

# Lentivirus-mediated RNAi knockdown of *LMP2A* inhibits the growth of the Epstein-Barr-associated gastric carcinoma cell line GT38 *in vitro*

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**Abstract.** In this study, lentivirus-mediated RNA interference (RNAi) was applied to inhibit latent membrane protein 2A (*LMP2A*) gene expression, in order to explore the effects of *LMP2A* silencing on the growth of an Epstein-Barr virus-associated gastric carcinoma (EBVaGC) cell line *in vitro*. Lentivirus-mediated RNAi technology was employed to specifically knock down the *LMP2A* gene in the EBV-positive gastric carcinoma cell line GT38. After infection, reverse transcription-quantitative polymerase chain reaction, western blotting, flow cytometry and colony formation assays were conducted to evaluate the expression of *LMP2A* and the biological behavior of the GT38 cell line *in vitro*. The results showed that the expression of the *LMP2A* gene was clearly downregulated in the infected cells, which indicated that a highly efficient and stable lentivirus vector was successfully constructed. In the GT38 cells in which the expression of *LMP2A* was downregulated, the proliferation and colony formation of the cells was significantly inhibited. In addition, it was found that the cell cycle of the GT38 cells was arrested in the G0/G1 phase and the apoptosis rate was increased. These results indicate that lentivirus-mediated RNAi knockdown of *LMP2A* inhibits the growth of the EBVaGC cell line GT38 *in vitro*, and suggests that *LMP2A* is a potential target for gene therapy in the treatment of EBVaGC.

## Introduction

Epstein-Barr virus (EBV) is a human gamma herpes virus that can exist in humans for a long time without producing any symptoms (1). A variety of human malignancies such as Burkitt's lymphoma, nasopharyngeal carcinoma and gastric cancer (GC) are reported to be associated with EBV (2). EBV-associated gastric carcinoma (EBVaGC) accounts for 10-18% of gastric carcinoma cases, and there are estimated to be >75,000 new cases of EBVaGC each year worldwide (3,4). As a type of GC, which is one of the most common types of malignant tumors, EBVaGC is very difficult to remove and the complete elimination of tumor cells by surgical, radiotherapeutic and chemotherapeutic methods is challenging. Therefore, the development of new therapeutic approaches for the inhibition of tumor cell growth or survival to treat EBVaGC is of critical importance.

EBVaGC is closely associated with EBV infection, and EBVaGC cells have been found to express a well-defined set of latent viral genes, including latent membrane protein 2A (*LMP2A*) (5). It has been revealed that ~40% EBVaGC expresses *LMP2A*, whose expression closely correlates with a poor survival outcome (6). *LMP2A* consists of a 27-amino-acid carboxy-terminal cytoplasmic domain, a 119-amino-acid tyrosine-rich amino-terminal cytoplasmic domain and 12 hydrophobic transmembrane domains, with the cytoplasmic domain playing a role as a signaling effector (7). *LMP2A* has various functions, one of which is to activate the phosphoinositide 3-kinase (PI3K)/Akt, nuclear factor (NF)- $\kappa$ B,  $\beta$ -catenin, signal transducers and activators of transcription (STAT) and Syk tyrosine kinase pathways in epithelial cells (8-10). *LMP2A* also plays an important role in cell transformation activities, such as the induction of cell growth, enhancement of cell adhesion and cell motility, as well as the inhibition of epithelial cell differentiation (11,12). Thus, the aforementioned findings indicate that *LMP2A* may be a potential target for gene therapy for EBVaGC treatment.

RNA interference (RNAi) is an efficient tool that can cause post-transcriptional silencing of gene expression and induce

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loss-of-function phenotypes (13). Lentivirus vectors have been developed to be a powerful technology for the achievement of a significant level of gene transfer *in vitro* (14). In the present study, lentivirus-mediated RNAi was used to inhibit *LMP2A* gene expression, and the effects of *LMP2A*-silencing on cell growth, cell cycle and cell apoptosis of EBVaGC cell line GT38 were studied.

