# Autophagy is involved in recombinant Newcastle disease virus (rL-RVG)-induced cell death of stomach adenocarcinoma cells *in vitro*

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Abstract. Oncolytic viruses can kill malignant cells while sparing normal cells. Multiple pathways are involved in this action. The antitumor effects of viral infection on SGC-7901 and AGS cells were investigated. We measured endoplasmic reticulum stress and autophagy caused by the recombinant avirulent Newcastle disease virus (NDV) LaSota strain expressing the rabies virus glycoprotein (rL-RVG) and the NDV wild-type strain. The dose-response curves were analyzed using the MTT assay. The expression of RVG was detected by western blotting, RT-PCR and immunofluorescence analyses. Cell death and autophagy were observed using transmission electron microscopy, TUNEL and western blotting. Endoplasmic reticulum stress and the mitochondrial transmembrane potential were detected by western blotting and immunofluorescence, respectively. Immunofluorescence, western blot and RT-PCR analyses indicated that RVG gene and protein were expressed in SGC-7901 and AGS cells infected by rL-RVG. MTT and TUNEL analyses showed that the growth of SGC-7901 and AGS cells in the rL-RVGinfected group was significantly inhibited compared with the wild-type NDV-infected group (p<0.05). Western blot analysis indicated that rL-RVG and NDV induced increases in apoptosis, endoplasmic reticulum stress, and autophagy in the SGC-7901 and AGS cells. However, apoptosis and autophagy decreased in these cells after the application of the autophagy pathway inhibitor 3-MA or ATG-5-specific siRNA. Immunofluorescence analysis showed that the mitochondrial membrane potential collapsed. Taken together, these results indicate that the rL-RVG virus group is much more powerful compared with the NDV-infected group (p<0.05). rL-RVG and NDV are potent antitumor agents that induce autophagy.

## Introduction

In recent decades, an integrated approach to malignant cell treatment that includes surgery, chemotherapy (1), radiotherapy, gene therapy (2,3) and immune therapy (4), has become a reasonable therapeutic strategy. Oncolytic virotherapy using agents such as Newcastle disease virus (NDV) is one of the new biological strategies for gene (2) and immune therapy (4-6) and has been tested against many different cancers, including gastric cancer (4), skin tumors (7) and solid cancers (8).

NDV, which is a member of the *Paramyxoviridae* family, is a non-segmented negative-strand RNA virus (NNSV). The NDV genes encode six major structural proteins, including nucleoprotein (NP), phosphoprotein (P), matrix protein (M), fusion protein (F), hemagglutinin-neuraminidase (HN) and large (L) RNA-dependent RNA polymerase, in the following order: 3'-NP-P-M-F-HN-L-5'. Oncolysis (9), apoptosis (10,11), and enhanced innate immunity (4,12,13) are the known mechanisms by which NDV kills malignant cells.

The rabies virus (RV), which is a member of *Rhabdoviridae* family, is an enveloped, bullet-shaped virus that is also associated with host cell apoptosis (13,14). Some clinical trials as early as the 1950s and 1960s demonstrated the effectiveness of RV in treating melanomatosis (15).

Autophagy is a catabolic and highly conservative cellular process. Its basic role is to recycle cellular components during nutritional starvation and other stressful conditions (16) and is considered a cytoprotective event (17). Despite its protective effects, autophagy can also induce a type of cell death (18) called 'type II programmed cell death' (19). Whether autophagy contributes to cell death or to protection during chemotherapy remains controversial (20). However, increasing evidence has demonstrated that autophagy represents cell death (21). Additionally, many chemicals, such as avicin D (22) and arsenic trioxide (23), as well as cellular stresses (24), and mitochondrial dysfunctions (25) can contribute to autophagy. Autophagy is a known potential anticancer strategy. Previous studies have shown that recombinant NDV-expressing rabies virus glycoprotein (rL-RVG) has the ability to spread from cell to cell without the help of trypsin (26), to alter its self-replication, and to induce cell death via the apoptosis pathway (27).

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The present study examined the contribution of autophagy to apoptosis induced by NDV or rL-RVG and raised a hypothesis regarding cross-talk among autophagy, apoptosis, mitochondrial dysfunction and endoplasmic reticulum stress (ERs).

