VTA = ventral tegmental area.

Nicotine can directly or indirectly activate the central reward pathway by increasing dopamine through a number of mechanisms. Activation of VTA dopaminergic neurons through binding of nicotine to nicotinic acetylcholine receptors (nAChR) results in a direct increase in the extracellular dopamine concentration within the NAc. Nicotine also can indirectly modulate dopaminergic transmission in the central reward circuit by binding to nAChRs on glutamatergic and GABAergic neurons in the VTA.[65](#page-8-9),[66](#page-8-10) In our study, nicotine showed a strain-specific pattern of gene expression involved in GABAergic neurotransmission in the PFC–VTA circuits. Chronic nicotine treatment decreased the mRNA of genes encoding GABAergic receptors in HIV-1Tg rats, whereas the mRNA of these genes was increased in F344 rats. Further, a synergistic nicotine effect was found in dopaminergic transmission in the NAc, in that the mRNA level of D2-like dopamine recepto*r Drd5* was upregulated in HIV-1Tg rats. In contrast, no significant change in mRNA concentrations of dopaminergic receptors was found in the NAc of F344 rats. Further, genes encoding the glutamatergic receptor subunits that were altered by nicotine in the F344 rats were altered in the same direction in HIV-1Tg rats, including upregulation of *Grm2* in the PFC and downregulation of *Grm1* and *Grm2* in the VTA. These results indicate that chronic exposure to nicotine selectively disrupts the inhibitory control of the PFC on the VTA by acting on nAChRs and on local GABA interneurons. These interactions may lead to disinhibition of the dopaminergic neurons in the VTA, which could cause increased dopaminergic transmission in the NAc.^{[67](#page-8-11),68} Thus, our results indicate a nicotine-induced uncoupling of PFC-mediated inhibitory control over VTA–NAc reward circuitry, providing a critical neural mechanism for the loss of cognitive control that is observed in HIV-1-infected patients with nicotine dependence. To our knowledge, the present study provides the first demonstration of the synergistic effects of HIV-1 proteins and nicotine on the PFC–VTA–NAc neural circuits, which play a key role in dopamine-driven phenotypes, such as drug addiction and novelty-seeking behavior.

There are a few limitations of this study. First, in our expression analysis, we could not include as many addiction-related genes as we would have preferred. This limit was primarily the result of the small number of genes that could be included in each assay; further, only a small amount of RNA could be extracted from each specific brain region. We did not wish to pool tissues from multiple animals, as this method would increase the number of animals required for the study and fail to detect expression differences among individual animals. Second, for the same reasons, we did not examine expression differences at the protein level for those genes, although we realize that it is important to do so, because the

RNA differences we detected may not translate into differences in the amounts of various proteins.

In sum, our current study constitutes another step in the investigation of different implications of nicotine dependence in HIV-1 infected patients. Specifically, we focused on differentiating the effects of nicotine on the PFC–VTA circuits of GABAergic transmission in both the diseased and healthy states. Our results imply that treatments that enhance GABAergic transmission may provide a more effective smoking cessation method for HIV-1-positive smokers. However, further studies should be performed to clarify possible sex- and age-related difference, as well as to account for the activity of other neurotransmitter systems. Additionally, future research utilizing other drugs of abuse may provide useful information regarding the effects of viral proteins on the molecular mechanisms of reward-related behaviors.

Supplementary Material

Supplementary data are available at *Nicotine & Tobacco Research* online.

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Declaration of Interests

None declared.

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