The construction and application of a cell line resistant to novel subgroup avian leukosis virus (ALV-K) infection

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Abstract A novel avian leukosis viruses (ALV) subgroup named ALV-K was recently isolated from Chinese indigenous chickens which is different from the subgroups (A to E and J) that have previously been reported to infect chickens. More and more ALV-K strains have recently been isolated from local breeds of Chinese chickens. However, there are no more effective diagnostic methods for ALV-K other than virus isolation followed by envelope gene sequencing and comparison. Viral infection can be blocked through expression of the viral receptor-binding protein. In this study, we have engineered a cell line, DF-1/K, that expresses ALV-K env protein and thereby confers resistance to ALV-K infection. DF-1/K can be used in combination with the ALV-K susceptible cell line DF-1 as a specific diagnostic tool for ALV-K and provides a good tool for further research into the molecular mechanisms of interaction between ALV-K env protein and the host cell receptor.

Introduction

Avian leukosis viruses (ALVs), which are taxonomically classed as members of the genus *Alpha retrovirus* of the family *Retroviridae*, are divided into 10 subgroups (A to J) based on viral envelope glycoprotein antigenic structure, host range and receptor interference [1]. Members of only six of these subgroups (A-E and J) infect chickens. The main characteristics of avian leukosis (AL), which has resulted in huge economic losses to the poultry industry in China, are serious immunosuppression, growth retardation and tumor-induced mortality. In recent years, a novel ALV subgroup named ALV-K was isolated from indigenous Chinese chickens [2–4]. ALV subgroups are determined based on the gp85 envelope protein [5–7], and based on gp85 amino acid sequence comparisons ALV-K is different from all other known subgroups that infect chickens (A-E and J) [2–4]. Interestingly, several fowl glioma-inducing viruses (FGV) [8–11] reported in Japan and several ALV-K strains (JS11C1, JS11C2, JS11C3, GDFX0601, GDFX0602 and GDFX0603) isolated from native Chinese chicken breeds [2, 3] show over 90 % identity to TW3593 [12] in their gp85 amino acid sequences. Based on phylogenetic analysis of their gp85 genes, these viruses form one large clade that is parallel to other ALV groups. The gp85 gene of TW3593 is different from all other known subgroups that infect chickens (A-E and J) [2–4]. These observations suggest that ALV-K might have existed locally in Chinese chickens for a long time [2–4]. Recently, more and more ALV-K strains have been isolated...
from local Chinese chicken breeds [13]. These ALV-K isolates replicate more slowly and are less pathogenic than some other ALV strains [4]. The low replication of the ALV-K strain results in a low level of p27 antigen expression, so that chickens infected with exogenous ALV-K are difficult to detect and cause widespread spread, causing interference to ALV control. Therefore, further study of the biological characteristics, genome structure and function of ALV-K, as well as the molecular mechanisms of infection are necessary.

Retroviruses efficiently infect only cells that express specific receptors that interact with the viral envelope glycoproteins [14]. Viral infection and pathogenesis are initiated by the binding of this viral surface protein to its cellular receptor. Indeed infection can be blocked by over expression of the viral receptor-binding protein [15]. Different ALV subgroups use distinct cellular receptors [16]. The ALV envelope protein, a glycoprotein encoded by the virus envelope (env) gene, is the main viral surface protein and determines the virus host range and induces the production of host neutralizing antibodies [16]. Specific interactions between the viral env protein and cell surface receptors are necessary for viral entry into host cells. After being infected with a specific ALV subgroup, viral receptors on the cell surface can be blocked by the viral envelope (env) glycoprotein. These infected cells show superinfection resistance and can specifically resist infection with the same subgroup virus again [16].