## Materials and methods

**Cell line and culture.** The EBV-positive human gastric carcinoma cell line GT38 was bought from American Type Culture Collection (Manassas, VA, USA). Cells were cultured in RPMI-1640 (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 10% fetal bovine serum (FBS; ExCell Bio, Shanghai, China), 50 U/ml penicillin G and 50 U/ml streptomycin (Gibco) in a humidified atmosphere of 5% CO<sub>2</sub> at 37°C. The medium was changed every 2 days and the cell line was passaged every 4-5 days.

**Construction of lentivirus vectors.** In order to produce lentivirus expressing RNAi specific for the *LMP2A* gene (GU979791), RNAi Designer software (rnaidesigner.thermo-fisher.com/rnaiexpress/; BLOCK-iTTM RNAi; Thermo Fisher Scientific, Inc.) was used to identify the RNAi sequence for human *LMP2A* (CTC CCA ATA TCC ATC TGC T), and then a scrambled sequence (TTC TCC GAA CGT GTC ACG T) was created as a negative control construct (control RNAi) that should have no homology with the human genome. DNA oligos with the target sequence were chemically synthesized, annealed, double digested with *Age*I and *Eco*RI, and inserted into the pLenR-GPH expression vector (GeneChem Co., Ltd., Shanghai, China) using T4 DNA ligase (Invitrogen; Thermo Fisher Scientific, Inc.), following the manufacturer's guidelines. The ligated vector was transformed into competent *Escherichia coli* DH5a cells (Invitrogen; Thermo Fisher Scientific, Inc.). Restriction enzyme analysis and DNA sequencing were performed to identify the correct transformant. The sequences were cloned into the pGCSIL-Green Fluorescent Protein (GFP) vector (GeneChem Co., Ltd.) to generate lentivirus vectors. 293T cells (Shanghai Research Institute of Chinese Academy of Sciences, Beijing, China) were used to generate lentiviruses after their transfection into the expression vectors and package vectors with the use of Lipofectamine 2000 (Thermo Fisher Scientific, Inc.). After 48 h, supernatants containing the lentiviruses pGCSIL-LMP2A-shRNA-LV and pGCSIL-neg-shRNA-LV were harvested and the remaining cells were removed by filtering with 0.45 μm filters. Ultracentrifugation (4,000 × *g* at 4°C for 10 min) was then performed to concentrate the lentiviruses and the titer was finally determined by 293T cell infection assay.

**Infection of lentivirus.** In this assay, 5×10<sup>3</sup> GT38 cells in the logarithmic growth phase were seeded in each well of a 96-well microplate and cultured overnight. The lentiviruses were then diluted with 0.2 ml RPMI complete medium containing Polybrene (10 μg/ml) and added to infect the seeded cells for 12 h at 37°C. The virus-containing medium was then changed with fresh culture medium. Fluorescence

microscopy (IX-53; Olympus Corporation, Tokyo, Japan) was used to detect GFP in the successfully infected cells, and the percentage of GFP-positive cells was used to measure the infection efficiency of the cells. At 5 days after the infection, analysis of *LMP2A* expression, cell proliferation and cell apoptosis was performed. GT38 cells treated differently were divided into three groups in subsequent assays: Blank control group (CON group; cells without infection), negative control (NC group; cells were infected with pGCSIL-neg-shRNA-LV) and the *LMP2A* knockdown group (KD group; cells were infected with pGCSIL-LMP2A-shRNA-LV).

**Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) analysis.** Total RNA was extracted from cells using TRIzol reagent (Invitrogen, Shanghai, China) and reverse transcribed using the SuperScript III First-strand Synthesis System kit (cat. no. 18080051; Invitrogen; Thermo Fisher Scientific, Inc.), according to the manufacturer's instructions. PCR amplification was carried out using Platinum SYBR Green qPCR SuperMix UDG (Invitrogen). The reaction volume contained DNA Polymerase 10X buffer (2 μl), MgCl<sub>2</sub> (1.5 mM), dNTP mix (2.5 mM), 10 pmol each primer, DNA template (1 μg), Taq DNA Polymerase (2-3 μl) and nuclease-free water to a final volume of 20 μl. The integrity of RNA was detected by the parallel amplification of glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) mRNA. The specific primer pairs were as follows: *LMP2A* (280 bp), sense: ATGACTCATCTC AACACATA and antisense: CATGTTAGGCAAATTGCAA; *GAPDH* (450 bp), forward: 5'-CTCAGACACCATGGGGAA GGTGA-3' and reverse: 5'-ATGATCTTGAGGCTGTTGTCA TA-3'. The following thermal cycling conditions were used: 95°C for 5 min, followed by 40 cycles of 95°C for 15 sec, 60°C for 15 sec and 72°C for 1 min, and the final extension was 72°C for 5 min. The 2<sup>-ΔΔC<sub>q</sub></sup> method was applied to analyze the data (15).

