

NT4(Si)-p53(N15)-antennapedia induces cell death in a human hepatocellular carcinoma cell line

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Abstract

AIM: To construct the recombinant lentivirus expression plasmid, pLenti6/V5-NT4 p53(N15)-antennapedia (Ant), and study its effect on HepG2 cells.

METHODS: Plasmid pLenti6/V5-NT4 p53(N15)-Ant was constructed incorporating the following functional regions, including signal peptide sequence and pro-region of neurotrophin 4, N-terminal residues 12-26 of p53 and 17 amino acid drosophila carrier protein, Ant. Hepatocellular carcinoma (HepG2) cells were used for transfection. 3-[4,5-dimethyl-thiazol-2yl]-2,5 diphenyl tetrazolium bromide (MTT) assay, lactate dehydrogenase (LDH) release assay, transmission electron microscopy (TEM) and flow cytometric analysis (FCM) were employed to investigate the effects of LV-NT4(Si)-p53(N15)-Ant *in vitro* on HepG2 cells. *In vivo* experiment was also performed to investigate the inhibitory effect of LV-NT4(Si)-p53(N15)-Ant on tumor growth in nude mice.

RESULTS: LV-NT4(Si)-p53(N15)-Ant significantly suppressed the growth of HepG2 cells. MTT assay showed that the growth of HepG2 cells was much more significantly inhibited by LV-NT4(Si)-p53(N15)-Ant than by LV-EGFP. The inhibition rate for HepG2 cell growth in the two groups was 46.9% and 94.5%, respectively, 48 h after infection with LV-NT4(Si)-p53(N15)-Ant, and was 33.9% and 95.8%, respectively, 72 h after infection with LV-NT4(Si)-p53(N15)-Ant ($P < 0.01$). Light microscopy and TEM showed morphological changes in HepG2 cells infected with LV-NT4(Si)-p53(N15)-Ant, but no significant changes in HepG2 cells infected with LV-EGFP. Changes were observed in ultra-structure of HepG2 cells infected with LV-NT4(Si)-p53(N15)-Ant, with degraded membranes, resulting in necrosis. LDH release from HepG2 cells was analyzed at 24, 48, 72 and 96 h after infection with LV-NT4(Si)-p53(N15)-Ant and LV-EGFP, which showed that LDH release was significantly higher in LV-NT4(Si)-p53(N15)-Ant treatment group (682 IU/L) than in control group (45 IU/L, $P < 0.01$). The longer the time was after infection, the bigger the difference was in LDH release. FCM analysis showed that LV-NT4(Si)-p53(N15)-Ant could induce two different kinds of cell death: necrosis and apoptosis, with apoptosis being the minor type and necrosis being the main type, suggesting that LV-NT4(Si)-p53(N15)-Ant exerts its anticancer effect on HepG2 cells by inducing necrosis. The *in vivo* study showed that LV-NT4(Si)-p53(N15)-Ant significantly inhibited tumor growth with an inhibition rate of 66.14% in terms of tumor size and weight.

CONCLUSION: LV-NT4(Si)-p53(N15)-Ant is a novel recombinant lentivirus expression plasmid and can be used in gene therapy for cancer.

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Key words: Gene therapy; Lentivirus vector; Anticancer; Necrosis; LV-NT4(Si)-p53(N15)-Ant; Hepatocellular carcinoma cell line

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INTRODUCTION

Hepatocellular carcinoma (HCC) is one of the most common causes of death worldwide, especially in Asian countries^[1]. To date, none of the conventional treatment modalities can completely eradicate HCC cells. In recent years, gene therapy has been evaluated as a novel treatment modality for HCC^[2-5], while conventional treatment modalities, including surgery, chemotherapy and liver transplantation, are still used^[6-8].

p53 is an important tumor suppressor gene which regulates many important cellular activities, including apoptosis. Mutations and deletions of the *p53* gene are common in HCC in a number of geographic regions. Since the mutation incidence ranges 5%-50%, *p53* has become an ideal target for gene therapy. Kanovsky *et al*^[9] and Do *et al*^[10] reported that a *p53* peptide, synthesized from residues 12-26 and fused with the *Drosophila* carrier protein antennapedia (Ant), can induce rapid tumor cell necrosis in all breast and pancreatic cancer cell lines irrespective of its status and exhibits a low cytotoxicity to normal cells, which is uncommonly observed in traditional cancer therapy. Human HCC cell line, HepG2, contains the wild-type *p53* gene and can thus be used in research of the relation between HCC and *p53* peptide gene therapy.

It is important to study the transfer of fusion gene and the secretion of expressed protein for the assessment of enhanced cancer-killing effects of a protein. In this study, NT4 signal peptide and its pro-region were used as the regions responsible for protein and peptide secretion from cells. Lentivirus gene expression plasmids were constructed for NT4(Si)-ADNF-9 and NT4(Si)-NAP fusion proteins containing the NT4 signal peptide and pro-region to enhance their expression. The restriction enzyme site *NaeI* at the NT4 signal peptidase fissure site and two restriction enzyme sites, *BamH I* and *Xho I*, in NT4(Si)-*p53*(N15)-Ant could ensure the correct construct. A novel recombinant lentivirus expression plasmid, pLenti6/V5-NT4 *p53*(N15)-Ant, was constructed, containing a signal peptide sequence and pro-region of neurotrophin 4 (NT4) fused to *p53*(N15)-Ant peptide. Its effect on HepG2 cells, both *in vitro* and *in vivo*, was investigated.