Different methods have been established for detecting exogenous ALV, including PCR, real-time PCR, immunofluorescence assay (IFA), traditional virus isolation plus an antigen-capture enzyme-linked immunosorbent assay (ELISA) for group-specific p27 antigen of ALV, as well as loop-mediated isothermal amplification (LAMP), and quantitative competitive reverse transcription PCR (QC-RT-PCR) [17–21]. However, each of these methods has limitations. Some methods were only developed for the detection of ALV-J and ALV-A/B. At present, ALV-K diagnosis relies mainly on virus isolation followed by envelope gene sequencing, and there is lack of other diagnostic methods. For instance, IFA-based diagnostics are based on a specific monoclonal antibody, which is not available for ALV-K. Therefore, IFA cannot be used as a specific diagnostic tool for ALV-K. A genetically engineered cell line resistant to ALV-J infection has been developed and applied to screen large numbers of ALV-J field samples [22] and to efficiently identify chicken Annexin A2 (chANXA2) as a novel receptor for retrovirus ALV-J [15]. In this study we induced super infection resistance to the ALV env protein through stable expression of the ALV-K env protein in DF-1 cells, obtaining a cell line that can resist ALV-K infection. This cell line was used to evaluate clinical plasma samples and investigate the distribution of ALV-K infection in native Chinese chickens. This is the first report of a cell line resistant to infection by a novel ALV subgroup (ALV-K).

**Materials and methods**

**Viruses and cell lines**

The DF-1 fibroblastic cell line (American Type Culture Collection, Manassas, VA) was used for virus culture. The cells were grown in Dulbecco’s modified Eagle medium (DMEM; GIBCO, USA) supplemented with 10% FBS (Fetal bovine serum, GIBCO, USA) and maintained in DMEM supplemented with 2% FBS at 37 °C in a humidified atmosphere containing 5% CO₂. The ALV strains GDFX0601 (GenBank accession number KP686142), GDFX0602 (GenBank accession number KP686143) and GDFX0603 (GenBank accession number KP686144) were isolated from a native Chinese yellow feather broiler breeder in southern China in 2014, propagated in DF-1 cells and maintained in our laboratory. The ALV-J subgroup strain CHN06 (GenBank accession number HQ900844) was isolated and identified by our laboratory [23]. Two field ALV-K isolates, GD-147 and GD-179, were isolated from a native Chinese yellow feather broiler breeder in southern China in 2015. The titers of the GDFX0601, GDFX0602, GDFX0603 and CHN06 strains were determined by ELISA and are presented as TCID₅₀ml⁻¹ calculated using the Reed-Muench method [24].

**Antibodies and reagents**

The Flag M2, mouse anti-GFP, mouse Anti-β-actin, and FITC-labeled goat anti-mouse IgG antibodies used in this study were purchased from Sigma (CA, USA). The primary antibodies used for ALV-K detection were single factor antisera of ALV-K and were provided by Dr Peng Zhao, Shandong Agricultural University. Zeocin was purchased from Invitrogen (Shanghai, China). BamHI and NotI restriction enzymes and T4 ligase were purchased from New England BioLabs.

**Plasmid construction**

The ALV-K env gene was amplified by PCR using the following gene-specific primers: P1: 5’-CGCGGATCCGCG ACCATGGAAGCGGATCATAAAGGCGATTCTGACTGGA TAC-3′ and P2: 5’-AAGGAAAAAGCGGCCGCTTACAC TGCTCCATTTCGG-3′. The primers contain protective bases, restriction enzyme cutting sites (italicized letters), a *Kozak* sequence (underlined letters) and a leader sequence (italicized letters) to improve translational efficiency. PCR was performed in a 50 μl reaction mixture that consisted of template DNA (5μl), 10×PCR buffer (TaKaRa,
Dalian, China), 1µM each of forward and reverse primers, 2mM MgCl₂, 100mM of each deoxynucleoside triphosphate (dNTP), and 1 unit of LA TaqTM DNA Polymerase (TaKaRa, Dalian, China). PCR thermocycling profiles included an initial denaturation for 3 min at 94 °C, followed by 30 cycles of amplification (94 °C for 30 s, 55 °C for 1 min, and 72 °C for 2 min), as well as a final extension of 72 °C for 8 min. The ALV-K env PCR product was purified using the QIAEX II gel extraction kit (Qiagen, Hilden, Germany), sequenced (Invitrogen, Shanghai, China) and subcloned into the pMD-18T vector (TaKaRa, Dalian, China). The ALV-K env gene was then cloned into the eukaryotic expression vector pcDNA3.1 using the BanHI and NotI sites. Three pcDNA-env-K vectors were sequenced and all had the predicted nucleotide sequences. The pcDNA-env-K-flag-EGFP vector, which contains flag and EGFP tags, was constructed by PCR amplification of the EGFP fragment which was ligated with the env gene using NotI and XbaI restriction enzyme sites. The fusion fragment was then cloned into the pcDNA3.1 vector.