**Western blot analysis.** Protein was extracted from the cells using lysis buffer (150 mM NaCl, 0.1% SDS, 0.5% sodium deoxycholate, 1% Nonidet P-40, and 50 mM Tris, pH 8.0), with the addition of 2 mM phenylmethylsulfonyl fluoride. The six-well plates of lentivirus-infected cells were put on ice, freshly prepared lysis buffer (100 μl/well) was added and the cells were incubated for 10 min. A protein assay kit was used to determine the protein concentration. Equal amounts of protein samples (10 μg per condition) were prepared in loading buffer and boiled for 10 min, then the boiled samples were loaded to 15% SDS-PAGE gels and separated. Separated proteins were transferred from the gel to polyvinylidene difluoride (PVDF) membranes at 100 V for 1 h. The PVDF membrane with protein samples was blocked using blocking buffer which was freshly prepared with PBS containing 5% skimmed milk powder. After blocking, anti-LMP2A (cat. no. ab59026; Abcam, Cambridge, UK) and anti-GAPDH (cat. no. 5174; Cell Signaling Technology, Inc., Danvers, MA, USA) primary antibodies and anti-rabbit horseradish-peroxidase (HRP)-conjugated IgG (cat. no. 7071; Cell Signaling Technology, Inc.) and anti-mouse HRP-conjugated IgG (cat. no. 7072; Cell Signaling Technology, Inc.) secondary antibodies were diluted to 1:1,000 concentration with PBS and Tween 20 (PBST) buffer before they were successively incubated with the PVDF membrane

for 60 min at room temperature. Prior to each step, the PVDF membrane was washed 3 times with PBST. Protein bands were then detected using enhanced chemiluminescence (Pierce ECL Substrate; Thermo Fisher Scientific, Inc.), and gel analysis was performed using Image J software (version 1.8.0; National Institutes of Health, Bethesda, MD, USA).

**Proliferation assay.** Cell proliferation was measured using a Cell Counting kit-8 (CCK-8) assay (Dojindo Molecular Technologies, Inc., Kumamoto, Japan). After lentivirus-infected cells had been seeded in 96-well plates for 20, 44, 68, 92 or 116 h, 10  $\mu$ l CCK-8 solution was added to each well and incubated for 1 h at 37°C. Absorbance was measured at 490 nm using an enzyme immunoassay analyzer (model 680; Bio-Rad Laboratories, Inc., Hercules, CA, USA).

**Colony formation assay.** A colony formation assay was performed to analyze the effect of *LMP2A* silencing on the colony formation ability of GT38 cells. In this assay,  $8 \times 10^2$  cells were seeded in 6-cm dishes and cultured with RPMI-1640 supplemented with 10% FBS in an atmosphere of 5% CO<sub>2</sub> and 95% humidity at 37°C. After 2 weeks, the cell colonies were washed twice with PBS, and were fixed with 4% paraformaldehyde for 15 min. The fixed colonies were then stained with Giemsa for 20 min, and washed twice with ddH<sub>2</sub>O. The colonies that consisted of  $\geq 50$  cells were manually counted.

**Cell cycle analysis.** Cells were harvested with 0.25% trypsin, fixed with 70% ethanol, and then resuspended in 20 mg/ml propidium iodide (PI). Flow cytometry (FCM) using a FACSCalibur™ flow cytometer (BD Biosciences, Franklin Lakes, NJ, USA) was used to detect DNA content, and the FCM data were used to determine the relative proportions of cells in the individual cell-cycle phases.

**Apoptosis assay.** In order to differentiate intact cells from apoptotic cells, the cells were stained with Annexin V-fluorescein isothiocyanate (FITC) and PI. Briefly, a total of  $1.0 \times 10^6$  cells were washed twice with ice-cold PBS and put into binding buffer (5  $\mu$ l Annexin V-FITC and 5  $\mu$ l PI) for a 30 min incubation. Finally, FCM was used to analyze the Annexin V-FITC/PI staining.