MATERIALS AND METHODS

Cells and cell culture

Human hepatoma HepG2 cells containing wild-type *p53* were cultured in RPMI 1640 containing 10% fetal bovine serum (FBS). 293T human kidney cells were

grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% FBS, 100 U/mL penicillin and 100 µg/mL streptomycin at 37°C in a humidified atmosphere containing 50 mL/L CO₂. All cell lines were supplied by Xi'an Huaguang Bioengineering Company (Xi'an, China).

Plasmid construction and virus titration

p53(N15)-Ant was synthesized by Beijing Sun Biotechnology Co, according to Kanovsky *et al*^[9] and Genbank. Two restriction enzyme sites (*Nae I* and *Xho I*) were introduced upstream and downstream of this sequence, respectively. *p53*(N15)-Ant was subcloned into pGEM-T-easy for sequencing, then into pBV220/NT4. Signal peptide sequence and pro-region of NT4 were cloned from human genomic DNA and subcloned into the *BamH I* / *Nae I* sites of pBV220. After digestion with restriction enzymes, the resulting NT4 *p53*(N15)-Ant gene with *BamH I* and *Xho I* sites from the pBV220/NT4-*p53*(N15)-Ant plasmid was inserted into the multiple cloning site of the expression plasmid, pLenti6/V5-D-TOPO (Invitrogen) containing a cytomegalovirus (CMV) promoter, upstream of the inserted gene. The resulting plasmid was named pLenti6/V5-NT4 *p53*(N15)-Ant.

To generate a control plasmid containing green fluorescent protein, EGFP was amplified from EGFP-C2 by PCR. The sequences of sense and anti-sense primers are 5'CGGGATCCATGGTGAGCAAGGGCGAGG-3' and 5'CGCTCGAGTCAAGTCCGGCCGGACTTGTAC-3', respectively. The resultant PCR fragments were digested with *BamH I* and *Xho I* (underlined), subcloned into pLenti6/V5-D-TOPO, and verified by DNA sequencing.

The four-plasmid-based lentiviral expression system was purchased from Invitrogen. Briefly, four kinds of plasmid including 4.2 µg of pLP1, 2 µg of pLP2, 2.8 µg of pLP/VSVG and 3 µg of lentivector pLenti6/V5-NT4 *p53*(N15)-Ant or pLenti6/EGFP, were cotransfected into 293T cells using the calcium phosphate coprecipitation method. The conditioned medium was harvested 72 h after transfection and filtered through a 0.45 µm filter. Concentrated viral stocks were prepared by ultra-centrifugation of 3 mL conditioned medium at 50 000 *g* for 1.5 h at 4°C in a SW41 rotor (Beckman). The pellet was resuspended in 30 µL complete medium and stored at -80°C. The resulting recombinant lentivirus was named LV-NT4 *p53*(N15)-Ant and LV-EGFP, respectively.

Viral titer was measured by assessing the viral p24 antigen concentration using ELISA (Beckman Coulter, Fullerton, California, USA), showing that one microgram per milliliter of p24 corresponds to approximately 2×10^6 transducing units per milliliter of EGFP virus, as assessed by titration in 293 T cells.

Lentivirus-mediated gene transfer and expression in HepG2 cells

HepG2 cells were seeded into 6-well plates at a density of 1×10^6 cells/well, allowed to adhere for 24 h, and then infected with EGFP lentivirus at multiplicities of infection (MOI) of 4 transducing units (TU)/cell. EGFP-positive cells were observed and counted under a fluorescence microscope 48 h after infection.

Expression of the *NT4 p53(N15)-Ant* gene was detected by reverse transcription-polymerase chain reaction (RT-PCR). HepG2 cells were seeded in 100 mL culture flasks at a density of 1×10^6 cells/flask and infected with NT4-p53(N15)-Ant lentivirus at a MOI of 4 TU/cell. Forty-eight hours after infection, HepG2 cells were scraped from the flask surface with a brush (Paro-Isola, Thalwil, Switzerland). Total RNA was extracted using Trizol reagent (Gibco). A reverse transcription (RT) step was carried out for 60 min at 37°C using 1 µg total RNA treated with DNase and M-MLV reverse transcriptase (Invitrogen, USA) in the presence of random primers. PCR was performed in 50 µL reaction volume containing 10 µL (about 2.5 ng) cDNA, 1 µL of 10 mmol/L dNTP, 1 U of Taq DNA polymerase, 5 µL of $10 \times$ Taq buffer, and 1 µL 50 pmol of forward primer (5'-CG GATCCATGCTCCCTCTCCCTCATGC-3') and reverse primer (5'-CCTCGAGTCATCCGCGCTGTACCTTTACC-3') of the *NT4 p53(N15)-Ant* gene. Samples were subjected to RT-PCR at the following conditions: pre-denaturation for 5 min at 94°C, followed by 30 cycles of denaturation for 60 s at 94°C, annealing for 60 s at 60°C, extension for 90 s at 72°C, and a final extension for 5 min at 72°C. The RT-PCR products were separated by electrophoresis on 2% agarose gels (Qiagen, USA).

Immunohistochemical detection of p53(N15)-Ant expression in lung cancer H1299 cells

H1299 cells (null for p53) were seeded onto cover slips and infected with LV-NT4(Si)-p53(N15)-Ant at a MOI of 4 TU/well. Polybrene was added to a final concentration of 8 µg/mL. Twelve hours after infection, 2 mL of a new DMEM medium was added. Forty-eight hours after infection, H1299 cells were washed and fixed with cold acetone. Protein expression of p53(N15)-Ant in H1299 cells was determined by immunocytochemical staining with anti-p53 antibody (DO-1) (Santa Cruz Biotechnology, Santa Cruz, CA, USA) using the Vectastain Elite ABC kit following its manufacturer's instructions (Vector Laboratories, Burlingame, CA, USA).