#### Materials and methods

Materials. The NDV strain LaSota, rL-RVG, and anti-NDV antibody were provided by Harbin Veterinary Research Institute, Chinese Academy of Agricultural Sciences. The human cancer cell lines SGC-7901 and AGS were purchased from the Cancer Cell Repository (Shanghai Cell Bank, 2010-02-20) and ATCC (MD, USA), respectively. 3-[4,5-dimehyl-2-thiazolyl]-2,5-diphenyl-2H-tetrazolium bromide (MTT) was purchased from Amresco (PA, USA). 3-Methyladenine (3-MA) and Hoechst 33342 were obtained from Sigma-Aldrich (St. Louis, MO, USA). The siRNA specific for human beclin-1 was from GenePharma (Shanghai, China). All PCR primers were purchased from Shanghai Sangon Biological Engineering Technology & Services Co., Ltd. (Shanghai, China). TRIzol and Lipofectamine 2000 were from Invitrogen (CA, USA). Rabbit polyclonal anti-caspases 3, 8 and 9, anti-bax, antibeclin-1, anti-caspase 12, anti-HSP90, and anti-cytochrome c; and mouse monoclonal anti-Bcl-2 and anti-grp 78 antibodies were from Boster (Wuhan, China). Rabbit monoclonal anti-LC3 antibody was from Epitomics (Burlingame, CA, USA), and mouse monoclonal anti-rabies virus was obtained from Santa Cruz (CA, USA). HRP-conjugated goat anti-rabbit, HRP-conjugated goat anti-mouse, and FITC-conjugated goat anti-mouse antibodies were purchased from CWBio (Shanghai, China). HRP-conjugated rabbit anti-chicken antibody was from EarthOx Life Science (Millbrae, CA, USA), and Cy3-conjugated rabbit anti-chicken antibody was purchased from KPL (CA, USA). The PVDF membrane and Luminata were provided by Millipore (MA, USA). DMEM, trypsin, and EDTA-2Na were offered by Gibco (NY, USA). Fetal bovine serum (FBS) was supplied by Hyclone (UT, USA). A mitochondrial transmembrane potential analysis kit was from KeyGen Biotech (Nanjing, China). All other supplies for cell culture were obtained from Costar Corning (NY, USA).

Cell culture and interference test. SGC-7901 and AGS cells were maintained in DMEM medium with 10% (v/v) FBS, antibiotics (100 U/ml penicillin and 100 U/ml streptomycin) at 37°C with 5% CO<sub>2</sub> and 100% humidity. When the cells reached ~50-70% confluence, the cells were infected with NDV or rL-RVG.

*3-MA application.* When the SGC and AGS cells grown in 6-well plates reached ~50-70% confluence, the cells were treated with 2.5 mM 3-MA. The next day, the cells were infected with NDV and rL-RVG at a multiplicity of infection (MOI) of 10. At 24 h after being infected, the cells were harvested and analyzed by western blotting.

*siRNA application*. When the SGC and AGS cells grown in 6-well plates reached ~50-70% confluence, the cells were transfected with 50 nM siRNA (beclin-1) according to the manufacturer's instructions. The next day, the cells were infected with NDV and rL-RVG at a MOI of 10. At 24 h after

being infected, the cells were harvested and analyzed by western blotting.

*MTT analysis*. SGC-7901 or AGS cells (1x10<sup>4</sup>) were cultured in 96-well plates for 24 h with 10% FBS at 37°C with 5% CO<sub>2</sub>. On the second day, the cells were incubated in serum-free DMEM with rL-RVG or NDV at concentrations of 10<sup>-2</sup>, 10<sup>-3</sup>, 10<sup>-4</sup> and 10<sup>-5</sup>. After 1 h, 10% FBS was added to each well, and then the cells were cultured as described previously, with a phosphate-buffered saline (PBS) group as the negative control. On the third day, each well was treated with 20  $\mu$ I MTT (5 mg/ml, PBS) and incubated for an additional 4 h. Then, 150  $\mu$ I of DMSO was added into each well, and the cells were shaken for 10 min. Finally, the absorbance was measured in triplicate using a standard spectrophotometer. The doseresponse curve for cell survival was made based on data from the MTT assay. Simultaneously, the morphological changes in the infected cells were monitored under a microscope.

*Immunofluorescence analysis.* Cells were cultured in 24-well plates for 24 h as previously described, infected with NDV or rL-RVG at a MOI of 10, or treated with PBS as a negative control. Then, at 24 h post-infection, the cells were fixed in 4% paraformaldehyde at 4°C overnight, followed by immunofluorescence staining with antibodies against NDV and RV and Hoechst 33342 staining. These stained SGC and AGS cells were monitored using immunofluorescence microscopy.

*RT-PCR analysis*. The cells were cultured and infected as described previously. At 24 h after infection, total RNA was extracted from the cultured cells using TRIzol reagent. First-strand cDNA synthesis was performed using oligo(dT) primers and M-MLV reverse transcriptase. The primer set for the NDV HN gene, the RV G gene and the human GAPDH gene are shown in Table I. The RT-PCR protocol was as follows: an initial denaturation at 94°C for 5 min, followed by 30 cycles at 94°C for 30 sec, annealing at 53°C (RVG) or 55°C (NDV and GAPDH) for 30 sec, and extension at 72°C for 30 sec, with a final extension at 72°C for 10 min. Then, 5  $\mu$ l of each PCR product was loaded onto 1% agarose gels for electrophoresis and visualized with ethidium bromide. The resulting bands were analyzed using Quantity One software (Bio-Rad). The primers for PCR are shown in Table I.

