

Short Communication

Human immunodeficiency virus type 1 Tat protein induces death by apoptosis in primary human neuron cultures

Deborah R New^{1,2}, Meihui Ma³, Leon G Epstein^{5,6,7}, Avindra Nath^{3,4}, and Harris A Gelbard^{5,6,8}

Departments of ¹Neurobiology and Anatomy, ²Dental Research, ⁵Neurology, ⁶Pediatrics, ⁷Microbiology and Immunology, ⁸Pharmacology and Physiology, University of Rochester Medical Center, 601 Elmwood Ave., Rochester, New York, 14642, USA; Departments of ³Medical Microbiology, and ⁴Internal Medicine (Section of Neurology), University of Manitoba, 523-730 William Ave., Winnipeg, Manitoba, R3E OW3, Canada

Neuronal loss in HIV encephalopathy remains a mystery since HIV-1 productively infects macrophage and microglia and only rarely infects neurons in the central nervous system. Apoptosis is a mechanism which may account for the loss of neurons in HIV-1 infected brain. Putative toxic factors that result in neuronal cell death in HIV-1 infection include the regulatory protein Tat, since this protein is known to be released from HIV-1 infected cells. Here we show that Tat induces cell death by apoptosis in cultured human fetal neurons producing characteristic morphological and biochemical features associated with apoptosis. These findings suggest that Tat may play an important role as a secreted, soluble neurotoxin in HIV-1 associated dementia.

Keywords: apoptosis; brain; central nervous system; dementia; HIV-1; pathogenesis; Tat; human cortical neuron cultures

Introduction

Neurodegeneration associated with infection of the central nervous system (CNS) by HIV-1 results in a CNS dysfunction termed HIV-1 associated dementia (Navia *et al*, 1986). The pathogenesis of HIV-1 associated dementia is not clear, but it is unlikely that productive infection of neurons with HIV-1 is responsible. In fact, HIV-1 infected neurons have rarely been demonstrated in post-mortem tissue, while macrophages and microglia are primarily found to be productively infected with the virus (Takahashi *et al*, 1996; Bernton *et al*, 1992; Wiley *et al*, 1986, 1991; Watkins *et al*, 1990). HIV-1 infection in the brain results in widespread myelin pallor, reactive astrogliosis, alterations of neocortical dendritic processes and neuronal loss without necrosis (Epstein and Gendelman, 1993; Sharer, 1992; Wiley *et al*, 1991). Several studies have demonstrated that HIV-1 infection of macrophages and microglia results in the release of soluble toxic factors that mediate neuronal death in *in vitro* models of HIV-1 neurotoxicity (Gelbard *et al*, 1994; Genis *et al*, 1992; Merrill and Chen, 1991; Giulian *et al*, 1990).

Recent evidence in post-mortem AIDS brain tissue indicates that neuronal apoptosis was present in cases with productive HIV-1 infection (Adle-Biassette *et al*, 1995; Gelbard *et al*, 1995; Petito and Roberts, 1995), but the cause of this pathological finding remains to be determined. Several authors have cited indirect mechanisms of neurotoxicity via macrophage-astroglial interactions mediated by a number of soluble factors including the cytokine tumor necrosis factor- α , and the phospholipid mediator platelet-activating factor (Gelbard *et al*, 1994; Genis *et al*, 1992; Merrill and Chen, 1991). Also *in vitro* studies indicate that the HIV-1 regulatory protein known as Tat, is released from infected cells (Ensoli *et al*, 1992). Several studies have shown Tat₁₋₈₆ and Tat peptides which contain the basic region within the first exon amino acid sequence are neurotoxic (Nath *et al*, 1996; Magnuson *et al*, 1995; Philippon *et al*, 1994; Hayman *et al*, 1993). Tat is a viral transactivator, and is expressed early in the viral life cycle (Sodroski *et al*, 1985). It activates transcription directed by the HIV-1 long terminal repeat (LTR), which in turn leads to increased expression of all viral genes, (Arya *et al*, 1985; Sodroski *et al*, 1985) and some cellular genes (Ensoli *et al*, 1992; Buonaguro *et al*, 1992; Helland *et al*, 1991; Vogel *et*