MTT assay

To determine the effect of lentivirus-NT4(Si)-p53(N15)-Ant on HCC, HepG2 cells were seeded in 96-well plates at a concentration of 5×10^3 cells, grown for 24 h, infected with lentivirus-NT4(Si)-p53(N15)-Ant or lentivirus-EGFP at a MOI of 4 TU/cell with polybrene (8 µg/mL) for 12 h. The culture medium was then replaced with a fresh medium. MTT assay was performed at different time points (24, 48 and 72 h) after infection with lentivirus according to its manufacturer's instructions (Xi'an, China).

Examination of HepG2 cell morphology under light microscope

After infection with NT4(Si)-p53(N15)-Ant, morphology of HepG2 cells was observed under an inverted light microscope and recorded.

Examination of ultra-structure of HepG2 cells under electron microscope

Forty-eight hours after infection with NT4(Si)-p53(N15)-Ant, HepG2 cells were trypsinized, fixed with 2.5% glutara-

ldehyde for 2 h at 4°C, washed with 0.1 mol/L dimethyl arsenic trioxide buffer, fixed with 1% osmium acid for 2 h, dehydrated, and then embedded in 618 domestic epoxy resin, followed by polymerization for 24 h. The embedded block was cut into ultra-thin sections. The sections were double stained with uranyl acetate and lead citrate, and observed under a transmission electron microscope.

LDH release assay

To determine LDH leakage into the extra-cellular fluid, supernatant was collected at different time points (24, 48, 72, and 96 h) after infection with NT4(Si)-p53(N15)-Ant. LDH in the supernatant was detected with a RA-100 automatic biochemical analyzer.

Annexin V and PI double staining

When HepG2 cells were grown to 2×10^5 /mL, they were seeded into 9-well plates at a volume of 3 mL/well, and left to adhere for 24 h. The cells were then divided into control group and lentivirus treatment group. The cells in control group were grown in a serum-free medium. Forty-eight hours after infection with lentivirus, the cells were washed twice with PBS at 4°C, re-suspended in 250 µL of a combination buffer solution, and the cell concentration was adjusted to 1×10^5 /mL. Five microliter of Annexin V/FITC and 5 µL of 20 µg/mL propidium iodide were added to 100 µL of cell suspension. After incubation in the dark for 15 min, the cells were analyzed by flow cytometry (FCM).

Inhibition of HepG2 cells in nude mice by LV-NT4(Si)-p53(N15)-Ant

Twenty 6-8 wk old male and female BALB/c (nu/nu) mice, weighing 20.3 ± 2 g, were purchased from Laboratory Animal Center, Fourth Military Medical University (Xi'an, China).

HepG2 cells (2×10^5) were injected subcutaneously on the right side of each mouse's back to induce tumor formation. The animals were divided into three groups when the tumor volume reached 30 mm³ with 0.1 mL LV-NT4(Si)-p53(N15)-Ant (2×10^6 TU/mL) injected into their tumor. Mice treated with PBS and LV-EGFP served as controls with 0.1 mL LV-EGFP (2×10^6 TU/mL) injected into their tumor (injection at different directions). Infection efficiency of lentivirus was observed under a confocal laser scanning microscope one week after infection.

The animals were sacrificed 3 wk after injection of lentivirus. Tumors were removed and weighed with their size measured using a vernier caliper and their volume calculated according to the following formula: $(3.14 \times L \times W \times H)/6$. Tumors were fixed in 4% formaldehyde and embedded in paraffin. Tumor tissue was cut into 5-µm thick sections which were stained with HE.

Tumor growth inhibition rate = $(A-B)/A \times K$, where A is the average tumor weight of control group, B is the average tumor weight of treatment group, K represents 100%.

Statistical analysis

The data were expressed as mean \pm SD. LDH release and

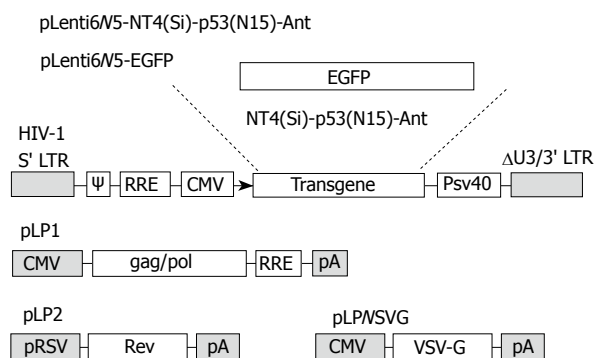


Figure 1 Packaging system of LV-NT4(Si)-p53(N15)-Ant and LV-EGFP. The packaging system: lentiviral vector, pLenti6/V5-D-TOPO, contains HIV-1 5'LTR, packaging signal (ψ), RRE sequence, CMV promoter and 3'LTR from which some regulatory sequences are deleted, resulting in "self-inactivation" of lentivirus after transduction of the target cells. pLP1, pLP2 and pLP/VSVG are three helper plasmids, which provide structural and replication proteins to produce lentivirus.

MTT were analyzed by *t*-test and χ^2 test, respectively. The data for animal experiment were processed by analysis of variance. Data analysis was performed using the SPSS 15.0. $P < 0.05$ was considered statistically significant.

RESULTS

Construction of lentivirus vector expressing NT4-p53(N15)-Ant or EGFP

The lentivirus expression vector, pLenti6 V5-D-TOPO we constructed, contains the elements required for virion packaging, such as 5' and 3' long terminal repeats and ψ packaging signal. The packaging system of LV-NT4(Si)-p53(N15)-Ant and LV-EGFP is shown in Figure 1.

