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Downregulation of microRNA-30a-5p contributes to the replication of duck enteritis virus by regulating Beclin-1-mediated autophagy

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Abstract

Background: MicroRNAs (miRNAs) is increasingly recognized as an important element in regulating virus-host interactions. Our previous results showed that cellular miR-30a-5p was significantly downregulated after duck enteritis virus (DEV) infection cell. However, whether or not the miR-30a-5p is involved in DEV infection has not been known.

Methods: Quantitative reverse-transcription PCR (qRT-PCR) was used to measure the expression levels of miRNAs(miR-30a-5p) and Beclin-1 mRNA. The miR-30a-5p - Beclin-1 target interactions were determined by Dual luciferase reporter assay (DLRA). Western blotting was utilized to analyze Beclin-1-mediated duck embryo fibroblast (DEF) cells autophagy activity. DEV titers were estimated by the median tissue culture infective dose (TCID₅₀).

Results: The miR-30a-5p was significantly downregulated and the Beclin-1 mRNA was significantly upregulated in DEV-infected DEF cells. DLRA confirmed that miR-30a-5p directly targeted the 3'-UTR of the Beclin-1 gene. Overexpression of miR-30a-5p significantly reduced the expression level of Beclin-1 protein ($p < 0.05$), leading to the decrease of Beclin-1-mediated autophagy activity, which ultimately suppressed DEV replication ($P < 0.05$). Whereas transfection of miR-30a-5p inhibitor increased Beclin-1-mediated autophagy and triggered DEV replication during the whole process of DEV infection ($P < 0.01$).

Conclusions: This study shows that miR-30a-5p can inhibit DEV replication through reducing autophagy by targeting Beclin-1. These findings suggest a new insight into virus-host interaction during DEV infection and provide a potential new antiviral therapeutic strategy against DEV infection.

Keywords: Duck enteritis virus, Autophagy, miR-30a-5p, Beclin-1

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Background

Duck enteritis virus (DEV) is the causative pathogen of duck viral enteritis disease, causing considerable economic losses in the duck industry due to high mortality and low egg production [1, 2]. In addition, DEV can cause variable morbidity and mortality in geese, swans and other wild waterfowl and poses a severe threat to waterfowl groups [3–5]. DEV is classified into the family *Herpesviridae*, sub-family *Alphaherpesvirinae*, genus *Mardivirus* and *anatid herpesvirus 1* [6]. Its genome is a linear double-stranded DNA molecule that is composed of a unique long region (UL) and a unique short region (US) flanked by a short internal repeat sequence (IRS) and a short terminal repeat sequence (TRS) [7–10].

Autophagy is an essential self-digestion process that degrades protein and waste in cells to maintain cellular metabolic balance and homeostasis [11, 12]. Growing evidence has shown that viral infection can induce cellular autophagy. For example, some viruses, such as Newcastle disease virus (NDV) [13], classical swine fever virus (CSFV) [14], porcine circovirus type 2 (PCV2) [15], porcine reproductive and respiratory syndrome virus (PRRSV) [16], dengue virus, foot-and-mouth disease virus (FMDV) and varicella-zoster virus (VZV) [17], can induce cell autophagy to enhance their replication. However, autophagy can suppress viral replication and eliminate viral infection [18, 19]. For example, cellular autophagy can inhibit replication of vesicular stomatitis virus (VSV) by regulating the P13K/AKT signaling pathway [20]. Recent research has reported that DEV induces autophagy to enhance its replication in duck embryo fibroblast (DEF) cells [21]. Nevertheless, the regulatory relationship of autophagy remains poorly understood.

MicroRNAs (miRNAs) are important small (18–24 nt), noncoding, endogenous RNAs that can negatively regulate gene expression by binding fully or partially to the 3'-untranslated region (3'-UTR) [22, 23]. Accumulating evidence has demonstrated that miRNAs participate in a wide range of biological processes, including cellular proliferation, differentiation, signal transduction, metabolism apoptosis and cellular autophagy [24–27], and play important roles in regulating virus-host interactions [28–30]. Our previous high-throughput sequencing results revealed that 13 cellular miRNAs (mir-125-2-3p, mir-124a-3p, mir-215-5p, mir-29b-3p, etc) were significantly upregulated and 25 miRNAs (mir-1a-3p, mir-133a-5p, miR-30a-5p, miR-16c-5p, etc) were significantly downregulated after CHV infection [31]. Therefore, we speculate that these miRNAs may play crucial roles in DEV infection.

In this study, we first confirmed that miR-30a-5p directly targeted the 3'-UTR of the Beclin-1 mRNA. Further study showed that overexpression of miR-30a-5p inhibited DEV replication by downregulating Beclin-1-

mediated autophagy in DEF cells. miR-30a-5p inhibitor triggered DEV replication, suggesting that miR-30a-5p plays important roles in the regulation of DEV-induced autophagy and viral proliferation. These data provide a basis for further understanding miRNAs' regulatory roles in cellular autophagy and should contribute to the development of anti-DEV drugs.