Correspondence: HA Gelbard
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al, 1988). This protein has been shown to induce neurotoxicity in human fetal neuron cultures which is mediated via excitatory amino acid receptors (Nath *et al*, 1996; Magnuson *et al*, 1995). The Tat protein has also been linked to apoptotic cell death in cultured peripheral blood mononuclear cells as well as a CD4-positive (CD4⁺) T cell line (Li *et al*, 1995; Purvis *et al*, 1995; Westendorp *et al*, 1995a, b).

In this study, we demonstrate that recombinant-Tat₁₋₈₆ and Tat₁₋₇₂ containing the basic region induces apoptosis in cultured human fetal cortical neurons in a dose-dependent fashion.

In situ detection of Tat induced apoptosis in human fetal neurons

Second trimester human cortical neuronal cultures, 28 days post-explantation, were treated with recombinant Tat₁₋₈₆ protein at various concentrations for 18 h then TUNEL stained for the free 3'-OH ends of cleaved DNA. Tat-induced apoptosis was found to be dose-dependent and was detectable at doses as low as 0.125 μ M (Figure 1). The dose response curve generated from our data indicates an LD₅₀ for neuronal apoptosis of 0.5 μ M. Doses of 1 μ M or greater resulted in detached cells (data not shown). Positively stained neurons were observed with chromatin aggregation, nuclear condensation, and apoptotic bodies, features consistent with apoptosis (Figure 2B). Similar apoptosis results were achieved with Tat₁₋₇₂. Neuron cultures were also treated with a vehicle control, where only rare TUNEL staining was observed (Figure 2A).

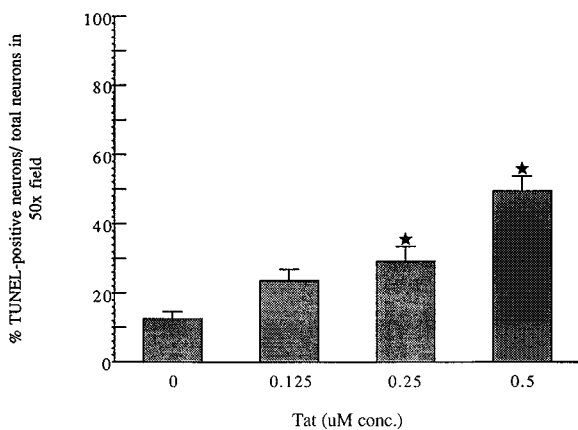


Figure 1 Primary human fetal neuron cultures were treated with Tat at the following doses; 0, 0.125, 0.25 and 0.5 μ M for 18 h. Cultures were fixed and TUNEL stained, the positive apoptosis immunostained neurons were analyzed in 16 random fields of each treatment by computerized morphometry. The histogram bars represent the average percent of TUNEL-positive apoptotic cells per total neurons per 50 \times field, SEM is shown by vertical lines (\star = p <0.001 vs control). This initial dose response curve at 18 h indicates an LD₅₀ at 0.5 μ M, higher doses at 1 μ M and greater were also analyzed, but resulted in detached cells.