**Statistical analysis.** All experiments were performed in triplicate, and data are shown as the mean  $\pm$  standard deviation where applicable. Statistically significant differences between the control and treatment groups were determined by one-way analysis of variance followed by Tukey's test using SPSS 17.0 software (SPSS, Inc., Chicago, IL, USA), and  $P < 0.05$  was considered to indicate a statistically significant result.

## Results

**Successfully construction of *LMP2A*-RNAi-LV.** A lentivirus vector system derived from HIV-1 was used to knock down the *LMP2A* gene by expressing short hairpin RNAs (shRNAs) directed against *LMP2A*. pGCSIL-*LMP2A*-shRNA-LV and pGCSIL-neg-shRNA-LV were constructed successfully, carrying GFP as a reporter gene. When GT38 cells were infected once with the constructed lentiviruses, >95% of the

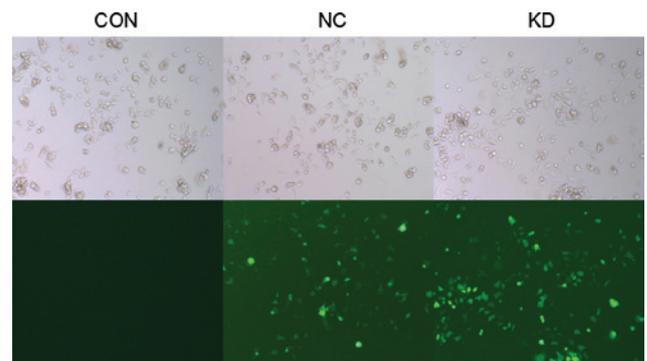


Figure 1. Lentiviral vectors encoding *LMP2A* RNAi or control RNAi were constructed and transfected into GT38 cells, with untreated GT38 cells serving as a blank control. The cells were observed using bright-field and fluorescence microscopy (magnification,  $\times 100$ ). *LMP2A*, latent membrane protein 2A; RNAi, RNA interference; CON, blank control; NC, negative control; KD, *LMP2A* knockdown.

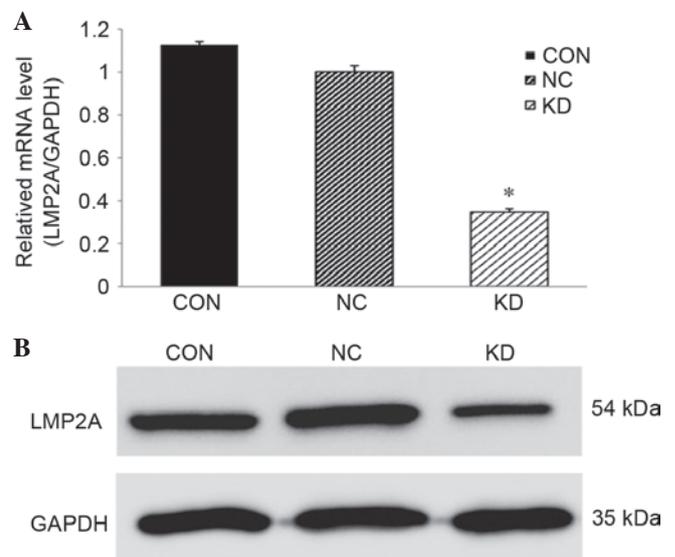


Figure 2. mRNA and protein levels of *LMP2A* in GT38 cells infected with *LMP2A*-RNAi-LV. (A) The mRNA level of *LMP2A* was detected by RT-qPCR [the  $2^{-\Delta\Delta Ct}$  method (14) was applied to indicate the relative mRNA level of the NC and KD groups, compared with the CON group]. The mRNA levels of *LMP2A* in the KD group were significantly reduced compared with that of the NC group ( $P < 0.05$ ). Data are presented as the mean plus standard deviation ( $n=6$ ). (B) The protein level of *LMP2A* was assessed by western blotting. *LMP2A*, latent membrane protein 2A; RNAi, RNA interference; CON, blank control; NC, negative control; KD, *LMP2A* knockdown.

infected GT38 cells expressed GFP 72 h after the infection (Fig. 1), indicating that highly efficient and stable lentivirus vectors had been successfully constructed.