Methods

Virus, cells, miRNA mimic and antibodies

The DEV CHV (Chinese virulent strain) (accession No. JQ647509) and mouse anti-UL41 serum were provided by the Avian Diseases Research Center, College of Veterinary Medicine, Sichuan Agricultural University. Duck embryo fibroblast (DEF) cultures were prepared from 10-day-old duck embryos for the propagation of CHV. The study was approved by the Animal Ethics Committee of Sichuan Agricultural University (approval No. XF2016–17). Cell monolayers were cultured in Dulbecco's Modified Eagle's Medium (DMEM, Gibco, Grand Island, NY USA) supplemented with 10% fetal bovine serum (FBS, Gibco, USA) and 1% penicillin-streptomycin (Gibco, USA) at 37 °C in a 5% CO₂ atmosphere. The miR-30a-5p mimic, mimic negative-control (mimic-NC), miR-30a-5p inhibitor and inhibitor-NC were synthesized by Ribobio (Guangzhou, China) and transfected into cells at a final concentration of 100 nM.

Quantitative real-time RT-PCR

Stem-loop qRT-PCR and general qRT-PCR methods were used to measure the expression levels of miRNAs and Beclin-1 mRNA, respectively. Total RNA from DEV-infected and uninfected DEF cells was extracted with TRIzol reagent (TIANGEN Biotech, Beijing) and quantified using a spectrophotometer (NanoDrop 2000). RNA (1000 ng) was reverse-transcribed to cDNA, and then 2 µl cDNA was used for real-time PCR amplification according to the kit manufacturer's (Thermo) instructions. The primers are listed in Table 1. Relative expression levels of miRNA and Beclin-1 mRNA were calculated using the 2^{-ΔΔCt} method. U6 and β-actin were used as respective endogenous controls.

Vector constructs and luciferase assay

MiR-30a-5p was predicted to target the DEF Beclin-1 3'UTR (nt 136,000–145,890) according to RNAhybrid and PITA software. The Beclin-1 3'UTR (nt 136,085–136,248), including the predicted miR-30a-5p binding site, was synthesized by TsingKe (Chengdu, China) and cloned into a pmirGLO vector (Promega, Madison, WI, USA) with SacI and XhoI sites, resulting in pmirGLO-WT-Beclin-1. Accordingly, the mutant 3'UTR of the Beclin-1 vector was constructed and named pmirGLO-MU-Beclin-1. For the luciferase assay, COS7 cells were

Table 1 Primers for analysis of gene expression by qRT-PCR

Primers	Sequence
RT-miR-146b-5p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACAACGCCTA
RT-miR-125b-5p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACCACAAGTT
RT-miR-30a-5p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACAGCTTCCA
RT-miR-27b-3p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACGCAGAACT
RT-miR-16c-5p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACCTCCAGTA
RT-miR-130b-3p	GTCGTATCCAGTGCCTGTCGTGGAGTCGGCAATTGCACTGGATACGACACGCCCTT
miR-146b-5p (F)	GCCGTGAGAACTGAATCCATA
miR-125b-5p (F)	GCCGTCCCTGAGACCCCTAA
miR-30a-5p (F)	GCCGTGTAACATCCTTGACTG
miR-27b-3p (F)	GCCGTTACAGTGGCTAAG
miR-16c-5p (F)	GCCGTAGCAGCACGTAATA
miR-130b-3p (F)	GCCGAGTCAATAATGAAA
UR-primer	CAGTGCCTGTCGTGGAGT
U6 (F)	CTCGCTTCGGCAGCACA
U6 (R)	GCGTGTCATCCTTGCGC
Beclin-1 (F)	AAGAGGTGCCTGGAGATCCT
Beclin-1 (R)	CGTCTCCAGTCTCTGAATC
β -Actin (F)	CCGGGCATCGCTGACA
β -Actin (R)	GGATTCATCATACTCTGCTTTGCT

seeded in 96-well plates and cotransfected with miR-30a-5p mimic, miR-NC, pmirGLO-WT-Beclin-1 and pmirGLO-MU-Beclin-1 with Lipofectamine 3000 (Invitrogen, Carlsbad, CA, USA). We performed a site-directed DLRA, and luciferase activity was measured at 36 h posttransfection according to the manufacturer's protocol (Promega, Madison, WI, USA).

Western blot analysis

The synthetic miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC were transfected into DEF cells with Lipofectamine 3000 (Invitrogen) according to the manufacturer's protocol. Meanwhile, the blank group (without) was set as the control group. Cells were infected with DEV at a multiplicity of infection (MOI) of 1.0 for 36 h. The cells were harvested and washed 3 times with cold PBS. The PBS was decanted and then 150 μ l RIPA lysis buffer (Solarbio, China) and 1.0 mM PSMF were added. After 30 min on ice and centrifugation at 12,000 g for 10 min, 25 μ l supernatant was mixed with 25 μ l 5 \times SDS loading buffer and boiled for 10 min. The protein samples were analyzed by 12% SDS polyacrylamide gel electrophoresis and transferred to a polyvinylidene difluoride (PVDF) membranes (Millipore, Billerica, MA) by electroelution. Membranes were blocked with 5% milk-TBS-Tween-20 for 2 h at room temperature and incubated with rabbit anti-LC3 (Proteintech, 14,600-1-AP), rabbit anti-p62/SQSTM1 (Cell