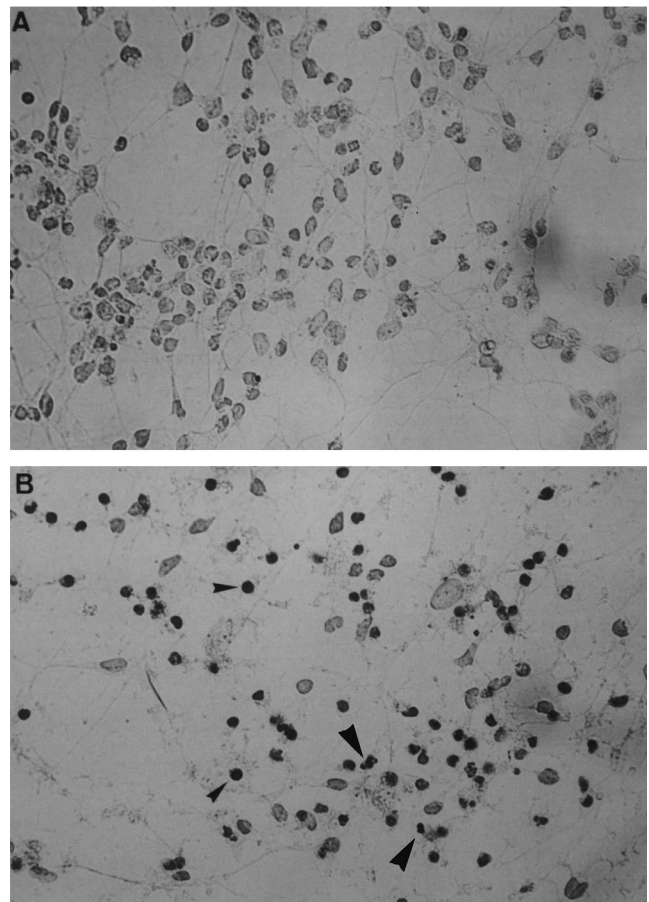


Figure 2 Primary human fetal neurons cultured for 28 days were treated with vehicle control (A) or 0.5 μ M of Tat (B) for 18 h. Neurons were subsequently fixed and stained *in situ* for new 3'-OH DNA ends generated by DNA fragmentation that results in chromatin condensation, a morphologic feature of apoptosis. The TUNEL assay or active labeling of cells by end labeling utilizing the 'Apoptag kit' (Oncor, Gaithersburg, MD) is indicated by the black precipitate seen as chromatin condensation (small arrows), and apoptotic bodies (large arrows).

Electron microscopy analysis of Tat treated neurons

Morphological indicators of apoptosis in the human fetal neurons treated with recombinant Tat₁₋₇₂ were analyzed by transmission electron microscopy (Figure 3). A number of changes consistent with apoptosis were observed in a subpopulation of neurons. Most commonly, condensation of the nuclear chromatin with relative preservation of the cytoplasmic structures, and the nuclear and cell membranes was noted (Figure 3A, B, D). Blebbing of the cell membrane was seen in some cells (Figure 3C). These cells showed only minimal changes in the nucleus and hence cell membrane blebbing was recognized as an early morphological change. Other cells showed partitioning and condensation of cytoplasm and nuclear material into membrane bound apoptotic bodies demonstrating end stage apoptotic changes. No phagocy-

tic cells were seen in these sections. The above morphological changes were absent in the untreated cultures or in cultures treated with solutions from which Tat had been immunoabsorbed (data not shown).

Cellular DNA analysis for Tat induced apoptosis

DNA was extracted from recombinant Tat₁₋₇₂ treated neurons and analyzed by agarose gel electrophoresis. A characteristic DNA cleavage ladder pattern specific for apoptosis was observed (Figure 4, Lane 2). No laddering was observed in the untreated cultures (Figure 4, Lane 3).

The phenomenon of apoptosis is usually associated with development, homeostasis, and aging processes. In HIV-1 infection, apoptosis occurs in the CD4⁺ subpopulation of lymphocytes in the peripheral blood as a consequence of the infection (Meyaard *et al*, 1992). The protein Tat also induces apoptosis in CD4⁺ lymphocytes (Ehret *et al*, 1996; Westendorp *et al*, 1995a,b). In the CNS, it has been suggested that pathological apoptosis occurs result-

ing in the loss of neurons (Adle-Biassette *et al*, 1995; Gelbard *et al*, 1995; Petit and Robert, 1995). This study focused on whether the HIV-1 regulatory protein Tat may play a role in mediating neuronal apoptosis.