**Cell transfection and cell screening**

The day before transfection, DF-1 cells grown in a monolayer were digested with 0.25% trypsin (GIBCO,USA), and the cells were then adjusted to a density of 1.7 × 10⁵ cells/mL in Dulbecco’s modified Eagle’s medium (GIBCO,USA) with 10% FBS (GIBCO,USA). These were plated in 6-well cell culture plates at 37 °C with 5% CO₂ until they reached approximately 80% confluence. Transfection of the pcDNA-env-K plasmid, pcDNA-env-K-flag-EGFP plasmid and pcDNA3.1/Zeo(+) plasmid into DF-1 cells was performed using Lipofectamine 3000 (Invitrogen, Shanghai, China) according to the manufacturer’s protocol. The empty plasmid pcDNA3.1/Zeo(+) served as a negative control. After 48 hours, the DF-1 cells grown in monolayer as well as cells in one of the 6-well cell culture plates were digested with 0.25% trypsin (GIBCO, USA), and the cells with the media (DMEM + 15%FBS + 200 µg/mL zeocin) were seeded into 24-well tissue culture plates (500 µL/well). The transfected DF-1 cells were selected for resistance to Zeocin. The following day, cells were treated with 500 µL/well of media containing Zeocin (DMEM+15%FBS+200µg/mL zeocin) and this media was replaced every three days. The Zeocin-resistant cells were passaged for 60 generations and then frozen. After 3 months, these cells were refreshed and cultured in medium free of Zeocin.

**Routine PCR and the real-time PCR assay**

Routine PCR tests were carried out with genomic DNA extracted from the ALV-K-resistant cell line, designated as DF-1/K cells, as well as DF-1 cells. The DF-1 cells served as a negative control. The specific primers, reaction mixture, thermocycling profiles were as described above. The PCR product was purified using the Omega Gel Extraction kit (Omega Bio-tek). Total cellular RNA was extracted from DF-1/K cells and DF-1 cells with the RNAfast200 kit (Fastagen, Shanghai, China), followed by cDNA synthesis with the RevertAid First strand cDNA synthesis kit (Fermentas, Canada) according to the manufacturer’s instructions. The cDNA was then used for routine PCR and real-time PCR amplification. Real-time reverse transcription (RT)-PCR was done with primers designed for the envelope gene and gene-specific primers synthesized by TaKaRa Company (Dalian, China): F: 5’-CCCCTGCTATTTAGGCAAAGCT-3’, R:5’-AGTTGGCAACCTTGGAGAA-3’, Probe:Fam-5’-CCATGTAGCACCACACAGAA-3’–Eclips.

DNA sequences were determined by Invitrogen (Invitrogen, Shanghai, China). For all reactions, PCR amplification and DNA sequencing were carried out at least twice, independently, to avoid PCR errors. Real-time PCR was performed on an ABI 7500 Real-time PCR System (Applied Bio systems) using Premix ExTaq (Probe qPCR) reagents (TaKaRa, Dalian, China) according to the manufacturer’s specifications. Fluorescent signals were recorded during the elongation step. The β-actin gene served as a reference gene (primers: F: 5’-CCAGCCATGTATGCTACCC-3’, R:5’-CAC CTACAGAAGTCACAC-3’, Probe: Fam-5’-CTGTGC TGTCCCTGTAGCCTGG-3’–Eclips). The relative expression level of the env gene was normalized to GADPH. Finally, real-time quantitative PCR analysis was carried out using the 2−ΔΔCT method [25].

**Indirect immunofluorescence assay (IFA)**

DF-1/K and DF-1 cells were washed with PBS, fixed with cold acetone–alcohol (3:2) for 20 min, washed with PBS again, and then allowed to air-dry. The cells were then incubated with a single factor anti-serum for ALV-K at 37 °C for 1 h, washed three times with PBS, and further incubated with goat anti-mouse IgG conjugated with FITC (Sigma, USA) at 37 °C for 1 h. After three washes with PBS, the cells were observed using fluorescence microscopy.

