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# Zika Virus Envelope Protein induces G2/M Cell Cycle Arrest and Apoptosis via an Intrinsic Cell Death Signaling Pathway in Neuroendocrine PC12 Cells

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#### Abstract

Clinical evidence suggests that there exists a strong correlation between Zika virus (ZIKV) infection and abnormal development of the nervous system. However, the underlying mechanisms remain to be elusive. In this study, recombinant lentiviral vectors coding for ZIKV structural proteins and truncations (prM-Env, M-Env and Env) were transduced into PC12 cells. Envelope (Env) overexpression induced significant inhibition of proliferation and triggered G2/M cell cycle arrest and apoptosis in PC12 cells. Flow cytometry and western blot analysis showed that the apoptosis was associated with up-regulation of both p53 and p21<sup>Cip1/Waf1</sup> and down-regulation of cyclin B1. Presence of aberrant nuclei clusters were confirmed by immunofluorescence staining analysis. The data indicate that overexpression of prM-Env, M-Env or Env led to apoptosis via an intrinsic cell death signaling pathway that is dependent on the activation of caspase-9 and caspase-3 and accompanied by an increased ratio of Bax to Bcl-2 in transduced PC12 cells. In summary, our results suggest that ZIKV Env protein causes apoptosis in PC12 cells via an intrinsic cell death signaling pathway, which may contribute to ZIKV-induced abnormal development of the nervous system.

Key words: ZIKV, envelope protein, G2/M arrest, mitotic catastrophe, apoptosis

# Introduction

The Zika virus (ZIKV), a member of the *Flaviviridae* family, is mainly transmitted by *Aedes aegypti* mosquitos [1]. ZIKV infections have become a major public health problem in the tropical and sub-tropical regions of the world with significant burden to ZIKV-infected patients and society as a

whole [2-4]. In Latin American countries, public recommendations to couples are to postpone pregnancy from six months to two years in ZIKV epidemic areas [5-7].

In ZIKV epidemic areas, most noticeably Brazil, a sharp increase in the incidence of microcephaly was

observed amongst newborns [8-10]. Additionally, a high incidence of Guillain-Barré syndrome was reported in Colombia [11] where ZIKV infections were highly prevalent from November of 2015 to March of 2016. Epidemiological data also showed that in 401 cases of ZIKV-associated secondary nervous system disease, 67% patients were diagnosed with Guillain-Barré syndrome [12]. These reports indicated that there exists a correlation between ZIKV infection and Guillain- Barré syndrome [13]. Recently, Souza, B.S and colleagues also found that ZIKV infection induces mitosis abnormalities and apoptotic cell death in human neural progenitor cells reinforcing the link between ZIKV and developmental neurological disorders abnormalities in the developing human brain, including such as microcephaly [14]. Despite much attention, however, the underlying mechanisms explaining this link remain elusive [15].

The ZIKV genome is presented as positive sense ssRNA encoding three structural proteins (Core (C), premembrane/membrane (prM/M) and envelope (Env)) and seven non-structural proteins (NS1, NS2A, NS2B, NS3, NS4A, NS4B, NS5) [16]. The prM/M protein acts as a chaperone for the correct folding of the Env protein during virus assembly and maturation [17, 18]. The Env protein plays a crucial role in viral infection and replication of *Flavivirus* [19]. Studies also have shown that the Env protein can induce cell-cycle G2/M accumulation [20]. In the present study, we first explored the effects of ZIKV prM-Env and two truncations (M-Env and Env) in neuroendocrine PC12 cells. Our results confirmed the effect of ZIKV Env protein in interfering with the cell cycle suggesting a role in abnormal nervous system development. The cell cycle arrest and apoptosis caused by Env was associated with increases in the expression levels of p21<sup>Cip1/Waf1</sup> and p53, as well as decreases in the levels of cyclin B1. Meanwhile, the apoptosis induced by Env was confirmed by upregulation of cleaved caspase-9, and caspase-3 in PC12 cells.