Signaling Technology, 5114), mouse anti- β -actin (Proteintech, 60,008-1-Ig), rabbit anti-Beclin-1 (Proteintech, 11,306-1-AP) and anti-CHV (UL41) antibodies overnight at 4 °C. Following incubation with HRP-conjugated goat anti-rabbit or anti-mouse IgG (Biodragon-Immunotech, China) as secondary antibody for 2 h at 37 °C, the immunoreactive bands were detected using an enhanced chemiluminescence kit (Solarbio, China). The amount of proteins was quantified by densitometry and normalized to β -actin, an internal standard.

Flow cytometry assay

DEF cells were seed in a 6-well plate at adensity of 1×10^6 cells per well. Cells were pretreated with control (without), miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then infected with DEV (MOI = 1.0) for 36 h. The cells were stained with Annexin V-fluorescein isothiocyanate (V-FITC) (BD Pharmingen, USA) and propidium iodide (PI) (BD Pharmingen, USA) according to the manufacturer's instructions, and the percentage of apoptotic cells was assayed by flow cytometry (FCM).

Cell viability analysis

miRNA toxicity tests were performed using the MTT assay kit (Sangon Biotech, Shanghai, China) according to the manufacturer's instructions. In brief, DEF cells were seeded in 96-well culture plates at a density of 1×10^5

cells per well. Cells were pretreated with control, miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then cultured in DMEM for 36 h, the cells were incubated in 100 μ l fresh culture medium containing MTT (0.5 mg mL⁻¹) for 4 h at 37 °C. The medium was replaced by 100 μ l formazan solubilization solution, and the absorbance was measured at 570 nm using a microplate reader (Bio-Rad).

DEV replication analysis

DEV viral copies were detected using qRT-PCR methods. DEF cells were seed in a 6-well plate at adensity of 1×10^6 cells per well. Cells were pretreated with control, miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then infected with DEV (MOI = 1.0). The cells were collected at the indicated times and stored at -80 °C for subsequent experiments. The DEV absolute quantitative curve was created as previously described methods [32, 33].

DEV titers were estimated by the median tissue culture infective dose (TCID₅₀). DEF cells were seed in 96-well plates at adensity of 1×10^5 cells per well. Cells were pretreated with control, miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then infected with DEV collected at the indicated times above. The plates were incubated for 5 days at 37 °C in a 5% CO₂ atmosphere. Cell pathological changes were observed under a light microscope and recorded. Viral titers were measured according to the Reed-Muench method [34].

Statistical analysis

Each assay was performed in three independent experiments. All experimental results are expressed as the mean \pm standard deviation (mean \pm SD) and were analyzed by the software GraphPad Prism (version 7.0). The statistical significance was assessed using Student's t-test. * $p < 0.05$ and ** $p < 0.01$ indicate significance.

Results

DEV infection induces downregulation of miR-30a-5p and promotes Beclin-1 mRNA expression

According to our previous high-throughput sequencing results [31], six miRNAs were chosen and detected using the stem-loop qRT-PCR method. The results showed that the expression levels of miR-30a-5p and miR-16c-5p were significantly downregulated in DEV-infected DEF cells at 24 hpi ($P < 0.01$) (Fig. 1a), and there were no obvious changes in the remaining four miRNAs (miR-146b-5p, miR-27b-3p, miR-130b-3p and miR-125b-5p) compared with uninfected cells (Fig. 1a). We detected the expression levels of miR-30a-5p and miR-16c-5p at 4, 12, 24, 36 and 48 hpi in DEV-infected group and uninfected group respectively. We found that the expression levels of miR-30a-5p and miR-16c-5p in the

uninfected group were 2.5 and 2.2 times as high as those in the infected group at 36 hpi respectively (Fig. 1b, c). In addition, qRT-PCR results showed that the expression level of Beclin-1 was upregulated in DEV-infected DEF cells and was 2.1- and 2.4-fold higher compared with the uninfected group at 36 hpi and 48 hpi (Fig. 1d).

Beclin-1 is a target of miR-30a-5p in DEF cells

A DLRA showed that pmirGLO-WT-Beclin-1 was significantly repressed by miR-30a-5p compared to the mimic-NC group ($p < 0.01$) (Fig. 2a, b). To further explore whether the downregulation of targets by miR-30a-5p was binding site-dependent, the binding sites of Beclin-1 were mutated to make the construct pmirGLO-MU-Beclin-1 vector (Fig. 2a). As expected, miR-30a-5p lost its repression effect on the mutant vector pmirGLO-MU-Beclin-1 (Fig. 2b). These results indicated that miR-30a-5p can directly target the DEF Beclin-1 mRNA at a 8-nucleotide complementary seed sequence.