Western blot assay. SGC and AGS cells with or without treatment were lysed in RIPA lysis buffer with a protease inhibitor cocktail (Santa Cruz). The protein concentration was measured using a BCA kit (Thermo Fisher Scientific, USA). An equal amount of protein from each sample was loaded onto a 10% polyacrylamide gel and separated by electrophoresis. Then, the proteins were transferred to a polyvinylidene difluoride (PVDF) membrane (Millipore, CA, USA). The membrane was blocked for 1 h in 5% BSA. Then, the membrane was incubated with primary antibodies against specific proteins (i.e., caspases 3, 8 and 9, bcl-2 and bax for apoptosis; NDV and RVG for infection, Beclin-1 and LC-3 for autophagy; anti-HSP90, anti-cytochrome c and anti-grp 78 for endoplasmic reticulum stress; and beta-actin as a control) and then incubated with HRP-conjugated secondary antibodies. The protein bands were scanned using a Typhoon

Primer		Sequence	Product size (bp)
rL-RVG	Upstream Downstream	5'-AGCCGATGCTCACTACAAG-3' 5'-CTGGAGGAGGGATGATTGC-3'	175
NDV	Upstream Downstream	5'-CTGGACGGTTTGGTGGGAA-3' 5'-TAATGCGACTGCGGGATGTG-3'	462
GAPDH	Upstream Downstream	5'-CAAGGTCATCCATGACAACTTTG-3' 5'-GTCCACCACCCTGTTGCTGTAG-3'	-

Table II. MTT results of the cellular dose-response curve for SGC cells after 24 h of infection.

Dilution	rL-RVG	NDV	t-value	p-value
10 <sup>1</sup>	0.18656±0.029626	0.283727±0.026238	3921	0.017
10 <sup>2</sup>	0.312080±0.04746	0.55666±0.024781	11.521	< 0.001
10 <sup>3</sup>	$0.54400 \pm 0.037857$	$0.62300 \pm 0.028032$	3.461	0.02
104	0.56130±0.038402	0.61522±0.61522	4.971	0.08

9400 Variable Mode Imager (Amersham Biosciences, UK) and detected by Pierce ECL Plus Substrate (Thermo Fisher Scientific).

*Transmission electron microscopy (TEM).* Standard TEM was performed to monitor the ultrastructure of the cells. Cells were infected with virus as described previously. At 24 h after infection, the cells were fixed and embedded in 4% paraformaldehyde and 2.5% glutaraldehyde. Thin sections were cut and examined at 200 kV using an H-600 transmission electron microscope. Autophagy, as defined by the presence of double-membraned vacuoles; apoptosis, as characterized by chromatin margination, pyknosis, and apoptotic bodies; and endoplasmic reticulum swelling were all observed.

*TUNEL assay.* Cells were cultured in 24-well plates with slides and 10% FBS DMEM at 37°C with 5% CO<sub>2</sub>. Then, the cells were infected with rL-RVG and NDV within the logarithmic growth phase and fixed in 4% paraformaldehyde. After fixing, the cells were stained according to the manufacturer's instructions. The slides were observed and imaged under an optical microscope. The apoptosis index (AI) was calculated as the number of apoptotic cells/(the number of apoptotic cells + the number of non-apoptotic cells) x 100%.

Detection of mitochondrial membrane potential (MMP). JC-1 fluorescence dye was used to determine the mitochondrial membrane potential. Cells were cultured in 24-well plates and treated with NDV or rL-RVG as described above for 24 h, followed by staining with JC-1 ( $2.5 \ \mu g/ml$  at 37°C for 30 min). The cells were monitored using fluorescence microscopy with excitation wavelengths at 527 nM for green and at 590-600 nM for red. The changes in the MMP could be accurately assessed by comparing the ratios of 590-600 nM (red)/527 nM (green). Statistical analysis. The data comparisons were performed using one-way analysis of variance (ANOVA) in SPSS V17.0 software. Differences were considered statistically significant at p<0.05. All experiments were repeated at least three times.

### Results

Expression of viral genes and proteins in infected stomach adenocarcinoma cells. The expression of viral proteins was monitored by fluorescence microscopy. Almost all of the SGC and AGS cells that were infected with NDV expressed the NDV HN protein, and almost all of the SGC and AGS cells infected with rL-RVG expressed both the NDV HN protein and RVG protein, while neither of these proteins was present in the control group. RVG and NDV protein expression was upregulated in the rL-RVG-infected group compared with the NDV-infected group, as shown in Fig. 1A and B. Western blot analysis confirmed these results, as shown as Fig. 1C. The PCR products of the NDV HN and RVG genes were detected to assess NDV HN and RVG mRNA expression in infected SGC and AGS cells. The results showed that the RVG gene (175 bp) was expressed in SGC and AGS cells infected with rL-RVG and that the NDV HN gene (462 bp) was expressed in the SGC and AGS cell infected groups. In contrast, neither the RVG gene nor the NDV HN gene was expressed in the control group (Fig. 1D).