# **Materials and Methods**

# **Cells and Plasmids**

Human embryonic kidney (HEK) 293T cells and rat pheochromocytoma PC12 cells (ATCC, Manassas, VA) were cultured in Dulbecco's modified Eagle's medium (DMEM) (Hyclone, Logan, UT) supplemented with 10% fetal bovine serum (FBS) (Hyclone, US), 100 units/ml penicillin, 100 mg/ml streptomycin (Invitrogen, US) and maintain in 5% CO<sub>2</sub> at 37°C.

pLV-eGFP containing Cytomegalovirus (CMV) promoter was constructed in this lab. The packaging plasmids pMD2.G and psPAX2 were obtained from

Dr. Junming Yue (Department of Pathology, University of Tennessee Health Science Center). The cDNA fragment of the ZIKV prM-Env (Haiti strain, GenBank: KU509998.3) was obtained by chemical synthesis (IGE Biotechnology, China) and the cDNA fragments of M-Env and Env were obtained using PCR method. Primers are listed in Table 1. A six-histidine coding region was fused to the 3' end of each cDNA fragment. All cDNA fragments were cloned into the lentiviral expression plasmid pLV-eGFP. The resulting plasmids were designated as pLV-eGFP-ZIKV-prM-Env, pLV-eGFP-ZIKV-M-Env and pLV-eGFP-ZIKV-Env, which were identified by restriction enzyme digestion and partial sequencing.

 Table 1 Gene-specific primer sets used in this study

Gene	Prime name	Sequence(5'-3')	Product
name			size (bp)
M-Env	M-Env-F	GCTAGCGCTGTGACGCTCCCC	1758
	M-Env-R	ACGCGTTTAATGGTGATGGTGATG	
Env	Env-F	GCTAGCATCAGGTGCATAGGAG	1533
	Env-R	ACGCGTTTAATGGTGATGGTGATG	
Bax	BAX-F	GTCAGCTGCCACACGGAAG	81
	BAX-R	AACAACATGGAGCTGCAGAGGAT	
Bcl-2	Bcl-2-F	TCAGAGACAGCCAGGAGAAATCA	131
	Bcl-2-R	CCTGTGGATGACTGAGTACCTGAA	
Cyclin B1	Cyclin B1-F	GCGTGTGCCTGTGACAGTTA	135
	Cyclin B1-R	CCTAGCGTTTTTGCTTCCCTT	
p53	p53-F	AGCCAGACAGATCTTGTGAG	134
	p53-R	TCCCGCACGGCAAACTTTATC	
p21	p21-F	CCTGGTGATGTCCGACCTG	103
	p21-R	CCATGAGCGCATCGCAATC	
$\beta$ -actin	β-actin-F	AAGATTCGAGAAGAATACCCTGA	121
	β-actin-R	CTACCAACTGGTGGACGGACA	

#### Generation of recombinant lentiviral vectors

The recombinant lentiviruses LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env and LV-eGFP were packaged and titered as described previously [21, 22].

#### Lentiviral transduction in vitro

A total of  $10^5$  PC12 cells / well were prepared in a 6-wells plate. On the following day, the cells in each well were transduced with packaged recombinant lentiviruses at a multiplicity of infection (MOI) of 100 in DMEM medium containing 10% FBS with 6 µg /ml hexadimethrine bromide (Polybrene, Sigma, US). After 24 h, transduction media was replaced with fresh DMEM with 10% FBS and incubated for 24 h at 37°C and 5% CO<sub>2</sub>.

#### **Cell proliferation assay**

For detection of cell proliferation, MTT assay were performed at 48 h post-transduction. Total 10  $\mu$ l of MTT regent (Sigma-Aldrich, US) at the concentration of 5 mg/ml was added to each well. Following 3 h of incubation, 100  $\mu$ l of dimethyl sulphoxide (DMSO) detergent (Sigma-Aldrich, US) was added to each well. The plate was then incubated on a shaker at 4 °C for 2 h, and absorbance was measured at a 570 nm with a microplate reader (Bio-Rad, US).

# Cell cycle analyses

For detection of cell cycle, lentiviral transducedcells were washed with phosphate buffer saline (PBS) and fixed with 70% ice cold ethanol at 4°C for overnight, followed by treatment with PI (Sigma-Aldrich, US) solution at a final concentration of 50  $\mu$ g/ml. The DNA content was measured using flow cytometry (Beckman Coulter, Epics XL).