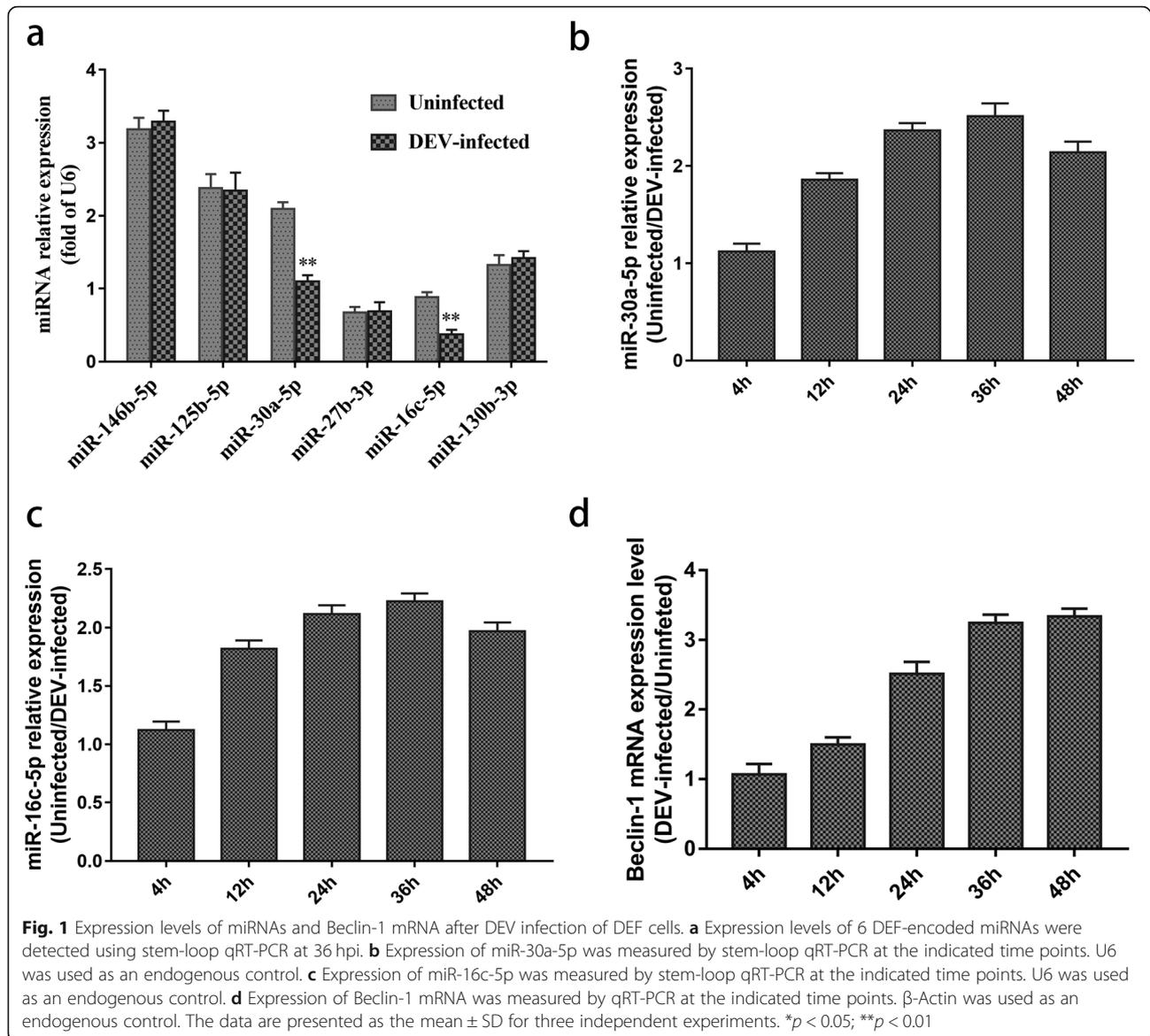
miR-30a-5p suppresses Beclin-1 expression and autophagy is induced by DEV

To further explore whether there is the negative regulation relationship between miR-30a-5p and Beclin-1 in DEV-infected DEF cells, DEF cells were transfected with control, miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then infected with DEV (MOI = 1.0) for 36 h. The mRNA expression levels of miR-30a-5p and Beclin-1 were evaluated by qRT-PCR. As expected, Transfection of the miR-30a-5p mimic significantly increased the miR-30a-5p expression and decreased the mRNA level of Beclin-1. Consistently, miR-30a-5p inhibitor had the opposite effects on expression in miR-30a-5p and Beclin-1 ($P < 0.01$) (Fig. 2c, d).