The DNA sequence of the *p53(N15)-Ant* gene was identical to those of *p53(N15)* and 17 amino acid Ant^[9]. The length of NT4-p53(N15)-Ant was 349 bp, including a 247 bp fragment of the signal peptide sequence and pro-region of neurotrophin 4 with the two restriction sites (*Bam*HI and *Nae*I). The DNA fragment of *p53(N15)-Ant* was 108 bp with *Nae*I and *Xho*I. Digestion of pLenti6/V5-NT4-p53(N15)-Ant with *Bam*HI and *Nae*I followed by electrophoresis verified that the NT4-p53(N15)-Ant secretory expression cassette was correctly inserted into the multiple cloning site (MCS) of pLenti6/V5-D-TOPO, and pLenti6/V5-EGFP was confirmed in the same way as pLenti6/V5-NT4-p53(N15)-Ant.

Gene transfer and lentivirus expression in HepG2 cells

Fluorescent microscopic analysis showed that EGFP was expressed in over 60% of HepG2 cells 48 h after infection with pLenti6/V5-EGFP at a MOI of 4 TU/cell (Figure 2A and B).

Expression of NT4-p53(N15)-Ant in lentivirus-delivered H1299 cells

Since H1299 cells lack of the *p53* gene (null-*p53*), H1299 lung cancer cells were used as target cells for the expression of *p53(N15)-Ant*. *p53* protein could not be detected with immunohistochemical staining before or after infection with LV-EGFP (Figure 2C). However, 48 h

after infection with LV-NT4(Si)-*p53(N15)-Ant*, *p53* protein was mainly expressed in cytoplasm of H1299 lung cancer cells (Figure 2D). Anti-DO-1 could identify and bind to amino acids 11-25 of *p53*, suggesting that LV-NT4(Si)-*p53(N15)-Ant* can effectively infect the target cells and express the *p53* fusion peptide.

Killing effect of lentivirus-NT4 p53(N15)-Ant on HepG2 cells in vitro

HepG2 cells, infected with pLenti6/V5-NT4-p53(N15)-Ant in our study, were significantly killed, whereas pLenti6/V5-EGFP had no effect on cell viability, suggesting that growth of HepG2 cells can be inhibited by the transferred exogenous gene rather by viral toxicity (Figure 3).

Morphological changes in HepG2 cells observed under light microscope

Significant morphological changes were observed in HepG2 cells at different time points after infection with LV-NT4(Si)-*p53(N15)-Ant* but not after infection with LV-EGFP and LV-NT4(Si)-*p53(N15)-Ant* (Figure 4). However, the cells became swollen and their boundary was blurred 48 h after infection with LV-NT4(Si)-*p53(N15)-Ant*, while their viability was significantly declined with noticeable cell shrinkage, fragments and detachment 72 h after infection with LV-NT4(Si)-*p53(N15)-Ant*, but without significant morphological changes 72 h after infection with LV-EGFP infection.

Changes in ultra-structure of HepG2 cells observed under electron microscope

Forty-eight hours after infection with LV-EGFP, the membranes and nuclear membranes of HepG2 cells were intact with some crimples in the nuclear membranes, some microvilli on the surface of HepG2 cells, normal structure of mitochondria and ER (Figure 5A), and 2 completely divided HepG2 cells (Figure 5B), suggesting that LV-EGFP has no significant effect on cell growth and division. Forty-eight hours after infection with LV-NT4(Si)-*p53(N15)-Ant*, significant changes in ultra-structure of HepG2 cells, as well as in cell membranes and mitochondria, were observed under electron microscope. Cell membranes were incomplete with several fractures and the cells appeared to leak contents. Mitochondria were swollen with vacuolization. Vacuoles were also seen under the nuclear membrane with bare nuclei (Figure 5C and D). These findings suggest that NT4(Si)-*p53(N15)-Ant* may exert its anticancer effect mainly by inducing necrosis of cell membranes.

Determination of LDH release

LDH was significantly increased in LV-NT4(Si)-*p53(N15)-Ant* treatment group, but not in LV-EGFP treatment group, with no significant difference between the two groups. The longer the time was after the infection, the greater the difference was between the groups, suggesting that cell membranes are damaged.