Neuronal loss in HIV-1 infection is not directly due to the virus but may be due to toxic viral products that are secreted from infected cells. The HIV-1 regulatory protein Tat is one of many viral and cellular products secreted from HIV-1 infected cells. It has been suggested that Tat may be transported transcellularly from either infected or transfected cells via direct cell to cell contact in amounts sufficient to transactivate Tat-responsive promoter elements (Frankel and Pabo, 1988; Helland *et al*, 1991; Marcuzzi *et al*, 1992a,b). This suggests that exogenous Tat may affect uninfected cells during the course of HIV-1 infection.

Here we show convincing evidence that the HIV-1 protein Tat induced the morphological and biochemical features of apoptosis in cultured human neurons. Tat-induced apoptosis was demon-

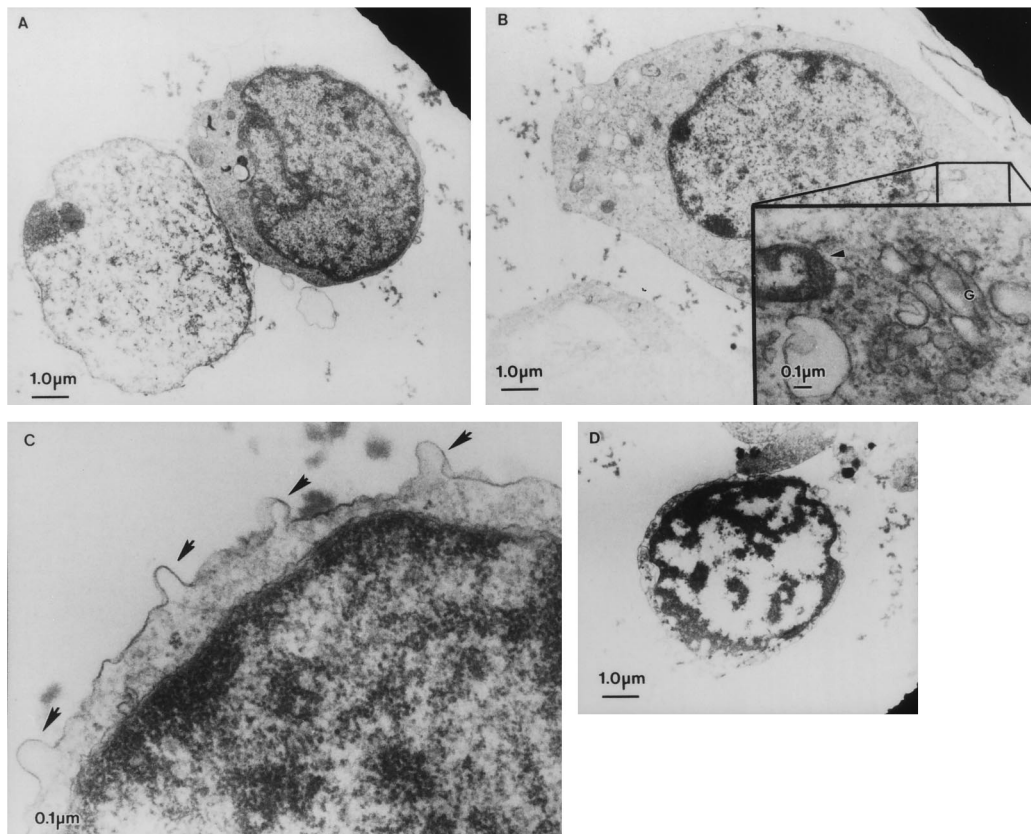


Figure 3 Morphological characteristics of Tat-induced apoptosis. (A) Two neurons are seen in close proximity. The one on the right shows normal morphological features while the other neuron shows degradation and condensation of the nuclear chromatin with preservation of the nuclear and cell membrane. (B) Condensation of nuclear chromatin with relative preservation of the cytoplasmic structures. Insert shows normal mitochondria and golgi apparatus. (C) Arrows show typical blebbing of the cytoplasmic cell membrane. (D) Another cell with clumping of the chromatin, degradation of cytoplasmic structures, but relative preservation of the nuclear and cell membranes.