The proliferation changes of SGC and AGS cells infected with *rL-RVG and NDV*. The dose-response curves for rL-RVG and NDV in SGC cells were monitored by MTT. After 24 or 48 h of rL-RVG or NDV infection, the infected SGC cells were assessed by MTT assay. SGC cell viability decreased with an increase in virus concentration or in incubation time. When the cells were infected simultaneously with NDV and rL-RVG, the OD of MTT increased in both the NDV- and rL-RVG.

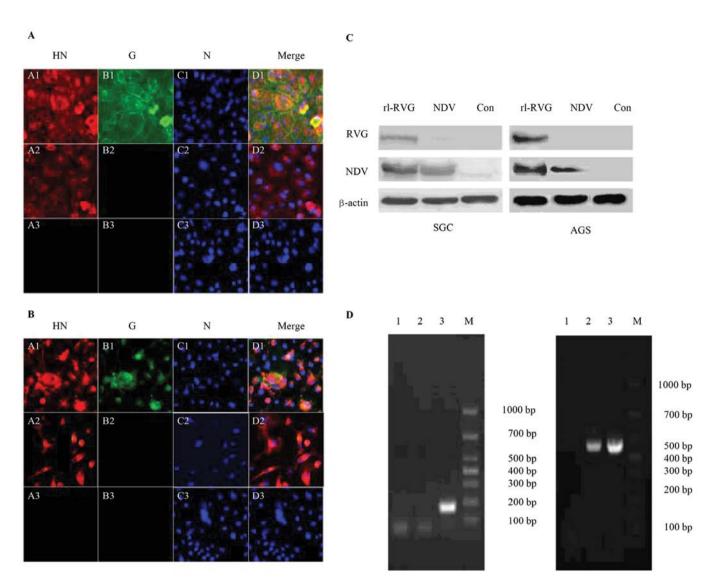
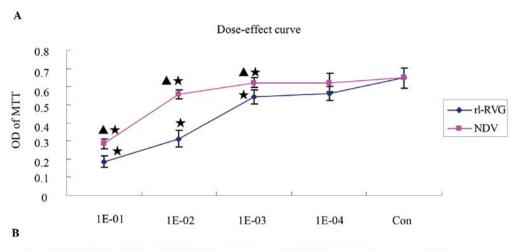


Figure 1. Expression of viral proteins and genes in infected SGC and AGS cells. RVG and NDV protein expression was monitored by immunofluorescence (x200 magnification) (A) for SGC and (B) for AGS. The RVG protein (green) was only expressed in the rL-RVG-infected group, while the NDV protein (red) was expressed in both the rL-RVG- and NDV-infected groups. Neither protein was expressed in the control group. (C) RVG and NDV protein expression in SGC cells and in AGS cells was detected by western blotting. The RVG protein was expressed in the rL-RVG-infected group, and the NDV protein was expressed in both the rL-RVG- and NDV-infected groups. Neither protein was expressed in the PBS group. (D) NDV and RVG mRNA expression. Lane 1, PBS group; lane 2, NDV-infected group; lane 3, rL-RVG-infected group; M, marker. (B) NDV HN mRNA expression. Lane 1, PBS group; lane 2, NDV-infected group; M, marker.

infected groups as the virus dilution decreased, and at the 10<sup>3</sup> dilution, the dose-response curve reached a plateau compared with the control group. Simultaneously, the OD of MTT in the rL-RVG-infected group was weaker compared with that in the NDV-infected group, suggesting that the rL-RVG-infected group had a greater inhibition ratio. The inhibition ratio increased over time after infection. Thus, the inhibition ratio caused by NDV and rL-RVG infection was dependent on both time and dose. The OD values of MTT in SGC cells after 24 h are shown in Fig. 2A and Table II. Additionally, the morphological changes in the infected or uninfected cells, as observed by microscopy, are shown in Fig. 2B; the cells had a greater decrease in size in the rL-RVG-infected group compared with the NDV-infected group, and the number of smaller cells was higher in the rL-RVG-infected group compared with the NDV-infected group. No changes were observed in the PBS group.

*rL-RVG and NDV infection affected the ultrastructure of SGC and AGS cells.* We monitored the cells by TEM to assess the ultrastructure of the cells after infection directly. The cells that were infected with NDV or rL-RVG showed increased autophagy (black arrow), apoptosis (white arrow) and ERs (white triangle), with a larger effect observed in the L-RVG group compared with the NDV-infected group, as shown in Fig. 3.