Annexin V-PI double staining

Flow cytometry (FCM) was employed to detect the

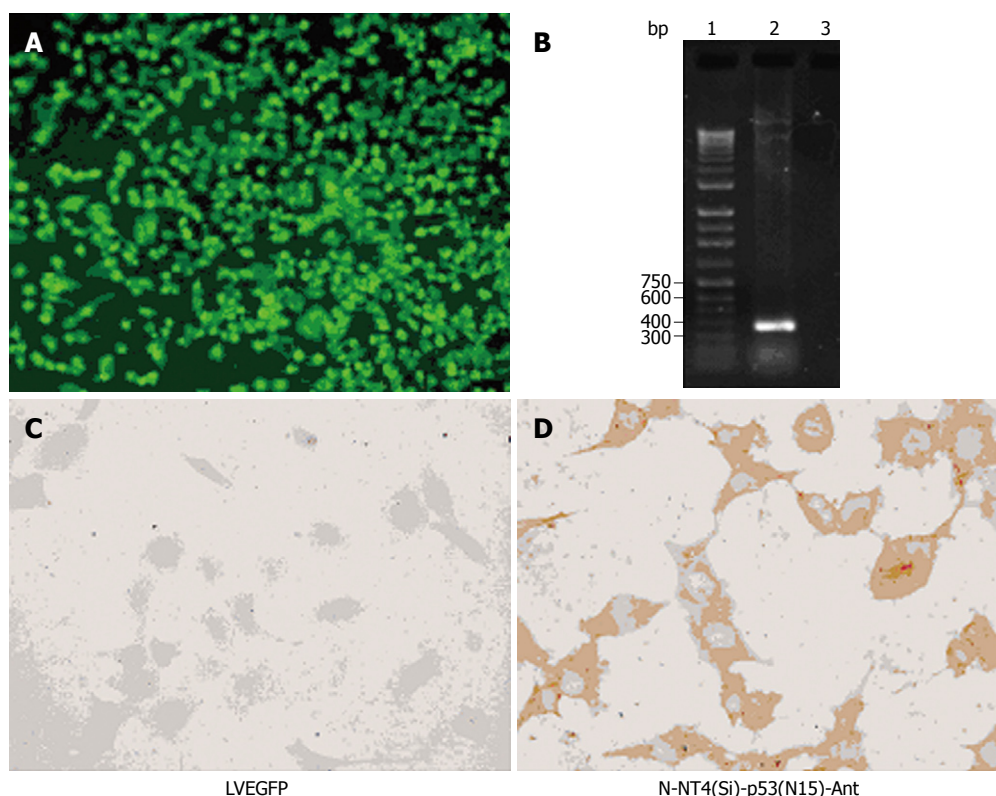


Figure 2 Fluorescent microscopic analysis and immunohistochemical staining. Fluorescence microscopy analysis showing EGFP expression in HepG2 cells (A), agarose gel electrophoresis of RT-PCR products from HepG2 cells (B) [lane 1: 1.0 kb DNA ladder; lane 2: RT-PCR products from cells infected with LV-NT4(Si)-p53(N15)-Ant; lane 3: negative controls infected with LV-EGFP], immunohistochemical staining showing no p53 protein (C), and p53 protein expression in cytoplasm of HepG2 cells (D) 48 h after infection with LV-NT4(Si)-p53(N15)-Ant.

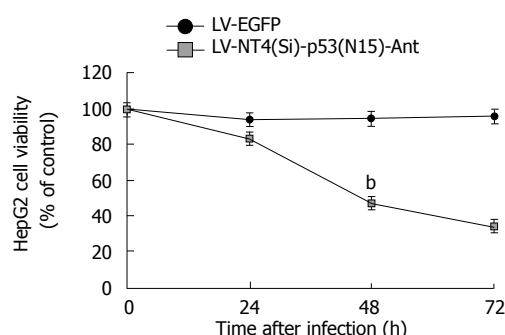


Figure 3 Cytotoxicity of LV-NT4(Si)-p53(N15)-Ant to HepG2 cells at different time points (24, 48 and 72 h) after infection with lentivirus. $^aP < 0.01$ vs LV-EGFP.

fluorescence intensity after HepG2 cells were double-stained with Annexin V and PI. A plot composed of 4 quadrants was detected by flow cytometry, which showed that the number of necrotic cells was significantly greater in LV-NT4(Si)-p53(N15)-Ant treatment group than in LV-EGFP treatment group 48 h after infection with lentivirus. The apoptosis level was also higher in cells expressing LV-p53(N15) than in controls. However, the number of necrotic cells was still much greater than that of apoptotic cells (Figure 6), suggesting that the NT4(Si)-p53(N15)-Ant gene kills tumor cells mainly by inducing necrosis of cell membrane.

Inhibition of HepG2 cell growth in nude mice by LV-NT4(Si)-p53(N15)-Ant

Nude mice developed tumor mass within 10 d after inoculation with HepG2 cells. Three weeks after infection with lentivirus, the size and weight of tumor mass were measured (Figure 7A and B). The inhibition

rate of tumor growth for LV-NT4(Si)-p53(N15)-Ant was 66.14%.

DISCUSSION

p53 protein plays an essential role in cell activities^[11,12]. Under various stress conditions, such as genotoxic damage, hypoxia, ribonucleotide depletion or oncogene activation, p53 molecule accumulates and is activated. Activated p53 acts as a transcription factor and activates or represses a variety of genes involved in cell-cycle regulation, induction of apoptosis or senescence. Therefore, the p53 gene represents an ideal target for cancer therapy. In non-stressed cells, p53 protein is present at a very low cellular concentration, because it interacts with the hdm2 protein, the human analogue of murine mdm2, which acts as a ligase of ubiquitin^[13,14] and promotes p53 degradation^[9,10]. Recent advances in regulation of p53 by hdm2 offer the possibility of generating new anticancer agents that activate wild-type p53 in tumors. Since hdm2 down-regulates p53, inhibition of this interaction would lead to an accumulation of p53 in tumor cells, eventually inducing their death^[10,15]. p53(N15)-Ant has been considered a novel cancer therapeutic peptide because it induces cancer cell death and does not seem to be cytotoxic to normal cells and may, therefore, prove useful as a general anticancer agent. In this study, we focused on how to improve the anticancer effect of peptide therapy by employing two strategies. The first was to enable the expressed therapeutic peptide to be secreted, and the second was to use the lentivirus gene transfer system. It is well-known that the signal peptide is located at the terminal of amino acid sequence of a secreted protein and plays an important role in protein targeting