Next, we observed the effect of miR-30a-5p on autophagy. Western blot analysis showed that miR-30a-5p over-expression significantly decreased Beclin-1 protein expression. The ratio of LC3-II/LC3-I was significantly downregulated and the p62 protein level was significantly upregulated in the miR-30a-5p group, suggesting that autophagy was inhibited by miR-30a-5p regulating Beclin-1 (Fig. 3a, b). While the miR-30a-5p inhibitor increased Beclin-1 protein level and decreased p62 expression. An increased ratio of LC3-II/LC3-I was observed for miR-30a-5p inhibitor-treated group compared to inhibitor-NC group, indicating that autophagy activity was promoted (Fig. 3c, d). Specific viral proteins with UL41 antibody indicated the process of viral infection. These results revealed that the miR-30a-5p inhibited autophagy by regulating Beclin-1 in DEV-infected cells.

miR-30a-5p promotes DEV-induced DEF cell apoptosis

A FCM was used to detect DEV-induced DEF cell apoptosis after transfecting control, miR-30a-5p mimic, miR-



NC, miR-30a-5p inhibitor and inhibitor-NC. The results revealed that the percentage of apoptotic DEF cells was significantly increased at 36 hpi in the miR-30a-5p mimic group. However, the transfection of miR-30a-5p inhibitor significantly decreased the percentage of apoptotic cells in DEV-infected cells (Fig. 4a). These results showed that miR-30a-5p promoted DEV-induced apoptosis of DEF cells.

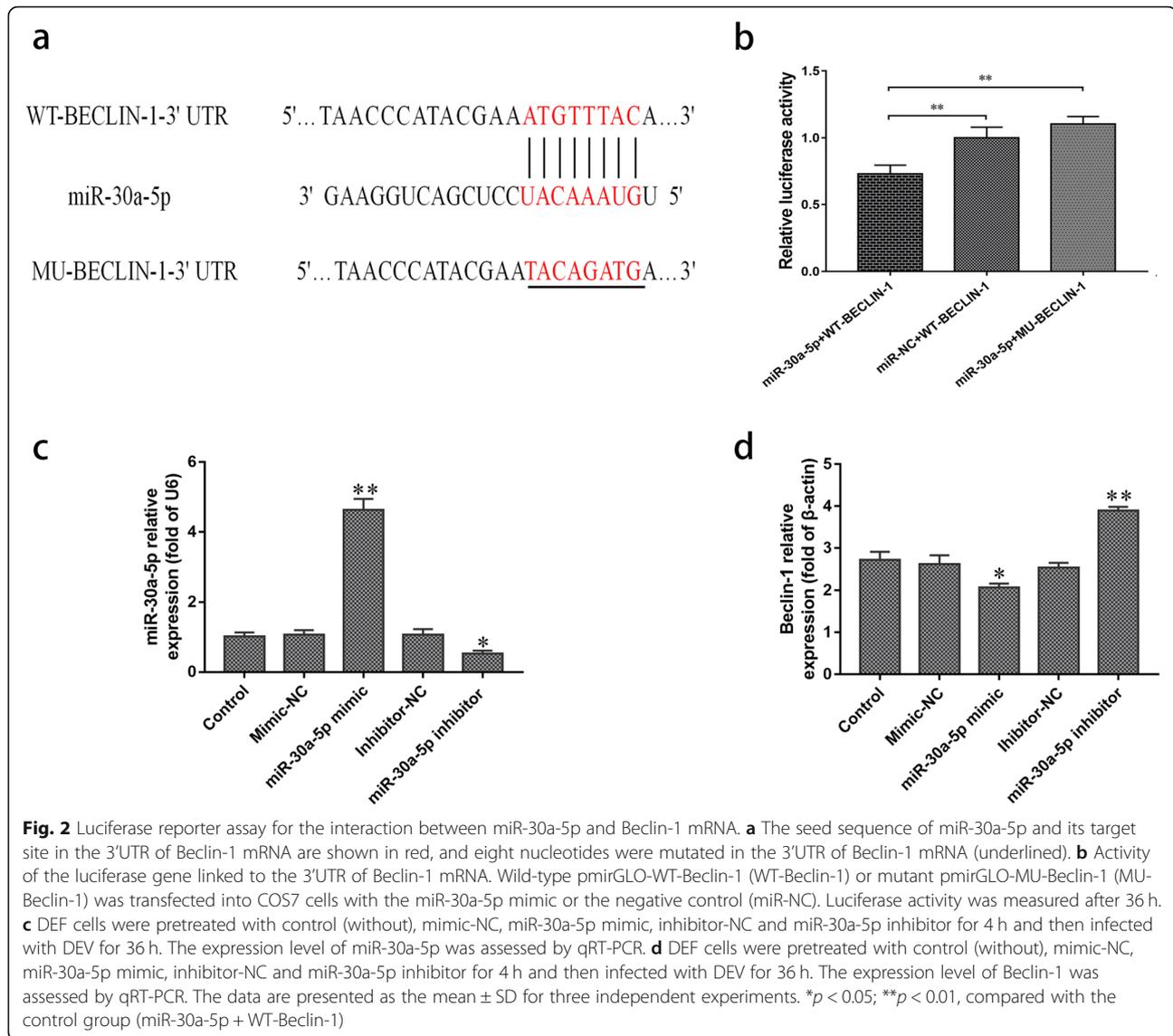
Cell viability unaffected by miRNAs treatment

The miRNAs might have influenced cell viability and affected our results. The effects on cell viability of the miRNAs used in this study were measured by MTT assays. The viability of treated cells was almost equal to that of control cells, so the miRNAs treatments did not affect DEF cell viability (Fig. 4b).

miR-30a-5p overexpression reduces DEV replication

To identify the potential function of miR-30a-5p in the viral replication process, DEF cells were transfected with control, miR-30a-5p mimic, miR-NC, miR-30a-5p inhibitor and inhibitor-NC for 4 h and then infected with DEV (MOI = 1.0). The genome copy number of DEV was measured by qRT-PCR at the indicated time points, and overexpression of miR-30a-5p strongly suppressed DEV replication at 12, 24, 36, 48 and 60 hpi, while the miR-30a-5p inhibitor significantly triggered DEV replication at the same time points (Fig. 4c).