**Western blot analysis**

DF-1/K cells and DF-1 cells in 150 mm dishes were harvested by scraping with a rubber policeman and homogenized with NP-40 lysis buffer containing 25 mM Tris, 150 mM NaCl, 1 mM EDTA, 1% NP-40, 5% glycerol (pH 7.4) and a protease inhibitor cocktail (Roche). The lysates were collected and incubated on ice for 10 min. Lysates were cleared by centrifugation at 10,000× g for 5 min at 4 °C. The supernatants were analyzed for total protein content with the BCA protein assay kit (Fermentas,
Life Technologies). Total protein (20μg) was resolved by 12% SDS-PAGE and transferred onto nitrocellulose membranes (Whatman, Maidstone, UK). Membranes were blocked with 5% (w/v) skim milk for 1 h at 37 °C, and then incubated overnight at 4 °C with a specific mouse anti-gp85 single factor anti-serum for ALV-K and β-actin (Santa Cruz, sc-1616-R). The β-actin protein served as a reference. After three rinses with PBS Tween20 (PBST) buffer, the membranes were incubated at 37 °C for 1 h with IRDye 800-conjugated anti-mouse IgG secondary antibody (1:10,000; Rockland Immunochemicals, Limerick, PA, USA) diluted in PBS. Membranes were washed three times with PBST, then visualized and analyzed with an Odyssey infrared imaging system (LI-COR Biosciences, Lincoln, NE, USA).

Immune electron microscopy

The DF-1 and DF-1/K cells were repaired and sliced with a slice thickness of 75 nm, according to the preparation method for ordinary electron microscopy samples. The ultra-thin slices were transferred to a nickel mesh with Fang Hua film (to ensure slice continuity) and then incubated with goat anti-mouse flag antibodies (Sigma, USA), goat anti-mouse EGFP antibodies (Sigma, USA) and single factor anti-serum for ALV-K (gifts from Dr. P. Zhao, College of Shandong Agricultural University) for 1 h at 37 °C. After washing the sections with PBSA solution 6 times, the samples were incubated with 10 nm colloidal gold-labeled goat anti-mouse IgG (Sigma, USA) for 1 h at 37 °C. The sections were washed with ultra-pure water and dried at room temperature. Finally, the samples were examined under a JEM-2010HR transmission electron microscope (JEPL, Japan).

Antiviral experiment

Five different virus titers ranging from $10^1$ TCID$_{50}$ to $10^5$ TCID$_{50}$ per 0.1 ml of ALV-K (GDFX0601), ALV-A (GD13), ALV-B (CD08) and ALV-J (CHN06) were inoculated per well in 24-well cell culture plates containing 1 mL 1.7 × 10^5 cells/well DF-1/K cells, all in triplicate. Three wells on each plate served as negative controls. Each dilution of virus was performed in triplicate. After the inoculum was removed, maintenance medium containing DMEM with 2% FBS was added, and the plates were incubated at 37 °C and 5% CO$_2$ for another 6 days. The supernatant fluid was then harvested for ALV p27 antigen ELISA detection (ALV-p27 Ag Test kit, IDEXX, Inc., Westbrook, MA). A mock-infected DF-1 cells group was established in parallel as a control.

Results

Cell line screening

To obtain Zeocin-resistant cells, cells transfected with the pcDNA-env-K plasmid or the pcDNA-env-K-flag-EGFP plasmid were grown in medium containing 200 μg/mL Zeocin. In our culture system, the DF-1 cells formed a single cell colony within 10-15 d (Fig. 1A), and this single cell colony appeared to increase in size over the next 6-10 d (Fig. 1B, C, D, E). The cells grew to near confluence after approximately 21 days in culture (Fig. 1F). After three weeks, the cells were washed with PBS, digested with 0.25% trypsin (Gibco), and then plated in 6-well cell culture plates at 37 °C and 5% CO$_2$ until they grew to confluence. The Zeocin-resistant cells were cultured in medium containing Zeocin, passaged continuously for 60 generations and then frozen. After 3 months, these cells were refreshed and cultured in medium free of Zeocin, and the env gene or env protein expression in the transfected DF-1 cell was examined.