***LMP2A*-RNAi-LV downregulates *LMP2A* expression in GT38 cells.** The expression of *LMP2A* in GT38 cells was examined by RT-qPCR and western blot analysis following infection with *LMP2A*-specific RNAi-expressing lentivirus. Compared with the NC group, the mRNA and protein levels of *LMP2A* in the KD group were significantly reduced ( $P < 0.05$ ,  $n=6$ ; Fig. 2). The mRNA level was decreased by 65.4%, while the protein level was reduced by 50.8%. The CON group exhibited no significant difference compared with the NC group. These

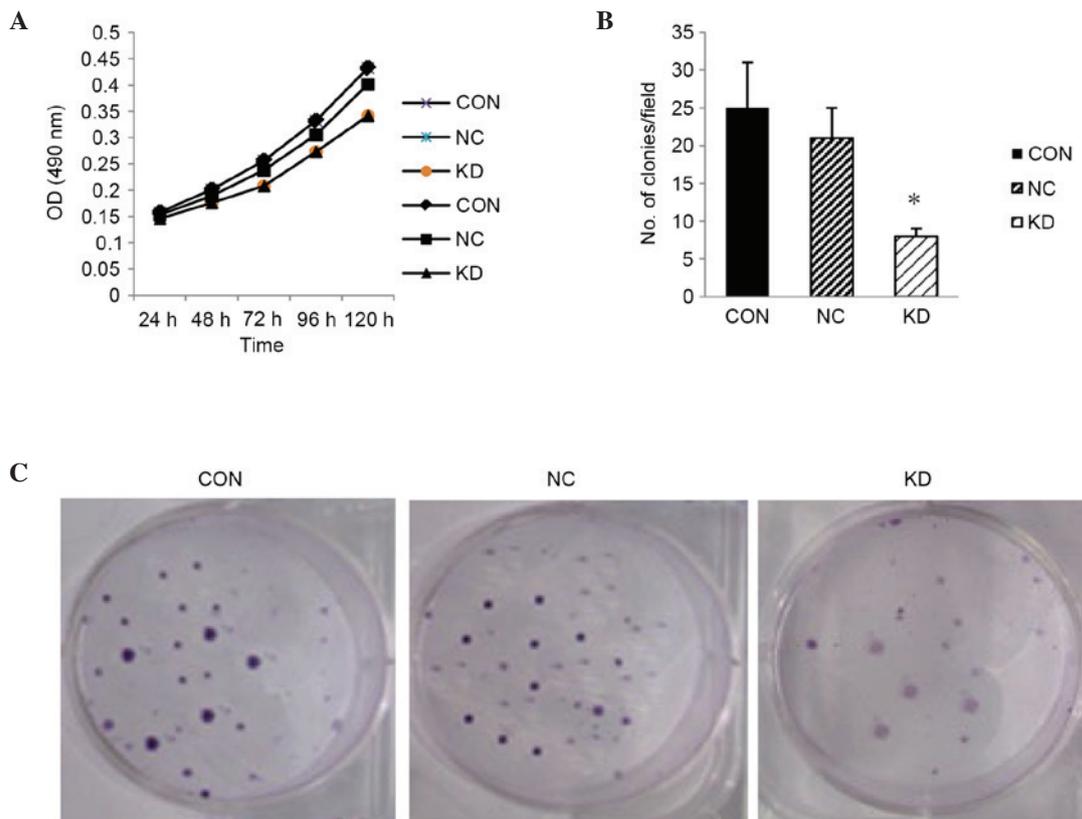


Figure 3. Suppression of cell growth in *LMP2A* RNAi-treated GT38 cells. (A) Cell proliferation was determined by CCK-8 assay. Cell proliferation in the KD group was inhibited at each time point (24, 48, 72, 96 and 120 h) compared with the CON and NC groups. (B) Colony formation assay was performed, and the cell proliferation inhibitory effect of *LMP2A* RNAi was confirmed ( $P < 0.05$  vs. CON and NC), indicating that the clone number in the KD group was greatly reduced after *LMP2A* RNAi. Data are presented as the mean plus standard deviation ( $n=6$ ). (C) Colonies of GT38 cells that were cultured in 6-cm dishes for 2 weeks and stained with Giemsa. *LMP2A*, latent membrane protein 2A; RNAi, RNA interference; CCK-8, Cell Counting kit-8; CON, blank control; NC, negative control; KD, *LMP2A* knockdown; OD, optical density.

data show that the expression of the *LMP2A* gene in GT38 cells was efficiently downregulated following the lentivirus infection.