Viral yields were determined by TCID₅₀ at the indicated time points, and titers of progeny viruses collected from miR-30a-5p mimic group were lower than from control cells at 24, 36, 48 and 60 hpi. While viral titers



in miR-30a-5p inhibitor group were significantly upregulated compared with the control group (Fig. 4d).

Discussion

There have been several reports about miR-30a-5p target genes and their involvement in pathophysiological processes. For example, miR-30a-5p promotes the replication of porcine circovirus type 2 through enhancing autophagy by targeting 14-3-3 [15]. MiR-30a-5p downregulation contributes to the chemoresistance of osteosarcoma cells by activating Beclin-1-mediated autophagy [35]. The expression of miR-30a-5p is significantly downregulated in human colorectal cancer (CRC) tissues and CRC cell lines and may be a potential candidate target for CRC therapy [36]. Low expression of miR-30a-5p induces the proliferation and invasion of oral cancer by promoting the expression of FAP (*Homo sapiens*

fibroblast activation protein α), and miR-30a-5p might be a new therapeutic target for oral cancer treatment [37]. In our previous study, miRNA expression profiles of virus and host were determined and analyzed in virulent DEV-infected DEF cells. The expression level of miRNA-30a-5p was significantly downregulated during the whole process of DEV infection [31], and similar results were confirmed by stem-loop qRT-PCR in this study (Fig. 1b). We speculate that miR-30a-5p plays an important role in regulating host-virus interactions.

Autophagy is an essential pathway for cellular homeostasis. Many studies have confirmed that some viruses can induce cells manipulate autophagy to promote their survival and replication [38, 39]. Examples include hepatitis C virus [40], egg drop syndrome virus (EDSV) [41], avian reovirus [42] and influenza A virus [43]. Recent study demonstrated that autophagy induced by DEV