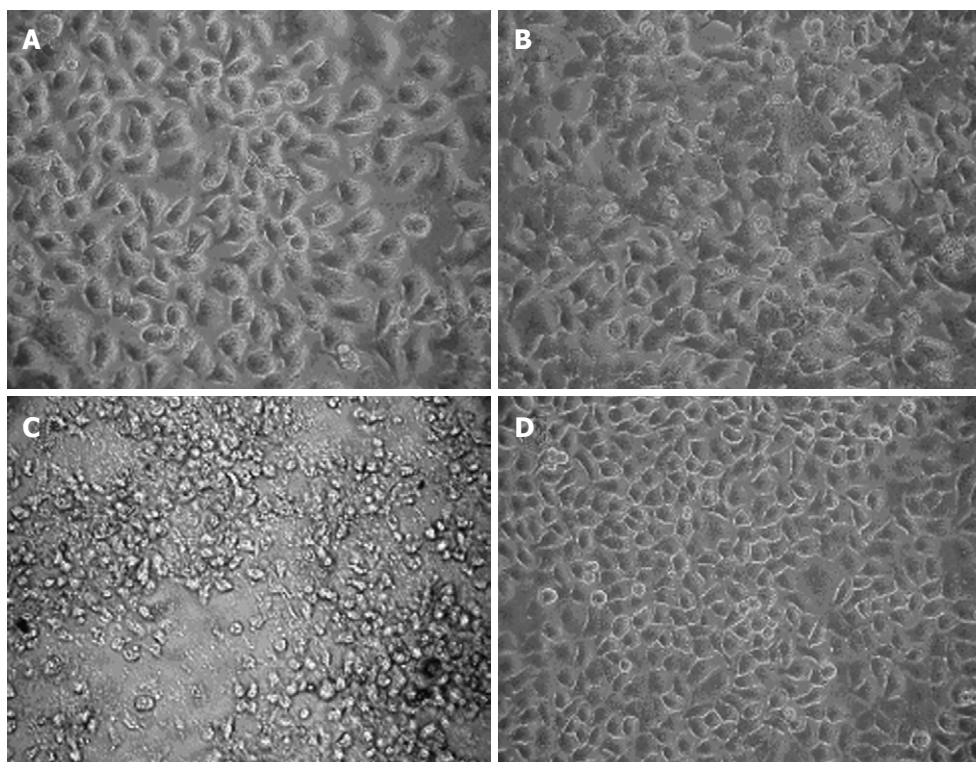


Figure 4 Morphological changes in HepG2 cells 24 h (A), 48 h (B), 72 h (C) after infection with LV-NT4(Si)-p53(N15)-Ant and 72 h (D) after infection with LV-EGFP ($\times 400$).

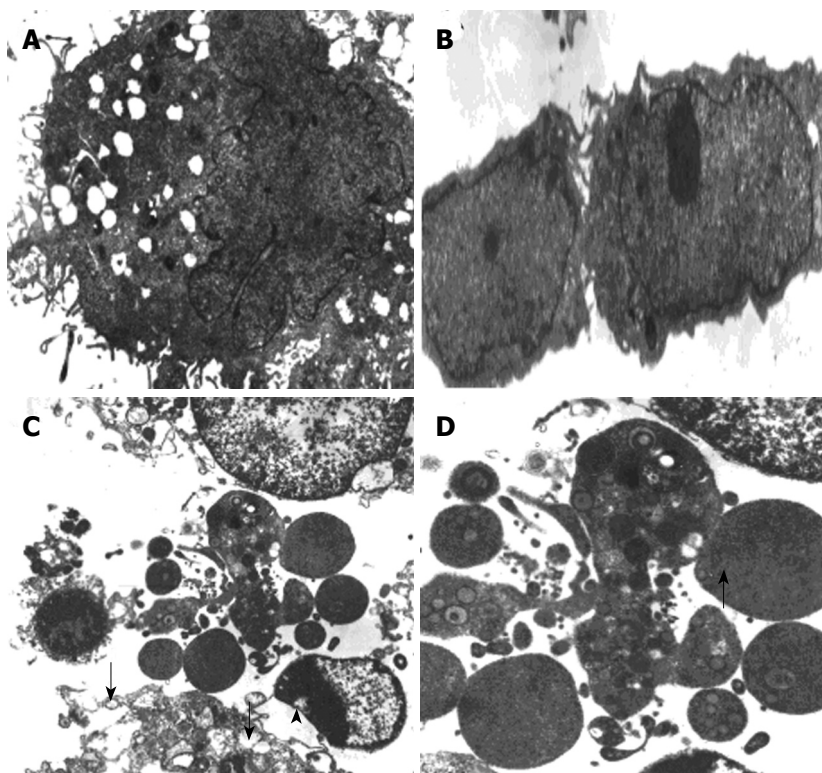


Figure 5 Ultra-structure of HepG2 cells 48 h after infection with LV-EGFP (A, B) and LV-NT4(Si)-p53(N15)-Ant (C, D) ($\times 5000$). A: Forty-eight hours after LV-EGFP infection, the cell membranes and nuclear membranes of HepG2 cells were intact; there were some crimples in the nuclear membranes and some microvilli on the surface of cells. The structure of mitochondria and ER was normal; B: Forty-eight hours after LV-EGFP infection, there were two HepG2 cells which had just completed division and this phenomenon suggested that LV-EGFP did not have a significant effect on cell growth and division; C: Forty-eight hours after LV-NT4(Si)-p53(N15)-Ant infection, it can be seen that cell membranes were incomplete and several fractures (arrows) were seen. Cell contents also appeared to be leaking. Mitochondria were swollen and showed vacuolization. Vacuoles were also seen under the nuclear membrane (arrowhead); D: Forty-eight hours after LV-NT4(Si)-p53(N15)-Ant infection, the remaining bare nuclei (arrow), after cytoplasm collapse can be seen.