# Immunofluorescence analyses

PC12 cells were collected 48 h post-transfection and fixed with 4% paraformaldehyde (Solarbio, China) for 20 min, permeabilized with 0.1% Triton X-100 (Biosharp, China) for 15 min. Immunostaining was then done on the fixed cells as detailed previously [23] using a monoclonal antibody against 6×His tag (1: 500, Abcam, US) followed by Alexa Fluor 633 anti-mouse IgG (1:1,000; ThermoFisher Scientific, US) as the secondary antibody. Nuclei DNA staining was performed with 4', 6-diamidino-2phenylindole (DAPI, Sigma-Aldrich, US). Images were captured using a fluorescence microscope (Nikon, Japan).

# Annexin V-Alexa 647 apoptosis assay

For detection of cell apoptosis, lentiviraltransduced PC12 cells were collected and incubated with 1 µg/ml Annexin V-Alexa 647 (Abcam, US) and 2.5 µg/ml propidium iodide (PI, Sigma-Aldrich, US) for 30 min at room temperature, and then analyzed using a BD FACSCanto II cytometer (Becton Dickinson). Specific inhibitors of caspase-3 (Ac-DEVD-CMK, Santa Cruz Biotechnology, US), caspase-8 (Z-LEHD-FMK, Abcam, US) and caspase-9 (Z-LEHD-FMK, Abcam, US) were used to confirm the assay specificity.

# Western blot analysis

Western immunoblots were run as described previously [21]. Primary antibodies and their sources were as follows. Anti-total p53, anti-cyclin B1, anti-Bax, anti-Bcl-2, anti-CDK1, anti-phospho-CDK1, anti-activated caspase 8, anti-activated caspase 9, and anti-activated caspase 3 were from Cell Signaling Technology. Anti-ZIKV Env monoclonal antibody was purchased from Zoongen Co., Ltd (Beijing, China). Anti-p21<sup>Cip1/Waf1</sup> and anti-6×His tag was from Abcam. Anti-β-actin and the secondary antibodies horseradish peroxidase-conjugated anti-rabbit IgG and anti-rabbit IgG were from Sigma-Aldrich. Anti-goat IgG was from Santa Cruz Biotechnology.

#### Quantitative real-time RT-PCR analysis

Total RNA was isolated from transduced PC12 cells using TRIzol® (Sigma, US), and then converted into cDNA with iScript<sup>TM</sup> Select cDNA Synthesis Kit (Bio-Rad, US). Quantitative real-time RT-PCR was performed on an ABI 7500 real-time PCR system (Applied Biosystems, US) as described previously [23]. Primers for target genes (*Bax, Bcl-2, Cyclin B1, p53, p21 and β-actin*) are listed in Table 1. The samples were quantified by the comparative  $\Delta\Delta C_T$  method by using rat β-actin as the internal standard.

# Statistical analysis

All continuous variables are presented as means  $\pm$  SEM. Parametric data were analyzed using unpaired two-tailed t tests, for comparisons between two groups, and 1-way ANOVA, followed by Bonferroni post hoc test for multiple-comparison test, using Prism 6.0 (GraphPad Software Inc, La Jolla, CA, USA). Values of p < 0.05 were considered statistically significant.

# Results

# Lentivirus- mediated overexpression of ZIKV prM-Env, M-Env and Env in PC12 cells

Cell supernatants and cells were collected at 12, 24, 48 h post-transduction to assess the expression of ZIKV prM-Env, M-Env and Env in transduced PC12 cells using Western blot analysis. The results showed that they were detectable as early as 24 h post-transduction in cell lysates (Figure 1A), but not in supernatants (data not shown). The expressions of prM-Env, M-Env and Env were further confirmed by indirect immunofluorescence at 48 h post-transduction. The results revealed that these proteins were localized in the nucleus and cytoplasm of transduced PC12 cells (Figure 1B).

#### prM-Env, M-Env and Env inhibited PC12 cell growth and induced G2/M arrest in lentivirus-transduced PC12 cells

MTT assays were performed to determine the effect of prM-Env, M-Env and Env on transduced PC12 cell growth. We demonstrated that lentivirusmediated overexpression of prM-Env, M-Env and Env significantly inhibited PC12 cells growth when compared with LV-eGFP transduced cells, but not amongst themselves (Figure 2A).

To examine the effect of prM-Env M-Env and Env on cell cycle, flow cytometric analysis was performed at 48 h post-transfection. The results showed that overexpression of prM-Env, M-Env and Env significantly increased the percentage of cells in the G2/M phase in transduced PC12 cells (Figure 2B).