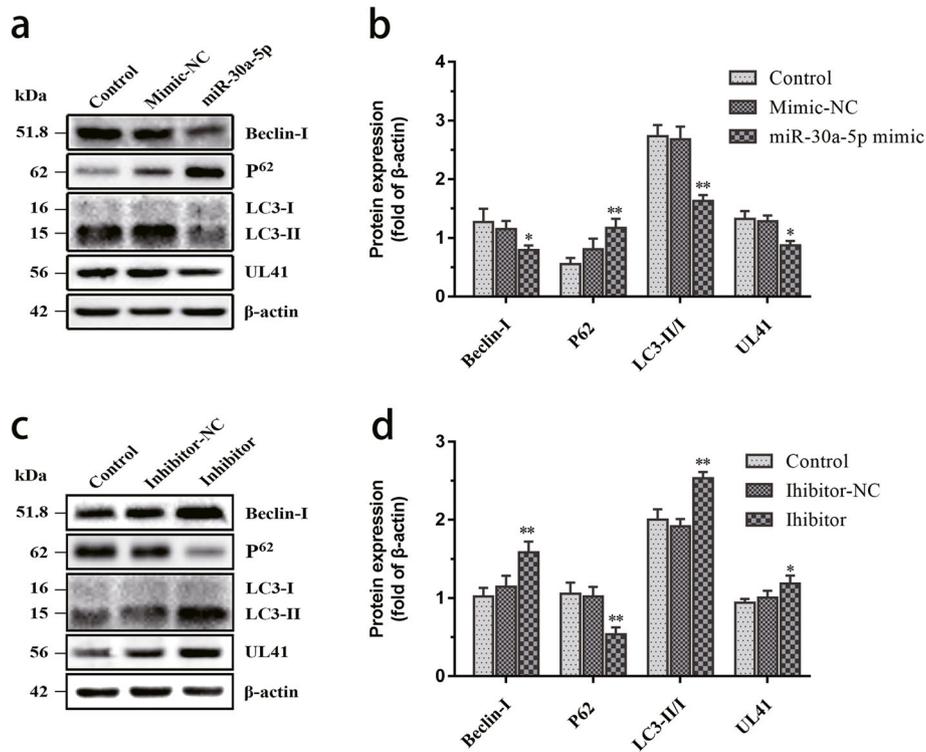
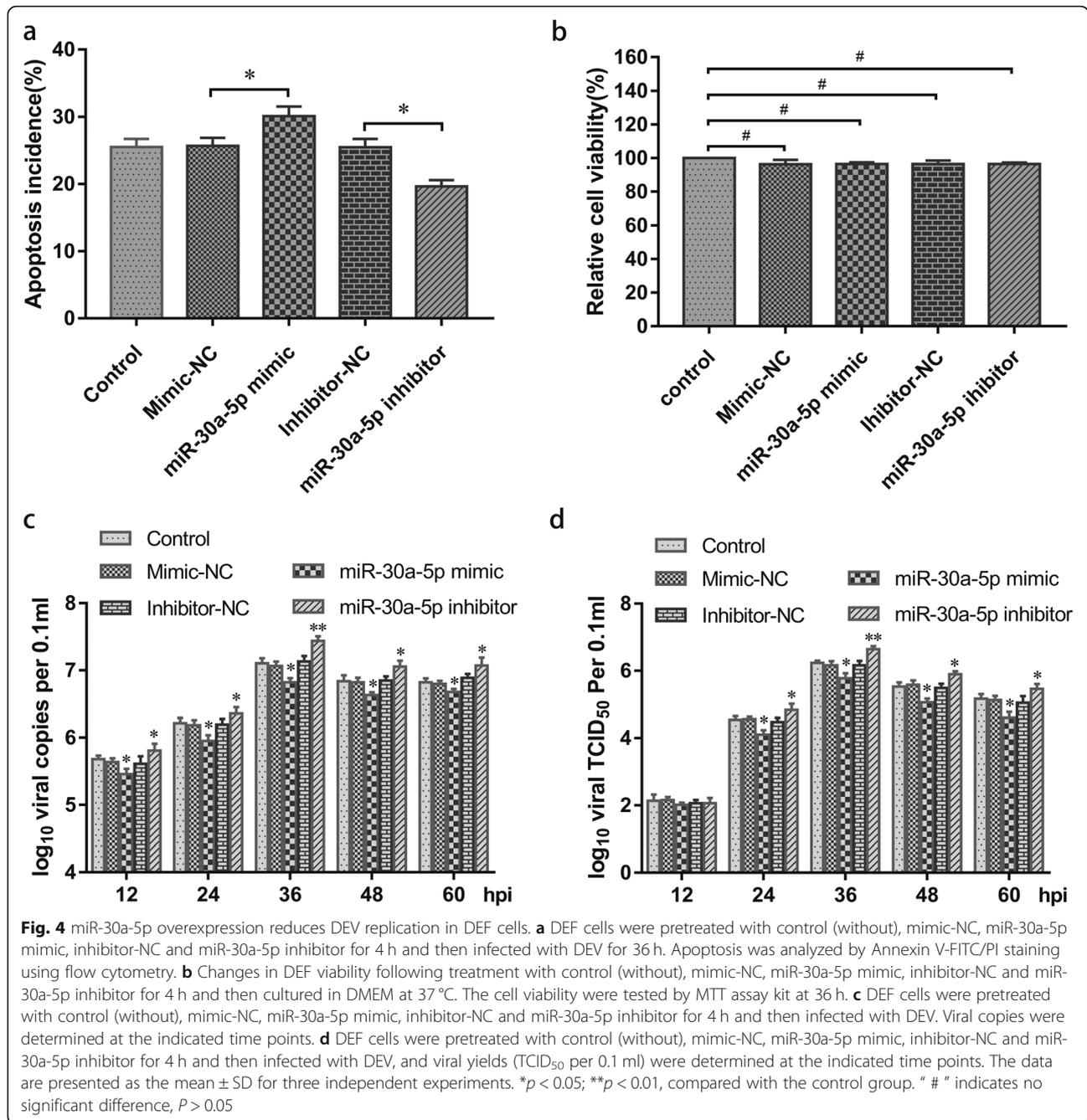


Fig. 3 miR-30a-5p overexpression suppresses Beclin-1 expression and DEV-induced autophagy. **a** DEF cells were pretreated with control (without), mimic-NC and miR-30a-5p mimic for 4 h and then infected with DEV. Cells were lysed and blotted with antibodies against Beclin-1, p62, LC3, β -actin and UL41 at 36 hpi. **b** The optical densities of each protein band from (a) were measured by densitometric scanning, and the optical density ratios of Beclin-1/ β -actin, P62/ β -actin LC3II/I, and UL41/ β -actin were calculated. **c** DEF cells were pretreated with control (without), inhibitor-NC and miR-30a-5p inhibitor for 4 h and then infected with DEV. Cells were lysed and blotted with antibodies against Beclin-1, p62, LC3, β -actin and UL41 at 36 hpi. **d** The optical densities of each protein band from (c) were measured by densitometric scanning, and the optical density ratios of Beclin-1/ β -actin, P62/ β -actin LC3II/I, and UL41/ β -actin were calculated. The data are presented as the mean \pm SD for three independent experiments. * $p < 0.05$; ** $p < 0.01$, compared with the control group