and translocation in both prokaryotic and eukaryotic cells. NT4 is a member of the neurotrophin family, involved in controlling survival and differentiation of vertebrate neurons^[16]. Besides the common features of neurotrophins, NT4 has many unusual features. Unlike other neurotrophins, the expression of NT4 is ubiquitous and less influenced by environmental signals^[17]. In the present study, NT4 structure analysis showed that the human NT4 initiation codon was followed by a signal

peptide sequence, a pro-region and an apparent dibasic cleavage site, which was followed by the sequence of the mature NT4. The signal peptide sequence and pro-region of human NT4 are notably shorter (by = 243 bp) than those of other neurotrophins^[18]. In our previous study, the pre-protein NT4 could be cleaved into the signal peptide and the mature protein by *Escherichia coli* (*E. coli*) endopeptidase. There is a native NaeI enzyme site in front of the site in the NT4 signal peptide,

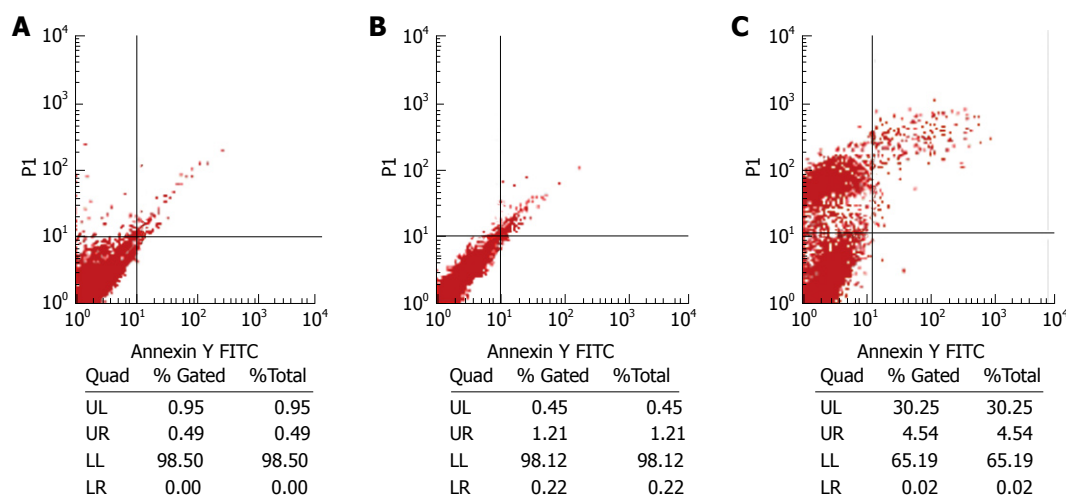


Figure 6 Flow cytometry analysis of HepG2 cells without treatment (A), 48 h after infection with LV-EGFP (B) and LV-NT4(Si)-p53(N15)-Ant (C) after stained with Annexin V and propidium iodide. The left lower quadrant represents normal cells (An-PI-), the right lower quadrant represents early apoptotic cells (An+ PI-), the right upper quadrant represents apoptotic cells and necrotic cells (An + PI +), the upper left quadrant represents early necrotic cells (An-PI +).

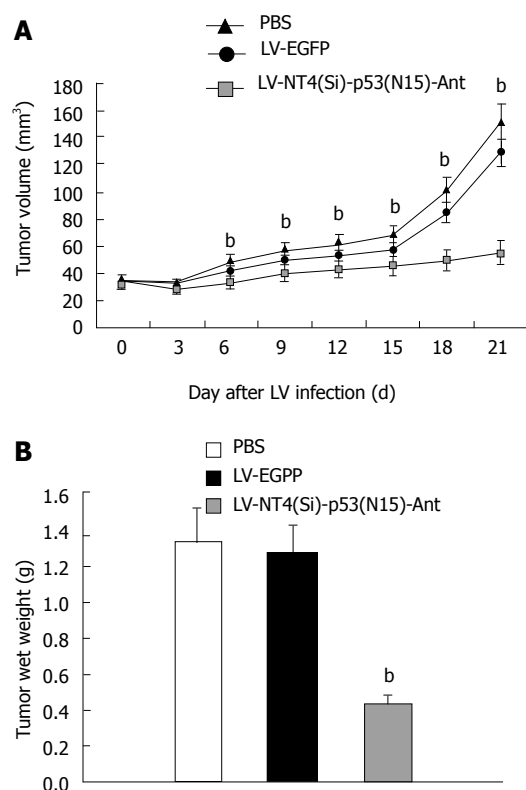


Figure 7 Tumor growth curve (A) and tumor wet weight (B) after treatment with LV-NT4(Si)-p53(N15)-Ant. A: Tumor growth curve after treatment with LV-NT4(Si)-p53(N15)-Ant. Tumors grew rapidly in the PBS and LV-EGFP groups, while the tumor growth was significantly inhibited in the LV-NT4(Si)-p53(N15)-Ant group 3 wk after infection. There was no statistical significance between the PBS group ($151.0 \pm 50.1 \text{ mm}^3$) and the LV-EGFP group ($128.5 \pm 45.3 \text{ mm}^3$) in terms of tumor volume ($P > 0.05$). But the tumor volume in the LV-NT4(Si)-p53(N15)-Ant group ($55.1 \pm 18.7 \text{ mm}^3$) was significantly less than that of the PBS and LV-EGFP groups ($^bP < 0.01$); B: Tumor wet weight after treatment with LV-NT4(Si)-p53(N15)-Ant. There were no significant differences in the wet weight of tumors between the two control groups: PBS group $1.33 \pm 0.93 \text{ g}$ vs LV-EGFP group $1.27 \pm 1.04 \text{ g}$ ($P > 0.05$). But the wet weight of the LV-NT4(Si)-p53(N15)-Ant treatment group ($0.43 \pm 0.77 \text{ g}$) was significantly less than that of the two control groups ($^bP < 0.01$).

convenient for the expression of other exogenous peptides in a secretory manner. It has been shown that it is not only feasible, but also improves the therapeutic effect^[19]. In this study, we successfully constructed a LV containing the NT4(Si)-p53(N15) fusion gene with *in vitro* recombinant DNA technology.