*NDV and rL-RVG induce apoptosis in SGC and AGS cells.* Apoptosis was monitored by TUNEL assay; apoptosis in the virus-infected SGC cells was obvious compared with the control cells, and the number of apoptotic cells in the rL-RVG-infected groups was greater than that in the NDV-infected group, as shown in Fig. 4A. Furthermore, to assess the apoptotic cell death pathway caused by NDV and rL-RVG infection, the expression of the apoptosis-associated



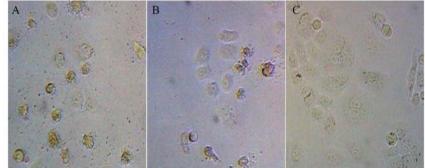


Figure 2. Proliferation changes in SGC cells after viral infection. (A) Dose-response curve of SGC cells after viral infection at 24 h. The OD of MTT after dilution (n=5). The cells infected with rL-RVG for 24 h showed significantly decreased absorbance when compared with OD of the NDV-infected group ( $^{\circ}p<0.05$ ). A significant difference was observed between different dilutions of the same virus ( $^{\circ}p<0.05$ ). (B) Images of SGC cells infected with virus at 24 h using microscopy (x200 magnification). (a) rL-RVG-infected group (10-2); (b) NDV-infected group (10-2); (c) control group. The cells infected with rL-RVG decreased in size compared with those of the NDV-infected group, and the number of smaller cells was higher in the rL-RVG-infected group compared with that of the NDV-infected group. No changes were observed in the control group.

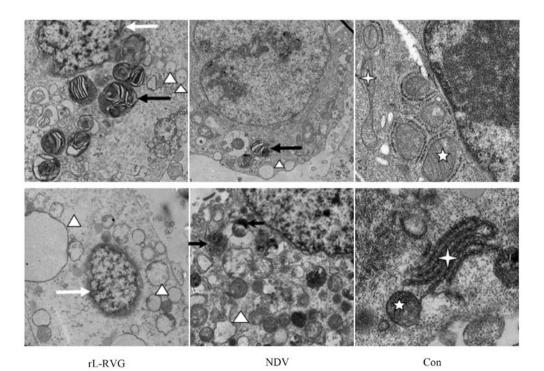


Figure 3. The ultrastructure of the SGC cells after viral infection. The black arrow shows autophagy, white arrows indicate apoptosis, and white triangles show ERs. Additionally, white stars and pentagons indicate the endoplasmic reticulum and mitochondria in the PBS group, respectively. Autophagy, apoptosis and ERs in cells from the rL-RVG-infected group significantly increased compared with those processes in cells from the NDV-infected group. No effect was

observed in the PBS group.

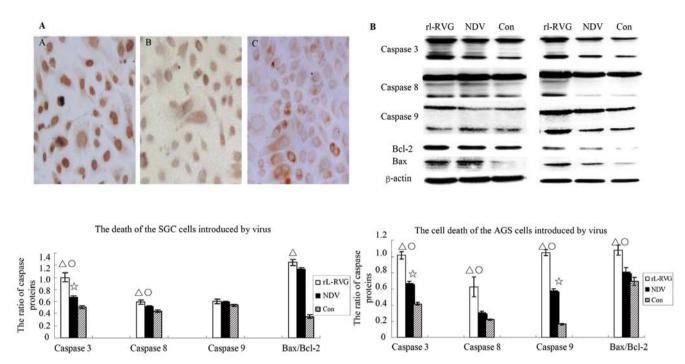
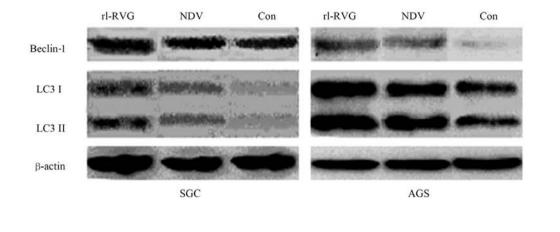


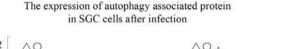
Figure 4. Apoptosis and its associated protein expression in SGC and AGS cells at 24 h after infection. (A) Apoptosis of infected SGC cells, as measured by TUNEL staining. Apoptosis of virus-infected cells was obvious compared with the controls. The apoptotic cells in the rL-RVG-infected groups were more abundant compared with those in the NDV-infected group. (B) The expression of apoptosis-associated proteins in SGC and AGS cells, as detected by western blotting. The proteins cleaved caspase 3 and 8 and bax/bcl-2 in the rL-RVG-infected group had higher expression levels compared with those in the other two groups ( $^{\circ}$ - $^{\circ}$ p<0.05); these proteins also had higher expression levels in the NDV-infected group compared with the controls ( $^{\circ}$ p<0.05). No significant differences were observed in the following pairs: in the SGC cells, the cleaved caspase 8 protein levels did not significantly differ between the NDV-infected group and the control group, the cleaved caspase 9 protein levels did not significantly differ between the NDV-infected group and the control group; in the AGS cells, the cleaved caspase 8 protein levels did not significantly differ between the NDV-infected group and the control group, and the protein ratio of bax of bcl2 did not significantly differ between the NDV-infected group and the control group and the control group, and the protein ratio of bax of bcl2 did not significantly differ between the NDV-infected group and the control group and the co