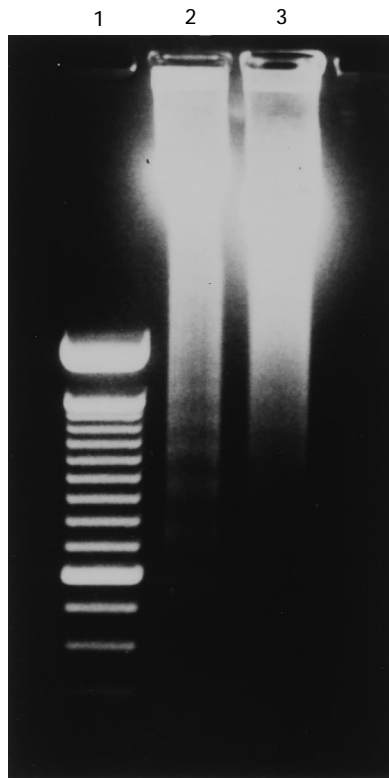


Figure 4 Tat induced DNA laddering. Lane 1. Molecular weight markers (1 kb) Lane 2. DNA extracts from Tat treated human fetal neurons analyzed by agarose gel electrophoresis show DNA laddering. Lane 3. Untreated neurons show absence of DNA laddering.

strated by several assays. First, we detected the presence of apoptotic neurons by demonstrating chromatin condensation and the formation of apoptotic bodies by *in situ* nick end labeling. We also demonstrated chromatin condensation and aggregation that abuts the inner surface of the nuclear membrane, and blebbing of the cytoplasmic cell membrane along with complete preservation of the integrity of the cytoplasmic organelles in the electron micrographs. Finally we demonstrated the typical DNA fragmentation ladder characteristic of apoptosis in the neuron cultures treated with Tat.

Recently, Tat was shown to activate non-NMDA excitatory amino acid (EAA) receptors and cause neurotoxicity in cultured neurons leading to increases in intracellular calcium and cell death (Nath *et al*, 1996; Magnuson *et al*, 1995). The neurotoxic domain resides within a conformationally dependent epitope within the first exon of Tat between residues 31 to 61 (Nath *et al*, 1996). These increases in intracellular calcium may induce apoptosis by activation of cellular enzymes (Conant *et al*, 1996).

Thus, recombinant Tat containing the basic region induces death in a subpopulation of cultured human fetal cortical neurons with the characteristic

morphological and biochemical features of apoptosis. Tat-induced necrosis of neurons was not observed at the light or electron microscope level. Our studies indicate that the first exon of Tat₁₋₇₂ is sufficient to cause cell death. Tat-induced apoptosis was only observed in cortical neurons and not in astrocytes (New, Angel, unpublished observations). We speculate that Tat-induced apoptosis can occur by more than one pathway, including glutamate receptor activation, oxidative stress and signaling via the pro-inflammatory cytokine TNF- α . The finding that Tat is a potent HIV-1-induced neurotoxin with an LD₅₀ of 0.5 μ M suggests that it may play a highly significant role in mediating neuronal apoptosis.

Methods

Human fetal brain tissue between gestational ages of 13 to 16 weeks were obtained and processed for culture as previously described (Gelbard *et al*, 1994). Cultured neurons were treated with full length recombinant Tat₁₋₈₆ protein. Once it was determined that only the basic region of the protein was required to elicit apoptosis, recombinant Tat₁₋₇₂ was used to complete these experiments. Since the pathological levels of Tat in the HIV-infected brain are unknown we initially ran dose response experiments with concentrations of Tat used for these experiments ranging from 0.125 μ M to 2.4 μ M.