В

Α



12 h

24 h

48 ł

ZIKV Env

β-actin

**Figure 1.** Lentiviral vectors-mediated ZIKV Env, M-Env and prM-Env overexpression in PC12 cells. Cells were infected with LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env or LV-eGFP at an MOI of 100. (**A**) At 12, 24 and 48 h after transduction, cell protein extracts were subjected to Western blot analysis using an anti-Env antibody. (**B**) At 48 h after transduction, ZIKV Env, M-Env or prM-Env immunoreactivity was also detected using a mouse monoclonal anti-His antibody followed by anti-mouse Alexa Fluor 633-conjugated antibody (Scale bar= 100 µm).

The percentage of G2/M phase LV-ZIKV-prM-EnV, LV-ZIKV-M-Env and LV-ZIKV-Env transduced cells increased by 20% compared to that of LV-eGFP infected PC12 cells (Figure 2C). Additionally, no significant difference was observed in G2/M phase ratio among LV-ZIKV-prM-Env, LV-ZIKV-M-Env and LV-ZIKV-Env transduced PC12 cells. The Env-induced G2/M arrest in transduced PC12 cells was further confirmed by downregulation of G2/mitotic-specific cyclin B1 and inhibition phosphorylation of cell cycle kinase CDK1 (Figure 2D&E).

# prM-Env, M-Env and Env induced mitotic catastrophe in lentivirus-transduced PC12 cells

Aberrant nuclei clusters were observed in LV-ZIKV-prM-Env, LV-ZIKV-M-Env and LV-ZIKV-Envtransduced PC12 cells (Figure 3A). This indicated that prM-EnrEnv, M-Env and Env induced significant mitotic catastrophe in transduced PC12 cells. Approximately 12 to 15% of PC12 cells, expressing prM-Env M-Env or Env showed different abnormal nuclei compared to the control cells (Figure 3B).

#### prM-Env, M-Env and Env induced apoptosis in lentivirus-transduced PC12 cells via an intrinsic cell death signaling pathway

We further examined whether prM-Env, M-Env and Env could elicit apoptosis in transduced PC12 cells since they caused the increment of G2/M phase and induction of mitotic catastrophe in these cells. Annexin V, an endogenous 36-kDa protein that binds to phosphatidylserine, have been identified and reported as potential apoptosis probes [24]. Annexin V-Alexa Fluor 647-conjugated staining was performed to evaluate apoptosis in PC12 cells at 48 h post-transduction. Flow cytometric results indicated a significantly higher proportion of annexin V positive



**Figure 2.** prM-Env, M-Env and Env inhibited PC12 cell growth and induced G2/M arrest in transduced PC12 cells. PC12 cells were transduced with LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env or LV-eGFP at an MOI of 100. After 48 h of incubation, cells were collected and subjected to MTT assay, flow cytometric analysis, real-time RT-PCR or Western blot. (A) Cell proliferation determined by MTT assay. \*: P<0.05 versus with LV-eGFP. (B) Representative data for lentivirus-induced G2/M cell cycle arrest measured by flow cytometry at 48 h after treatment. The y axis represents the cell count, and the x axis represents the DNA content. The experiments were performed at least three times. (C) Data showing percentages of PC12 cells in G2/M phase. \*: P<0.05 versus with LV-eGFP. (E) Cyclin B1 and CDK1 expression in transduced PC12 cells. Data are representative of three experiments.

cells in the prM-Env, M-Env and Env expressing groups when compared to the control group (Figure 4A & 4B). Apoptosis associated proteins were then detected by western blot to explore the mechanism for Env-induced apoptosis. Results indicate that p53, p21<sup>Cip1/Waf1</sup>, Bax, cleaved caspase-3, cleaved caspase-8, cleaved caspase-9 were increased, while Bcl-2 was decreased in PC12 cells transduced with LV-ZIKV-prM-Env, LV-ZIKV-M-Env and LV-ZIKV-Env when compared with LV-eGFP (Figure 4C). The mRNA relative levels of p53, p21<sup>Cip1/Waf1</sup>, Bax, and Bcl-2 were then measured by real-time RT-PCR in transduced PC12 cells. The results revealed significant up-regulation of p53, p21Cip1/Waf1, and Bax mRNA relative levels in LV-ZIKV-Env, LV-ZIKV-M-Env, and LV-ZIKV-prM-Env transduced PC12 cells while the mRNA levels of Bcl-2 (Figure 4D) in PC12 cells with

Env, M-Env and prM-Env overexpression were significantly lower than that of control PC12 cells.