infection positively promotes viral replication [21]. However, the regulating relationship of autophagy in DEV-infected DEF cells is still unclear. In this study, our data showed that the expression of miR-30a-5p was downregulated and that Beclin-1 was upregulated after CHV infection of DEF cells. The Beclin-1 gene is recognized as a critical regulatory gene during autophagosome formation and maturation [44] and plays important roles in the replication of some viruses. For example, the replication of these three viruses (NDV, CSFV and DEV) was inhibited by siRNA knockdown of Beclin-1 gene level, which is required for autophagy [13, 14, 21]. Previous studies have demonstrated that miR-30a-5p regulates autophagy activity by targeting Beclin-1 mRNA [45–47]. Therefore, we speculated that there is the negative regulation relationship between Beclin-1 and endogenous miR-30a-5p in DEV-infected DEF cells. To confirm the regulation of miR-30a-5p on Beclin-1, bioinformatics analysis was performed using RNAhybrid software [31]. The results showed that miR-30a-5p was predicted to target the 3'-UTR region of Beclin-1. DLRA confirmed

that overexpression of miR-30a-5p markedly reduced the luciferase level from Beclin-1 (Fig. 2a, b).

Western blot analysis demonstrated that overexpression of miR-30a-5p decreased Beclin-1 protein level and the ratio of LC3-II/LC3-I, enhanced the p62 protein level, which are related to autophagy. The expression of UL41 protein decreased (Fig. 3a, b). Whereas miR-30a-5p inhibitor attenuated DEV-induced cell autophagy and reversed the effect of miR-30a-5p. Both Beclin-1 and the ratio of LC3-II/LC3-I significantly increased in miR-30a-5p inhibitor group. The p62 protein level decreased and the expression of UL41 protein increased (Fig. 3c, d). The TCID₅₀ and the viral copy test confirmed that DEV titers were significantly decreased in the miR-30a-5p mimic group compared with control group. Whereas transfection of miR-30a-5p inhibitor promoted DEV replication during the whole process of DEV infection (Fig. 4c and d). Our results were consistent with a previous report on autophagy induced by DEV [21], suggesting a key role for the miR-30a-5p/autophagy loop in DEV infection. These results strongly



suggested that overexpression of miR-30a-5p decreased DEV replication by suppressing Beclin-1-mediated autophagy. Therefore, it is reasonable to conclude that downregulation of miR-30a-5p contributes to DEV replication by upregulating Beclin-1-mediated autophagy.

Apoptosis regulates embryonic development, cell turnover, and the immune response against tumor or virus-infected cells [48, 49]. Virus-induced cell apoptosis is involved in the pathogenesis of many viral infections [50, 51]. Our laboratory has discovered that DEV can induce apoptosis in the thymus, spleen and pancreatic lymphocytes

of adult ducks and can cause apoptosis in DEFs in vitro [52, 53], and further confirmed that the mRNA levels and enzymatic activities of caspase-3, caspase-7 and caspase-9 were significantly increased during DEV-induced cell apoptosis [53]. Recent study has also reported miR-30a-5p can promote doxorubicin-induced osteosarcoma cell apoptosis by increasing the expression of cleaved caspase-3, and further certified that miR-30a-5p promotes chemotherapy-induced osteosarcoma cell apoptosis via repressing Beclin-1-mediated osteosarcoma autophagy [35]. In our study, flow cytometry demonstrated that

overexpression of miR-30a-5p enhanced DEV-induced cell apoptosis (Fig. 4a). Nevertheless, miR-30a-5p inhibitor attenuated DEV-induced cell apoptosis and reversed the effect of miR-30a-5p (Fig. 4a). However, whether or not the miR-30a-5p increased DEV-induced cell apoptosis by suppressing Beclin-1-mediated autophagy requires further confirmation.

Conclusions

The results of this study showed that the miR-30a-5p/autophagy loop plays an important role in DEV infection, and this is the first report of miR-30a-5p in herpesvirus-induced autophagy. Our work confirms the existence of a novel regulatory pathway controlled by miR-30a-5p and its direct target, Beclin-1, that regulates DEV replication in DEFs. Therefore, miR-30a-5p and its target gene pathway may represent new treatment methods for duck viral enteritis disease.

Abbreviations

CHV: Chinese virulent; CSFV: Classical swine fever virus; DEV: Duck enteritis virus; EDSV: Egg drop syndrome virus; IRS: Inverted repeated sequences; NDV: Newcastle disease virus; PCV2: Porcine circovirus type 2; PRRSV: Porcine reproductive and respiratory syndrome virus; qRT-PCR: Quantitative real-time reverse transcription PCR; TRS: terminal repeated sequences; UL: Unique long region; US: Unique short region; VZV: Varicella-zoster virus

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Authors' contributions

Conceived and designed the experiments: RYJ XLW. Performed the experiments: XLW. Analyzed the data: RYJ, MSW, SC, MFL, DKZ, XXZ, QY, YW, ZQY, SQZ, JH, YLY, LCP, BT, MUR, XYZ, ACC. Contributed reagents and materials: LZ, YYL. Wrote the paper: XLW, RYJ. All authors read and approved the final manuscript.

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Availability of data and materials

The datasets used or analysed during this study are included within the article.

Ethics approval and consent to participate

The study was approved by the Animal Ethics Committee of Sichuan Agricultural University (approval No. XF2016-17).

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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