Recombinant HIV-1 Tat₁₋₈₆ was expressed and purified as described before (Herrmann and Rice, 1993). Briefly a glutathione S-transferase Tat₁₋₈₆ fusion protein (AIDS research and reference reagent program, NIAID, NIH; Herrmann and Rice, 1993) was expressed and purified by adsorption to glutathione-sepharose beads (Pharmacia). Recombinant Tat₁₋₇₂ was also prepared as described before (Magnuson *et al*, 1995). Tat₁₋₇₂ was expressed as a fusion protein with a naturally biotinylated protein at the N-terminus in *E. coli* DH5aF'IQ (Gibco BRL). The biotin portion of the fusion protein was adsorbed to Softlink™ soft-release avidin resin (Promega), then cleaved from the resin with factor Xa, a serine endopeptidase (Boehringer Mannheim). Dithiothreitol (DTT) was added in each step of the purification of Tat₁₋₈₆ and Tat₁₋₇₂. Tat protein was suspended in a buffer containing 50 mM Tris (pH 8.0), 100 mM NaCl, 1 mM CaCl₂ and 0.5 mM DTT. The Tat protein was 95% pure by gel electrophoresis. The purified product was further analyzed by Western immunoblot analysis with a polyclonal antibody (AIDS Research and Reference Reagent Program, NIAID, NIH; Hauber *et al*, 1987). Its biological activity was measured by β -galactosidase (β -gal) gene activation in an HIV-1 long terminal repeat (LTR)- β -gal plasmid which had been transfected into HeLa cells (AIDS Repository, NIH) (Conant *et*

al, 1996). An *E. coli* strain containing the GST expression plasmid without the Tat sequence was expressed and purified as a sham Tat and used as a vehicle control.

Human fetal neuron cultures were treated with recombinant Tat for 18 h then stained to detect free 3'-OH ends of newly cleaved DNA *in situ* as described previously (Talley *et al*, 1995). Tat-induced apoptotic cells were stained with the *in situ* terminal deoxynucleotidyl transferase-mediated digoxigenin-dUTP nick end labeling (TUNEL) assay (Oncor, Gaithersburg, MD). Digitized images of TUNEL stained neurons in 16 or greater microscopic fields were analyzed for numbers of positively stained neuronal nuclei divided by total numbers of neurons per 50 × field using computerized morphometry (Imaging Research Inc., Onatario, Canada). Data were expressed as means ± SEM, with significance determined by one-way ANOVA.

Electron microscopy analysis of neurons was performed following 18 h treatment with 1 μM Tat. The cells were scraped and fixed in 2% glutaraldehyde, washed in a 0.1 M sodium cacodylate solution, suspended in agarose. The cells in agarose were cut into 1 mm³ blocks and fixed in 1% osmium tetroxide. The cells were dehydrated in successively increasing concentrations of acetone followed by propylene oxide and then embedded in Epon 812/araldite 502 (Marivac, Halifax, NS). Ultra-microtome sections of cells were stained with Reynolds lead citrate and viewed on a transmission electron microscope (Hayat 1981). Control cultures were similarly processed for electron microscopy.

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- DNA extraction and electrophoresis analysis was performed on human fetal neuronal cultures (4 × 10⁶ cells) treated with 2.4 μM Tat_{1–72} for 18 h. The neurons were separated by vigorous shaking and harvested by centrifugation at 400 × g for 5 min. The cell pellet was lysed in 0.5 ml lysis buffer (20 mM Tris-HCl pH 7.5, 4 mM EDTA, 3% SDS, 0.5 mg proteinase K per ml) at 50°C for 1 h. RNase H (20 g/ml, Boehringer Mannheim) was then added and the incubation continued for another 1 h. The lysate was centrifuged at 12 000 × g for 10 min and the supernatant was extracted with phenol followed by chloroform-isoamyl alcohol (24:1) and precipitated with ethanol at –70°C. The DNA pellet was dried and resuspended in TE buffer (10 mM Tris-HCl pH 8.0, 1 mM EDTA) and analyzed by 1.8% agarose gel electrophoresis.

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