PCR detection of the env gene in DF-1/K cells

PCR was used to detect the ALV-K env gene in the first passage of DF-1/K cells (Fig. 2A) and to confirm ALV-K env gene remained stable in the genome after 20-60 passages (Fig. 2B). A 1791bp amplicon consistent in length with the ALV-K env fragment was amplified from all DF-1/K DNA samples. This PCR product was purified, and sequencing analysis verified that the fragment corresponded to ALV-K env (Fig. 2B). The results demonstrate that the ALV-K env fragment remained genetically stable in DF/K cells during passage.

Analysis of env gene transcription

Viral envelope gene transcription levels in DF-1/K cells was monitored by routine PCR and real-time RT-PCR with primers designed against the ALV-K env gene. RNA was extracted from DF-1/K and DF-1 cells and reverse transcribed into cDNA, which was then used for routine PCR and real-time PCR amplification. As shown in Fig. 3A, a 1791bp amplicon consistent in length with the ALV-K env fragment was amplified from all DF-1/K DNA samples. This PCR product was purified, and sequencing analysis verified that the fragment corresponded to ALV-K env (Fig. 2B). The results demonstrate that the ALV-K env fragment remained genetically stable in DF/K cells during passage.
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Indirect immunofluorescence (IFA) testing of ALV-K env gene expression

ALV-K env gene expression was confirmed by detecting the ALV-K env protein using IFA (Fig. 4). The cells were then incubated with a single factor anti-serum for ALV-K, and further incubated with goat anti-mouse IgG conjugated with FITC. No green fluorescence in the cytoplasm could be observed in DF-1 cells, but the green fluorescence signal was bright in DF-1/K cells. This indicates

Fig. 1 Zeocin selection of cell lines. (A) After Zeocin selection, the transfected DF-1 cells formed a single cell colony within 10-15d. The single cell colony appeared to increase in size over the following 6-10d. The differences in the size of the cell clones are shown on days 16 (B), 18 (C), 19 (D) and 20 (E). (F) The cells had nearly grown to confluence after approximately 21 days in culture. Magnification is 400× for (A) and 200× for (B-F)
that the exogenous ALV-K env gene was successfully expressed in DF-1/K cells.

**Western blot analysis of env protein expression in DF-1/K cells**

DF-1 and DF-1/K cells grown to monolayer were harvested and lysed. Env proteins within cell lysates were detected by Western blotting using mouse anti-gp85 single factor anti-serum for ALV-K, and IRDye 800-conjugated anti-mouse IgG was used as the secondary antibody. A protein of 90kDa was observed in DF-1/K cells but not DF-1 cells (Fig. 5).

**Antiviral experiment**

Representative viruses from ALV subgroups A (GD13), B (CD08), J (CHN06) and K (GDFX0601) were used to determine the ability of DF-1/K and DF-1 cells to resist infection. All of the viruses, ALV-A, ALV-B, ALV-J and ALV-K, were capable of infecting and replicating in DF-1 cells. In contrast, only subgroups A, B, J viruses infected the DF-1/K cells, while viruses from subgroup K were largely blocked from infecting these DF-1/K cells. As shown in Fig. 6A–D, DF-1/K cells inhibited the replication of ALV-K but not that of ALV-A, ALV-B, ALV-J viruses based on ELISA measurements of viral p27 protein expression. In order to further determine its antiviral effect on ALV-K, four different
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Field isolates of ALV-K viruses were used to infect DF-1 and DF-1/K cells. All four field isolates were capable of infecting DF-1 cells, but were largely blocked from infecting DF-1/K cells (Fig. 6E). The anti-virus assay showed that DF-1/K cells can have resistance to infection at a viral dose of 1 x 10^4 TCID50 ALV-K, and at lower doses infection was completely blocked. When the ALV-K infection dose reached 1 x 10^5 TCID50, the ability of ALV-K to infect DF-1/K cells was still strongly inhibited (Fig. 6A).

**Immune electron microscopy**

To further understand and monitor the molecular mechanisms underpinning the blocking of cell receptors by membrane-bound envelope glycoproteins, we determined the localization of the ALV-K env fusion protein, which contains flag and EGFP tags, in DF-1/K cells using colloidal gold immune electron microscopy (Fig. 7). The ALV-K fusion protein was detected with goat anti-mouse flag antibodies, goat anti-mouse EGFP antibodies and single factor anti-serum for ALV-K individually. As the secondary antibody, 10nm colloidal gold-labeled goat anti-mouse IgG was used. As a negative control, TBS (pH 7.4) was used in place of the primary antibody of control group. After uranyl acetate and lead citrate staining, images of the samples were obtained with a JEM-2000EX transmission electron microscope. Immunogold particles were observed in the cell membrane. In the controls, there were no immunogold particles in the cell membrane area.