To transfer the NT4(Si)-p53(N15)-Ant gene into the target cells and to achieve a consistent expression, selection of the gene transfer vector is a critical step. Lentiviruses, such as HIV and SIV, are a new generation of vectors and have many advantages over the traditional virus vectors, such as retroviral or adenovirus vector. These LVs can infect both dividing and non-dividing cells and efficiently integrate into the host DNA, thus providing stable transgene expression over several months^[20,21]. At the same time, LV is very safe because all lentivirus accessory genes have been removed, virus production components have been split into 3 or 4 separate parts, and self-inactivating deletions have been introduced into the vector^[21,22]. Since the aim of cancer gene therapy is to remove all cancer cells including those in a precancerous stage, only extensive gene transfer and long-term gene expression in cancer cells are appropriate. Thus, LV has a greater promise to become an ideal vector for gene delivery.

In this study, MTT assay showed that LV-NT4(Si)-p53(N15)-Ant could significantly induce HepG2 cell death 48 h after infection, LDH was significantly increased in the LV-NT4(Si)-p53(N15)-Ant treatment group but not in the LV-EGFP treatment group, suggesting that the cell membranes are damaged. Moreover, cytotoxicity occurred in a time-dependent manner. The ultra-structure of HepG2 cells, especially cell membranes and mitochondria, was significantly changed after infection with LV-NT4(Si)-p53(N15)-Ant, which was different from that during apoptosis and necrosis. In addition, FCM revealed that two cell death modes (apoptosis and necrosis) occurred 48 h after infection with LV-NT4(Si)-p53(N15)-Ant with necrosis being common and apoptosis being rare, revealing that LV-NT4(Si)-p53(N15)-Ant exerts its effect on HCC mainly by inducing necrosis of cell membranes.

thus ensuring the correct cleavage of therapeutic peptide, and making this signal peptide sequence highly

This is probably related to the process of LV infection, integration of the therapeutic gene into the host genome and gene expression.

The *in vivo* study also showed that LV-NT4(Si)-p53(N15)-Ant significantly inhibited tumor growth. Since p53(N15) contains overlapping sequences from the p53 mdm-2 binding domain, p53(N15)-Ant peptide may block the interaction of mdm-2 with other proteins. Recently, a study investigating the secondary structure of the p53 fusion peptide has revealed that p53(N15)-Ant peptide plays an important role in membrane interaction^[15], showing that p53(N15)-Ant peptide forms distinctive S-shape helix-loop-helix structures, which can rapidly disrupt cancer cell membranes by forming a toroidal-like pore, resulting in necrosis. The two mechanisms exist simultaneously.

In conclusion, we have established a way to express lentivirus-introduced p53(N15)-Ant 32-peptide. The growth of HepG2 cells can be significantly inhibited by LV-NT4(Si)-p53(N15)-Ant. LV-NT4(Si)-p53(N15)-Ant gene therapy may be used as a novel anticancer strategy. Further study on the biological characteristics of the p53 peptide is needed by transfecting LV-NT4(Si)-p53(N15)-Ant into other cancer cell lines.

COMMENTS

Background

Peptide has emerged as a new anticancer agent in recent years. Numerous reports suggest that many low molecular weight peptides possess an anticancer effect but have almost no cytotoxic effect on normal cells. p53(N15)-Ant has been considered a novel cancer therapeutic peptide because it induces cancer cell death and does not seem to be cytotoxic to normal cells.

Research frontiers

It has been reported that p53 peptide, synthesized from residues 12-26 and fused with *Drosophila* carrier protein antennapedia (Ant), induces rapid tumor cell necrosis in all breast and pancreatic cancer cell lines tested, irrespective of the p53 status, whereas it shows a low cytotoxicity to normal cells. In this study, the authors constructed a novel recombinant lentivirus expression plasmid LV-NT4(Si)-p53(N15)-Ant and demonstrated its anticancer effect.

Innovations and breakthroughs

The signal peptide plays an important role in protein targeting and translocation in both prokaryotic and eukaryotic cells. In this experiment, the authors used the NT4 signal peptide to enable the therapeutic peptide to be secreted. Lentivirus vectors were also employed. LV-NT4(Si)-p53(N15)-Ant was constructed and successfully cultured at a high titer and verified to induce necrosis of cancer cells.

Applications

This work has established a way to express lentivirus-introduced p53(N15)-Ant 32-peptide. LV-NT4(Si)-p53(N15)-Ant dependent gene therapy may be a promising novel anticancer strategy. p53(N15)-Ant may represent a promising agent of gene therapy for hepatocellular carcinoma.

Terminology

NT4 signal peptide and its pro-region are protein sequences that can direct the secretion of proteins and peptides from cells. Such a mechanism is thought to be helpful in fusion peptide gene therapy as it can enhance the therapeutic effect.

Peer review

In this study, Song *et al* showed that the constructed lentivirus vector expressed in cancer cells could inhibit the growth of HepG2 cells. This vector can induce necrosis and apoptosis, and inhibit tumor growth *in vivo*. The study in general is significant and interesting.

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