proteins caspase 3, 8 and 9, bcl-2, and bax in the SGC and AGS cells was examined routinely at 24 h after being infected with NDV or rL-RVG. Accumulations of cleaved caspase 3, 8 and 9 and of bax/bcl-2 were found in SGC and AGS cells infected with virus. The expression levels of cleaved caspase 3 and 8 and bax/bcl2 proteins were upregulated in the rL-RVG-infected group compared with the other two groups. The expression levels of these proteins were higher in the NDV-infected group compared with those in the control group. In the SGC cells, the expression levels of cleaved caspase 8 protein did not differ between the NDV-infected group and the control group. Additionally, the expression levels of cleaved caspase 9 protein did not differ among all the three groups, and the protein ratio of bax to bcl2 did not differ between the NDV-infected group and the control group. In the AGS cells, the expression levels of cleaved caspase 8 protein did not differ between the NDV-infected group and the control group, and the protein ratio of bax to bcl-2 did not differ between the NDV-infected group and the control group, as shown in Fig. 4B.

*NDV and rL-RVG induce significant autophagy in SGC and AGS cells.* Because TEM analysis indicated that autophagy significantly increased after SGC cells were infected with virus, we detected the autophagy activity of cells with or without NDV or rL-RVG infection. The ratio of lipid-bound LC3-II, which is converted to LC3-I, the soluble LC3-I is associated with the formation of autophagosomes and can indicate autophagic activity. In a preliminary experiment, we found that the autophagic activity peaked at least two times after infection from 30 min to 48 h. The first peak occurred at the 3rd hour after infection, while the second peak occurred at 24 h after infection (data not shown). Considering that autophagy protects against infection, we suggested that the second peak time contributes to cell death. In addition to monitoring the ratio of LC3-II to LC3-I in cells at 24 h post-infection, we also monitored another autophagy marker, beclin-1, in these cells. As shown in Fig. 5. the expression of the autophagy-associated protein beclin-1 and LC3 in the SGC cells of the rL-RVGinfected group showed a significant difference compared with the other two groups ( $^{\wedge}, ^{\circ}p < 0.05$ ); the expression of the autophagy-associated protein in the NDV-infected group showed a significant difference compared with the control group ( $\Rightarrow$ p<0.05). The trend is the same in the expression of beclin-1 and LC3 in the AGS cells and the same trend exists among the three group. However, beclin-1 in the AGS cells showed no significant differences between the rL-RVG and the NDV-infected group (p>0.05).

*NDV and rL-RVG cause endoplasmic reticulum stress in the SGC cells.* Recent evidence has demonstrated that ERs is a type of trigger for autophagy. In the present study, TEM images showed infected cells with endoplasmic reticulum swelling, indicating that the SGC cells had ERs. Therefore, we monitored the expression of the following proteins associated with ERs: grp-78, HSP-90 and cytochrome *c*. The expression of the grp-78, and HSP-90 proteins in SGC cells showed significant





The expression of autophagy associated protein in AGS cells after infection

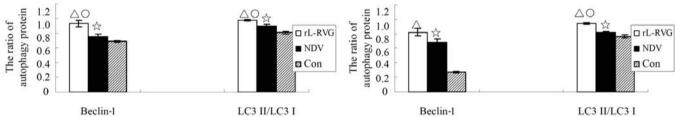


Figure 5. NDV- and rL-RVG-induced autophagy of SGC and AGS cells. The expression of the autophagy-associated proteins beclin-1 and LC3 in the NDV-infected group significantly differed compared with that of the control group ( $^{\circ}p$ <0.05) and compared with that of the rL-RVG-infected group and the NDV-infected group ( $^{\circ}p$ <0.05). The expression of these autophagy-associated proteins in the rL-RVG-infected group significantly differed compared with that of the control group ( $^{\circ}p$ <0.05).

differences among the three groups ( $^{\Delta, \dot{\pi}, O}$ p<0.05), and the expression of the cytochrome *c* protein in the control groups was higher compared with that in the two infected groups ( $^{\Delta, \dot{\pi}}$ p<0.05). However, no significant difference was observed between the rL-RVG and NDV-infected groups (p>0.05). These results suggested that the infected cells remained under ERs.