**Discussion**

ALVs are RNA viruses that have a high level of genetic variation due to the high error rate of their polymerases and a high recombination rate. They have spread all over the world and caused enormous economic losses in the international poultry industry. ALVs have been divided into 10 different subgroups, subgroup A to J, based on receptor interference patterns, host range, and neutralization by antibodies [26]. So far, most of the ALV in China that has been known to infect chickens belongs to subgroups A, B and J [7, 27–29]. In recent years, a novel ALV subgroup named ALV-K has been isolated from indigenous Chinese chicken breeds, based on gp85 amino acid sequence comparisons [2–4]. Notably, more and more ALV-K strains have been isolated from local Chinese chicken breeds and these isolates represent a new subgroup of ALV viruses that do not induce tumors in SPF chickens and replicate at a relatively slow rate in DF-1 cells [4]. Because the Nationwide Eradication Program (NEP) for ALV in chicken breeder farms was not initiated in China until 2008, ALV infection in chickens has caused serious problems. Over the past decade, many tumor cases induced by ALV have been reported [23, 27]. No effective drugs or
Vaccines are available for ALV, which is a major challenge for the control and eradication of this virus. The best way to control ALV infection is for all of the exogenous ALV subgroups to be eradicated from chicken breeder flocks. Currently, for large scale identification of these pathogens, a preliminary test is done to see whether ALVs exist and then, if necessary, primers specific for subgroups A, B and J are used to detect the specific ALV subgroup, however this system is not available for ALV-K. At present, ALV-K diagnostic methods mainly rely on virus isolation followed by envelope gene sequencing comparisons, and there is lack of other diagnostic methods.

Retroviruses only efficiently infect cells that express a specific receptor that can interact with their viral envelope glycoproteins [14]. Shortly after the virus has gained entry, it begins to produce viral proteins. As the envelope protein is produced, it saturates the cell surface receptors and blocks superinfection of viruses from the same subgroup [30]. Because saturation of the viral cell receptors of susceptible cells via the expression of viral receptor binding protein can block corresponding viral infection, genetically engineered cell lines resistant to ALV-J infection have been developed [22, 31, 32]. The host range of subgroups A through E is more restricted, and there are

![Fig. 6 The antiviral experiment results. Five different virus titers from $10^1$ TCID$_{50}$ to $10^5$ TCID$_{50}$ per 0.1 ml of ALV were inoculated per well in 24-well cell culture plates containing 1mL 1.7x10^5 cells/well DF-1/K cells, all in triplicate. A. ALV-K (GDFX0601) replication was inhibited in DF-1/K, but not DF-1 cells. B, C, D. Replication of ALV-A (GD13), ALV-B (CD08), ALV-J (CHN06) were not inhibited in either DF-1/K and DF-1 cells. E. Infection by four ALV-K field isolates was blocked in DF-1/K, but not DF-1 cells. In A-E, viral p27 protein levels determined by ELISA are reported. Black lines mean S/P = 0.2](image-url)
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lines of chickens that possess genetic resistance to one or more of these subgroups [33–35].