**Knockdown of *LMP2A* inhibits the proliferation and clonogenicity of GT38 cells.** To determine the effect of downregulation of *LMP2A* on cell growth *in vitro*, a cell proliferation assay was performed after *LMP2A* RNAi using the CCK-8 method. The results show that cell proliferation in the KD group was inhibited (Fig. 3A), compared with that in the CON and NC groups. A colony formation assay was then performed to confirm the inhibitory effect of *LMP2A* RNAi. The results show that when compared with the CON and NC groups, the colony formation ability of *LMP2A* RNAi cells was significantly decreased ( $P < 0.05$ ,  $n=6$ ; Fig. 3B and C), suggesting that knockdown of the *LMP2A* gene results in the inhibition of cell growth.

**GT38 cells are arrested in the G0/G1 phase after knockdown of *LMP2A*.** FCM analysis was conducted to evaluate whether downregulation of *LMP2A* affects the cell cycle of GT38 cells. The results show an increase in the proportion of cells in the G0/G1 phase in the KD group compared with the NC group ( $66.35 \pm 1.18$  vs.  $51.19 \pm 1.07\%$ , respectively;  $P < 0.05$ ,  $n=6$ ; Fig. 4). Compared with the control group, a greater proportion of *LMP2A* RNAi cells remained in the G0/G1 phase while

fewer cells were in the S phase. The results indicate that downregulation of *LMP2A* expression in GT38 cells arrests their cell cycle in the G0/G1 phase.

**Knockdown of *LMP2A* induces apoptosis of GT38 cells.** FCM analysis was applied to evaluate the rate of apoptosis, aiming to further investigate the effects of downregulation of *LMP2A* expression on the cell death of GT38 cells. Apoptosis rates in the CON and NC groups were  $15.89 \pm 0.41$  and  $17.15 \pm 0.39\%$ , respectively, while the apoptosis rate of the KD group was  $29.89 \pm 0.48\%$ ; the percentage of apoptosis in the KD group was significantly increased compared with that in the CON and NC groups ( $P < 0.05$ ,  $n=6$ ; Fig. 5). These data indicate that downregulation of *LMP2A* significantly increases the apoptosis rate of GT38 cells compared with that in the control group, demonstrating that suppression of cell proliferation may be caused by cell death resulting from *LMP2A* RNAi.

## Discussion

*LMP2A* has been detected in the majority of EBVaGC samples (16). EBVaGC has been reported constitute 10-18% of gastric carcinoma cases and has a very poor survival outcome (17). *LMP2A* has been found to play important roles in epithelial cells, such as the enhancement of cell adhesion and cell motility, inhibition of epithelial cell differentiation