Autophagy contributes to SGC and AGS apoptosis induced by NDV and rL-RVG. Recent research has indicated that autophagy can have both cytoprotective and cytotoxic effects (28,29). 3-MA, which is an inhibitor of autophagy, was administered to SGC and AGS cells with or without NDV or rL-RVG infection to determine whether autophagy contributes to the apoptosis of NDV- and rL-RVG-infected cells. The expression of the apoptosis-associated proteins caspase 3, 8 and 9 and calpain 1, which is related to autophagy and apoptosis (30), decreased in SGC and AGS cells with 3-MA treatment relative to control cells without 3-MA treatment. The effect was greater in the rL-RVG-infected groups than in the NDV-infected groups, as shown in Fig. 7A. Because 3-MA is a non-specific autophagy suppressor, it inhibits both PI3K-III (an autophagy inducer) and PI3K-I (an autophagy suppressor). We silenced beclin-1, which is an autophagy regulator, to confirm the 3-MA results. The potency of the siRNA treatment was determined by western blotting, as shown in Fig. 7B. We found that the expression of the beclin-1 protein decreased significantly in SGC cells upon beclin-1 siRNA treatment. Therefore, we monitored the apoptosis-associated proteins caspase 3, 8 and 9

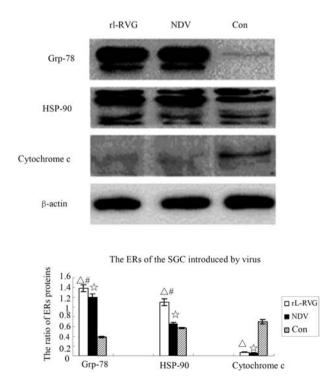


Figure 6. The expression of virus-induced endoplasmic reticulum stressassociated proteins in the SGC cells. The expression of ERs-associated proteins in the NDV-infected group significantly differed compared with that of the control group ( $^{\circ}p<0.05$ ) and the expression of these proteins in the rL-RVG-infected group significantly differed compared with that of the control group ( $^{\circ}p<0.05$ ). A significant difference ( $^{\circ}p<0.05$ ) in the expression of these proteins was observed between the rL-RVG-infected group and the NDV-infected group. In contrast, no significant difference in cytochrome *c* levels was observed between the rL-RVG- and NDV-infected groups.

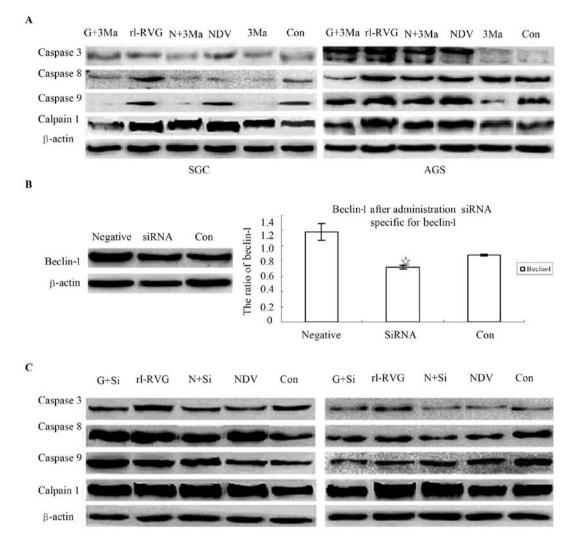


Figure 7. The contribution of downregulated autophagy to the expression of apoptosis-associated proteins in SGC and AGS cells after infection. (A) Apoptosisassociated protein expression after 3-MA treatment. The expression of the apoptosis-associated proteins caspase 3, 8 and 9 and calpain-1 decreased in cells after 3-MA treatment compared with the cells without 3-MA treatment, and the above proteins were more strongly expressed in the rL-RVG-infected groups compared with the NDV-infected groups. (B) The expression of beclin-1 in cells. After administering beclin-1-specific siRNA, the expression of the beclin-1 protein decreased. (C) Apoptosis-associated protein expression after siRNA treatment. The expression of the apoptosis-associated proteins caspase 3, 8 and 9 and calpain 1 decreased in cells after beclin-1-specific siRNA treatment compared to those cells without siRNA treatment. These proteins were more strongly expressed in the rL-RVG-infected groups compared with the NDV-infected groups.

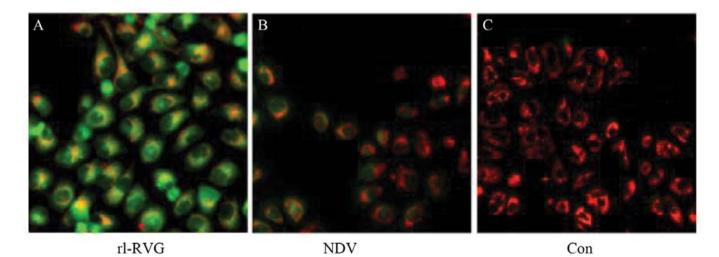


Figure 8. NDV and rL-RVG infection induced MMP collapse in SGC cells. (A) rL-RVG-infected group, (B) NDV-infected group, and (C) control group. A greater number of dysfunctional mitochondria are present in rL-RVG-infected cells than in NDV-infected cells, and the control group has few dysfunctional mitochondria compared with the NDV-infected group.

and calpain 1, and the results were similar to the 3-MA treatment experiment, as shown in Fig. 7C.