The host range of ALV-K and which chicken lines it is able to infect has not yet been characterized. The lack of genetically resistant cells makes differentiating ALV-K from other subgroups difficult. In addition the lack of genetically resistant chicken lines complicates the classification of field samples. In an attempt to overcome the lack of cell lines that are genetically resistant to ALV-K, we have developed a specific ALV-K-resistant cell line that expresses the subgroup K env protein. The env gene of one ALV-K isolate, GDFX0601, was cloned and expressed in the DF-1 cell line to create the ALV-K-resistant cell line (DF-1/K). This over-expressed env protein should theoretically bind to the cells' viral receptors and selectively interfere with ALV-K infection. The ALV-K-resistant cell line and its parental cell line DF-1 do not express other ALV proteins and thus the ALV p27 antigen ELISA can be used to monitor ALV infection. To test this system, the env gene in DF-1/K cells was detected by PCR, and the env protein itself was detected by IFA and Western blot. The results showed that DF-1/K cells could stably express the ALV-K env gene. Immune electron microscopy revealed that the ALV-K env fusion protein is localized in the cell membrane area of DF-1/K cells. The anti-virus assay showed that DF-1/K cells showed resistance to ALV-K infection at a viral dose of 1 × 10^4 TCID_{50}. ALV-K in field samples can be definitively identified by using ELISA to monitor levels of p27 in DF-1 and DF-1/K cells that have been inoculated with field samples. Only inoculation with ALV-K will result in the detection of p27 in DF-1 cells and no or a lower p27 response in DF-1/K cells. Tests for p27 antibodies, goat anti-mouse EGFP antibodies and single factor antiserum for ALV-K individually. These were then subsequently incubated with 10nm colloidal gold-labeled goat anti-mouse IgG. TEM ×250000. The arrows point to gold particles.

In conclusion, we have developed a specific DF-1/K cell line that expresses a viral receptor-binding protein and is resistant only to ALV-K infection. We have not only constructed a cell line that will be a useful diagnostic tool for the novel ALV-K subgroup. Others have demonstrated that cell lines that express viral receptor-binding protein can be efficient tools for isolating functional receptors; a cell line resistant to ALV-J infection was developed and applied to efficiently identify chicken Annexin A2 (chANXA2) as a novel receptor for retrovirus ALV-J [15]. Our DF-1/K cell line will be further applied to identify ALV-K functional receptors and to identify novel anti-viral targets. This study also is helpful to existing efforts to control and eradicate exogenous ALV in local Chinese chickens.

Compliance with ethical standards

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Conflict of interest  The authors declare that they have no competing interests. RMZ, CWS, LM and FM conceived and designed the experiments; RMZ and ZZJ performed the experiments. YLX, CJ and ZJ contributed reagents/materials/analysis tools. RMZ contributed to the writing of the manuscript. All authors read and approved the final manuscript.
Ethical approval  Our animal research was conducted under the guidance of the SCAU’s Institutional Animal Care and Use Committee. The chicken sampling procedures were approved by the Animal Care and Use Committee of Guangdong Province, China.

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References

1. Payne LN, Howes K, Gillespie AM, Smith LM (1992) Host range of Rous sarcoma virus pseudotype RSV (HPRS-103) in 12 avian species: support for a new avian retrovirus envelope subgroup, designated. J Gen Virol 73(Pt 11):2995–2997
2. Wang X, Zhao P, Cui ZZ (2012) Identification of a new subgroup of avian leukemia virus isolated from Chinese indigenous chicken breeds. Bing Du Xue Bao 28:609–614 (in Chinese, with English abstract)
3. Cui N, Su S, Chen Z, Zhao X, Cui Z (2014) Genomic sequence analysis and biological characteristics of a rescued clone of avian leukemia virus strain JS11C1, isolated from indigenous chickens. J Gen Virol 95:2512–2522
4. Li X et al (2016) Isolation, identification and evolution analysis of a novel subgroup of avian leukemia virus isolated from a local Chinese yellow broiler in South China. Arch Virol 161(10):2717–2725
5. Zavala G, Cheng S (2006) Detection and characterization of avian leukemia virus in Marek’s disease vaccines. Avian Dis 50:209–215
6. Silva RF, Fadly AM, Taylor SP (2007) Development of a polymerase chain reaction to differentiate avian leukemia virus (ALV) subgroups: detection of an ALV contaminant in commercial Marek’s disease vaccines. Avian Dis 51:663–667
7. Cui Z, Du Y, Zhang Z, Silva RF (2003) Comparison of Chinese field strains of avian leukemia subgroup J viruses with prototype strain HPRS-103 and United States strains. Avian Dis 47:1321–1330
8. Tomioka Y, Ochiai K, Ohashi K, Kimura T, Umemura T (2003) In vivo infection with an avian leukemia virus causing fowl glioma: viral distribution and pathogenesis. Avian Pathol 32:617–624
9. Toyoda T, Ochiai K, Ohashi K, Tomioka Y, Kimura T, Umemura T (2005) Multiple perineuromas in chicken (Gallus gallus domesticus). Vet Pathol 42:176–183
10. Tomioka Y (2004) Genome sequence analysis of the SCAU’s Institutional Animal Care and Use Committee. The animal research was conducted under the guidance of the Creative Commons license, and indicate if changes were made.