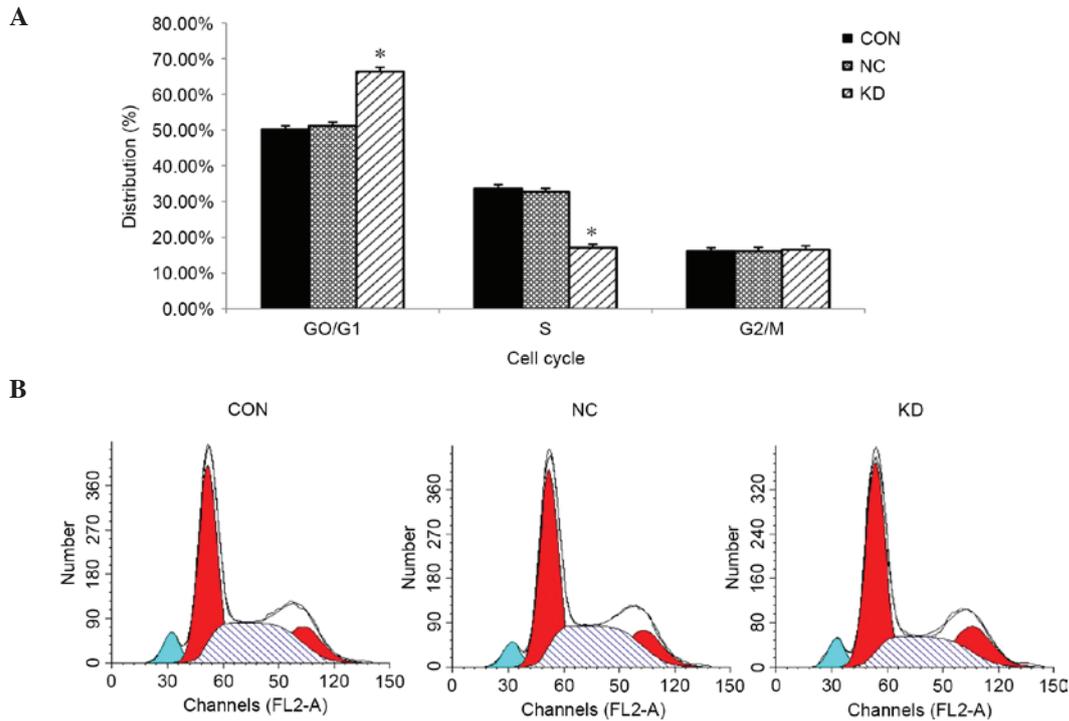


Figure 4. *LMP2A* RNAi treatment changed the cell cycle distribution of GT38 cells. (A) Proportion of cells in the G0/G1 phase significantly increased while the proportion in the S phase decreased in the KD group after treatment with *LMP2A* RNAi, compared with the CON and NC groups (\* $P < 0.05$ ). Data are presented as the mean plus standard deviation (n=6). (B) FCM analysis showed that *LMP2A* RNAi arrests GT38 cells in the G0/G1 phase of the cell cycle. *LMP2A*, latent membrane protein 2A; RNAi, RNA interference; FCM, flow cytometry; CON, blank control; NC, negative control; KD, *LMP2A* knockdown.