*NDV and rL-RVG induce mitochondrial dysfunction*. Mitochondria have essential roles in autophagy (31). We monitored the MMP with JC-1 staining. The greater the intensity of green in a cell, the more dysfunctional the mitochondria are. We found that the red fluorescence intensity decreased and the green fluorescence intensity increased in these cells. The ratio of red to green significantly decreased in the NDV- and rL-RVG-infected groups compared with the control group, and the ratio of red to green the in the rL-RVG-infected group decreased more than that in NDV-infected group, as shown in Fig. 8.

## Discussion

NDV, which is a member of the oncolytic virus family, has the ability to inhibit malignant cells via multiple mechanisms (15). In recent decades, NDV has been shown to induce tumor cell death via its oncolytic effects (9). NDV can also induce the immune system to eliminate tumor cells (6), and NDV can infect tumor cells more effectively compared with normal cells (12). The rL-RVG virus spreads more easily within cells (26), and RVG has a potential oncolytic effect (13,14); therefore, we propose that the recombinant virus rL-RVG has much more antitumor potency.

In the present study, we found that rL-RVG caused stomach adenocarcinoma SGC-7901 and AGS cell death, which is consistent with previous studies (27). Interestingly, in addition to apoptosis, both NDV and rL-RVG can induce autophagy, ERs and mitochondrial dysfunction. The results of the present study indicated that the net effects of autophagy, mitochondrial dysfunction and ERs contribute to tumor cell death. We considered that the observed mitochondrial dysfunction, ERs, autophagy, and apoptosis, indicate possible cross-talk with other pathways.

The autophagy pathway cross-talks with the apoptosis pathway; virus-induced autophagy can also contribute to apoptosis (32). Autophagy might also be a mechanism that contributes to virus-induced cell death (33). Although autophagy can promote tumor survival in some cases (29,34), autophagy can also be fatal to malignant cells (35). Autophagy and apoptosis work together in multiple ways to determine the fate of tumor cells (28,36). In the present study, both autophagy and apoptosis increased in the cells after viral infection. In the last decade, many researchers have confirmed that Bcl-2 plays an important role in both apoptosis and autophagy (37-39). Beclin-1 has an N-terminal BH3 domain, which this protein a subgroup of the Bcl-2 family (39,40); therefore, beclin-1 can participate in the activity of the Bcl-2/Bcl-xL complex and inhibit the formation of autophagosomes (41). We found that the virus-infected cells showed increase beclin-1 expression, thus decreasing the interaction between Beclin-1 and Bcl-2/Bcl-xL and promoting autophagy and apoptosis. In addition, caspase 8 and caspase 12 play important roles in autophagy (41,42). We also found that the expression of apoptosis-associated proteins increased in cells after viral infection. We suggest that rL-RVG and NDV cause tumor cell death via both autophagy and apoptosis.

Autophagy and ERs are related (43); both excess autophagy and excess ERs can cause cell death (44), and both moderate autophagy and moderate ERs are adaptive mechanisms when cells are under severe stress (45). Autophagy can recycle the organelles and provide the basic energy and material for cell survival (16,46). In addition, Meng et al (47) and Meng et al (48), reported that autophagy can enhance NDV replication in tumor cells. Considering that NDV and rL-RVG introduced massive autophagy in the present study, we propose that the autophagy induced by NDV and rL-RVG allowed the viruses to increase replication and that these increased amounts of the viruses induced even more autophagy, thus becoming a positive feedback loop, with more and more virus initiating the unfolded protein response and causing ERs. Finally, autophagy and ERs reach a certain threshold, inducing cell death.

Additionally, autophagy and mitochondrial dysfunction work together to cause cell death. Mitochondria contribute to autophagy during cell death and disease by the mitochondrial permeability transition pathway (46,49) and by reactive oxygen species (ROS) generation (50). The protein calpain, which is a member of the calmodulin family, can degrade ATG-5, and cleaved ATG-5 can anchor to mitochondria, causing cytochrome c release, followed by mitochondrial dysfunction (30). In the present study, we found that the viruses caused excess autophagy and mitochondrial dysfunction, indicating that these viruses participate in cell death. The above results suggest that autophagy and mitochondrial dysfunction contribute to cell death. Although autophagy dysfunction may result in abnormal mitochondrial function (51), further experiments will be required to establish crosstalk among autophagy, mitochondria and oxidative stress pathways.

In conclusion, this study demonstrated that NDV and rL-RVG induce stomach adenocarcinoma cell death via autophagy and apoptosis, in association with ERs and mitochondrial dysfunction. Cell death may involve multiple pathways. Although the causal relations among autophagy, apoptosis and ERs remain unclear, rL-RVG may become a powerful candidate for antitumor treatments.

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