Lastly, treatment of PC12 cells with the caspase-3 inhibitor Ac-DEVD-CMK (100  $\mu$ mol/L) significantly reduced the apoptosis elicited by LV-ZIKV-prM-Env, LV-ZIKV-M-Env or LV-ZIKV-Env transduction (Figure 5A & 5B). Furthermore, similar inhibition of ZIKV Env-induced apoptosis was obtained by treatment of the PC12 cells with the caspase-9 inhibitor Z-LEHD-FMK (100  $\mu$ mol/L) while no inhibition effects were observed with caspase-8 inhibitor Z-IETD-FMK (100  $\mu$ mol/L). These data suggest the involvement of a caspase-9-mediated intrinsic signaling pathway followed by downstream activation of caspase-3 in the apoptosis elicited by overexpression of ZIKV prM-Env, M-Env and Env in PC12 cells.



**Figure 3.** ZIKV Env, M-Env and prM-Env induced mitotic catastrophe in lentiviruses-transduced PC12 cells. (**A**) PC12 cells were transduced with LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env or LV-eGFP at an MOI of 100, immunostained with an anti-His antibody followed by anti-mouse Alexa Fluor 633-conjugated antibody, and then counterstained with DAPI for nuclear staining (Scale bar= 100 µm). Mitotic catastrophe was characterized by the appearance of aberrant nuclei clusters. The number of aberrant nuclei clusters in each treatment condition was counted from 5 random fields per well by an individual who was blinded to the treatment. (**B**) Data are presented as a percent of the total number of cells on the dish. \*\*: P<0.01 versus with LV-eGFP.

#### Discussion

The severe clinical symptoms caused by ZIKV infection have aroused world-wide attention. Significant breakthroughs and new questions have been made during the rapid attempt to understand the relationship between ZIKV and abnormal development of nervous system [15, 25, 26]. The neuronal apoptosis induced by ZIKV has been confirmed both in vitro [14] and in vivo [13, 27]. The data presented here indicate that overexpression of ZIKV prM-Env, M-Env and Env in PC12 cells induced inhibition of cell growth, G2/M arrest and mitotic catastrophe. Furthermore, the overexpression of these proteins led to apoptosis via an intrinsic cell death signaling pathway that is dependent on activation of caspase-9 and caspase-3. Finally, the apoptosis induced by ZIKV prM-Env, M-Env and Env

overexpression was accompanied by up-regulation of both p53 and p21<sup>Cip1/Waf1</sup>. Collectively, the observations presented here indicate the ability of increased ZIKV prM-Env, M-Env and Env expression to induce apoptosis in neuroendocrine PC12 cells *in vitro* suggesting that the ZIKV Env protein may play a crucial role in ZIKV-induced neuronal apoptosis.

The prM-Env protein of ZIKV is a major candidate antigen for vaccine development [28]. We had initially planned to produce this protein in HEK293T by transduction with LV-prM-Env, LV-M-Env and LV-Env but inadvertently observed significant apoptosis (data not shown). HEK293 cells are ubiquitously used for many purposes [29] but can also be used for analysis of neuronal synapse formation [30] and has even been argued that HEK293 cells are of neuronal lineage [31]. Our accidental finding in HEK293 cells led to this investigation of the



**Figure 4.** ZIKV Env, M-Env and prM-Env induced apoptosis in lentiviral vectors-transduced PC12 cells. PC12 cells were transduced with LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env or LV-eGFP at a MOI of 100. (**A**) At 48 h post- transduction, indicated cells were collected and double-stained with Alexa Fluor 647 conjugated annexin V and PI for flow cytometric analysis. The gate setting distinguished among living (lower left), necrotic (upper left), early apoptotic (lower right) and late apoptotic (upper right) cells. (**B**) Quantification of the annexin V -positive cells as a percent of the total number of cells in the dish. Columns, mean of three experiments; bars, SE. \*\*: P<0.01 versus with LV-eGFP. (**C**) Expression of p53, p21Cipl/Waf1, cleaved caspase-3, 8, and 9, Bax and Bcl-2 in lentiviral vectors-transduced PC12 cell lines. Western blots were performed at 48 h post-transduction.  $\beta$ -actin was used as a loading control. Western blots are representative of three different experiments. (**D**) The relative expression levels of p53, p21, Bax and Bcl-2 in transduced PC12 cells were measure by real-time RT-PCR at 48 h post-transduction. All data were normalized against levels of  $\beta$ -actin mRNA expression within the same sample. Columns, mean from three separate experiments; bars, SE. \*: P<0.05 versus with LV-eGFP.