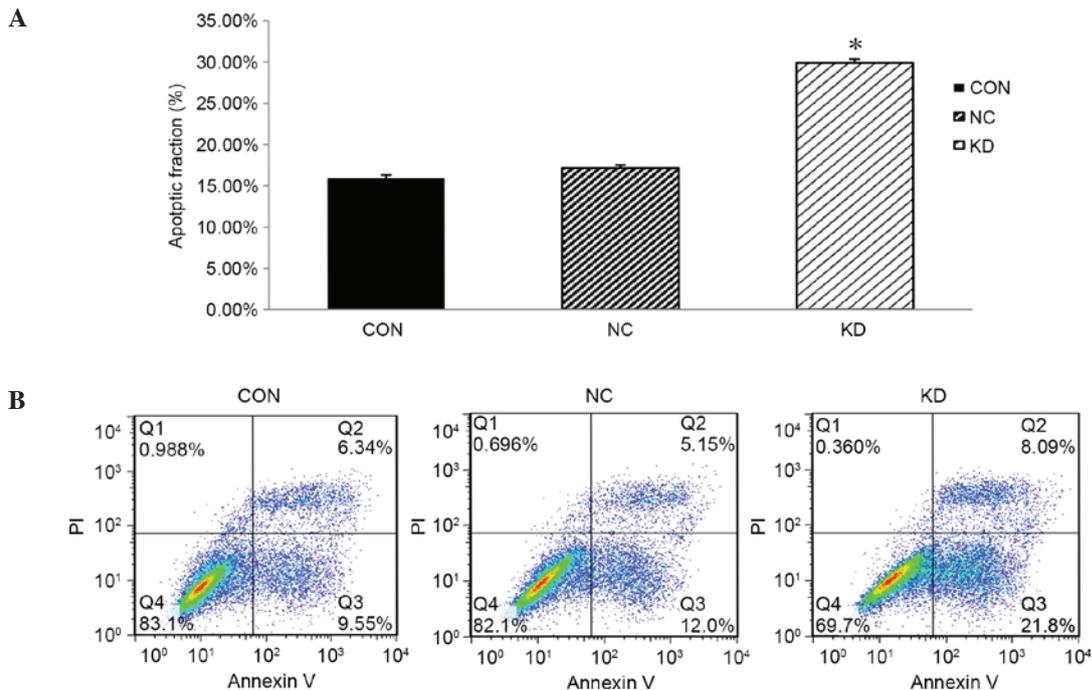


Figure 5. *LMP2A* RNAi treatment induced GT38 cells into apoptosis which was detected by FCM using Annexin V-FITC/PI double staining. (A) Compared with the CON and NC groups, *LMP2A* RNAi significantly promoted cell apoptosis in the KD group (\* $P < 0.05$ ). Data are presented as the mean plus standard deviation (n=6). (B) FCM data show that the apoptotic rate of cells with downregulation of *LMP2A* is significantly increased. *LMP2A*, latent membrane protein 2A; RNAi, RNA interference; FCM, flow cytometry; FITC, fluorescein isothiocyanate; PI, propidium iodide; CON, blank control; NC, negative control; KD, *LMP2A* knockdown.

and induction of anchorage-independent growth; these functions are reported to be correlated with the activation of the PI3K/Akt, NF- $\kappa$ B,  $\beta$ -catenin, STAT and Syk tyrosine kinase

pathways (18-20). In addition, *LMP2A* is also reported to be involved in the regulation of the proliferation, apoptosis and self-renewal ability of cancer stem cells (21). Through the

findings outlined above, *LMP2A* can be considered as a potential target of gene therapy for EBVaGC treatment. However, the detailed mechanism by which *LMP2A* expression down-regulation inhibits EBVaGC cell growth has not been clarified.

In the present study, an *LMP2A*-specific RNAi-expressing lentivirus was constructed and used to infect the human gastric carcinoma cell line GT38. This was conducted to downregulate *LMP2A* expression, in order to investigate the effects of *LMP2A* silencing on cell growth, the cell cycle and cell apoptosis of EBVaGC cells. The results indicate that when *LMP2A* is downregulated, the proliferation of GT38 cells is significantly decreased, while the cell apoptosis is markedly increased, and a greater proportion of GT38 cells are arrested in the G0/G1 phase of the cell cycle. The inhibition of cell growth may be caused by cell apoptosis resulting from *LMP2A* RNAi. Although, the mechanisms by which downregulation of the *LMP2A* gene results in growth inhibition and apoptosis promotion in the GT38 cell line are not clear, there is evidence that when *LMP2A* is activated, gastric carcinoma cells would be protected from apoptosis and their proliferating ability promoted, all of which are regulated by a variety of signaling pathways (22-25) Thus, it is hypothesized that *LMP2A* gene silencing suppresses cell proliferation of the EBVaGC cell line GT38 following such mechanisms, and *LMP2A* can be considered as a potential therapeutic target in the treatment of EBVaGC. For further investigation, future studies may be conducted to attempt to clarify the mechanisms by which *LMP2A* modulate those pathways to affect cell biological functions, which may indicate new methods for creating novel therapies against EBVaGC.

RNAi has become a powerful tool for molecular biological research with efficient ability for gene functional analysis, and also can be considered as a potential therapeutic strategy for various diseases, including cancers (26). RNAi is considered to be a novel method for gene therapy against cancer, particularly when combined with lentivirus. The exploration of target genes that when downregulated could efficiently inhibit EBVaGC cell growth may increase the possibility of clinical application of gene-silencing therapy. During the present study, a highly efficient and stable lentivirus vector system was successfully constructed, which efficiently knocked down the *LMP2A* gene in the infected cell line GT38. The results also revealed that after *LMP2A* was knocked down by lentivirus-mediated RNAi, the cell proliferation and growth potency of EBVaGC cells were decreased significantly. Thus, a new strategy involving reduction of the expression of *LMP2A* by lentivirus-mediated RNAi is provided with the potential for treating human EBVaGC.

In conclusion, the results of the present study show that the knockdown of *LMP2A* by lentivirus-mediated RNAi significantly suppresses cell growth and induces apoptosis in the EBVaGC cell line GT38. This provides an attractive anticancer therapeutic strategy for the treatment of human EBVaGC.

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