role of prM-Env in ZIKV induced neuronal apoptosis and further corroborates the relationship between ZIKV infection and neurological developmental disorders [13, 14, 26].

Mechanistically, our results showed that ZIKV prM-Env, M-Env and Env overexpression elicited not only up-regulation of p53, but also changes in downstream effectors: up-regulation of p21<sup>Cip1/Waf1</sup>, down-regulation cyclin B1 and an increase in the ratio of Bax/Bcl-2 (Figure 6). The p53 tumor suppressor is at the heart of several fundamental cellular signaling pathways. Arguably, the most important pathways for p53's tumor-suppressive role are those involved in the induction of cell cycle arrest and apoptosis [32, 33]. The cyclins are a family of proteins that control the progression of cells through the cell cycle by

activating cyclin-dependent kinase (Cdk) enzymes. Cyclin B1 and Cdk1 are responsible for the transition of the cell from the G2 to the M phase and their down-regulation will lead to G2/ M cell cycle arrest [34]. Recent evidence also suggests that p21 may participate in apoptosis in p53-dependent pathways [35]. p53-induced apoptosis results from overlapping downstream pathways that both suppress mitogenic and survival signaling and promote proapoptotic signaling. In this context, p53 can up-regulate the proapoptotic Bcl-2 family member Bax and possibly transcriptionally repress the antiapoptotic protein Bcl-2 [36]. Therefore, p53-mediated up-regulation of Bax and perhaps concomitant down-regulation of Bcl-2 are pivotal in shifting the ratio of Bax/Bcl-2 in the cell and ultimately favoring apoptosis.



Figure 5 Effects of caspase inhibitors on ZIKV Env-mediated apoptosis in PC12 cells. (A) Cells were transduced with LV-ZIKV-prM-Env, LV-ZIKV-M-Env, LV-ZIKV-Env or LV-eGFP (100 MOI) for 6 h followed by treatment with either caspase-3 inhibitor Ac-DEVD-CMK (100 µmol/L), caspase-8 inhibitor ZIETD-FMK (100 µmol/L), caspase-9 inhibitor ZIETD-FMK (100 µmol/L) or control solvent (DMSO) for 24 h. Caspase-3, caspase-8 and caspase-9 were detected by western blot. (B) Treated cells were double- stained with Alexa Fluor 647 conjugated annexin V and PI. The annexin V-positive cells as a percent of the total number of cells were measured from flow cytometric analysis. \*\*: P<0.01 versus with control (DMSO-treated) cells. Data are representative of three experiments.



Figure 6 Schematic diagram of ZIKV Env-induced cell cycle arrest and apoptosis in PC12 cells. prM-Env, M-Env and Env induced apoptosis in lentivirus-transduced PC12 cells via an intrinsic cell death signaling pathway. p53 may involve ZIKV Env-induced G2/ M cell cycle arrest and apoptosis.

In all, this may indicate the involvement of p53 in the G2/ M cell cycle arrest and apoptosis induced by ZIKA Env in PC12 cells. We also found that all of cleaved caspase-3, caspase-8, 9 expression levels were significantly increased in PC12 cells expressing prM-Env, M-Env or Env. However, only the inhibitors of caspase-3 and caspase-9, not caspase-8, could block the apoptosis in these cells, which indicates that the apoptosis was induced via an intrinsic cell death signaling pathway.

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#### **Author Contributions**

JL carried out lab work, analysis, and writing, QQ, XL, ZZ, AL, WH, HY, XS, XL, XY, JJ, WS, MY, YY and HY carried out lab work, and JL, HW, WW advised and contributed significantly to writing and analysis.

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# **Competing Interests**

The authors have declared that no competing interest exists.

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