Ethical approval  Our animal research was conducted under the guidance of the SCAU’s Institutional Animal Care and Use Committee. The chicken sampling procedures were approved by the Animal Care and Use Committee of Guangdong Province, China.

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References

1. Payne LN, Howes K, Gillespie AM, Smith LM (1992) Host range of Rous sarcoma virus pseudotype RSV (HPRS-103) in 12 avian species: support for a new avian retrovirus envelope subgroup, designated. J Gen Virol 73(Pt 11):2995–2997
2. Wang X, Zhao P, Cui ZZ (2012) Identification of a new subgroup of avian leukemia virus isolated from Chinese indigenous chicken breeds. Bing Du Xue Bao 28:609–614 (in Chinese, with English abstract)
3. Cui N, Su S, Chen Z, Zhao X, Cui Z (2014) Genomic sequence analysis and biological characteristics of a rescued clone of avian leukemia virus strain JS11C1, isolated from indigenous chickens. J Gen Virol 95:2512–2522
4. Li X et al (2016) Isolation, identification and evolution analysis of a novel subgroup of avian leukemia virus isolated from a local Chinese yellow broiler in South China. Arch Virol 161(10):2717–2725
5. Zavala G, Cheng S (2006) Detection and characterization of avian leukemia virus in Marek’s disease vaccines. Avian Dis 50:209–215
6. Silva RF, Fadly AM, Taylor SP (2007) Development of a polymerase chain reaction to differentiate avian leukemia virus (ALV) subgroups: detection of an ALV contaminant in commercial Marek’s disease vaccines. Avian Dis 51:663–667
7. Cui Z, Du Y, Zhang Z, Silva RF (2003) Comparison of Chinese field strains of avian leukemia subgroup J viruses with prototype strain HPRS-103 and United States strains. Avian Dis 47:1321–1330
8. Tomioka Y, Ochiai K, Ohashi K, Kimura T, Umemura T (2003) In vivo infection with an avian leukemia virus causing fowl glioma: viral distribution and pathogenesis. Avian Pathol 32:617–624
9. Toyoda T, Ochiai K, Ohashi K, Tomioka Y, Kimura T, Umemura T (2005) Multiple perineuromas in chicken (Gallus gallus domesticus). Vet Pathol 42:176–183
10. Tomioka Y (2004) Genome sequence analysis of the SCAU’s Institutional Animal Care and Use Committee. The animal research was conducted under the guidance of the Creative Commons license, and indicate if changes were made.

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References

1. Payne LN, Howes K, Gillespie AM, Smith LM (1992) Host range of Rous sarcoma virus pseudotype RSV (HPRS-103) in 12 avian species: support for a new avian retrovirus envelope subgroup, designated. J Gen Virol 73(Pt 11):2995–2997
2. Wang X, Zhao P, Cui ZZ (2012) Identification of a new subgroup of avian leukemia virus isolated from Chinese indigenous chicken breeds. Bing Du Xue Bao 28:609–614 (in Chinese, with English abstract)
3. Cui N, Su S, Chen Z, Zhao X, Cui Z (2014) Genomic sequence analysis and biological characteristics of a rescued clone of avian leukemia virus strain JS11C1, isolated from indigenous chickens. J Gen Virol 95:2512–2522
4. Li X et al (2016) Isolation, identification and evolution analysis of a novel subgroup of avian leukemia virus isolated from a local Chinese yellow broiler in South China. Arch Virol 161(10):2717–2725
5. Zavala G, Cheng S (2006) Detection and characterization of avian leukemia virus in Marek’s disease vaccines. Avian Dis 50:209–215
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7. Cui Z, Du Y, Zhang Z, Silva RF (2003) Comparison of Chinese field strains of avian leukemia subgroup J viruses with prototype strain HPRS-103 and United States strains. Avian Dis 47:1321–1330
8. Tomioka Y, Ochiai K, Ohashi K, Kimura T, Umemura T (2003) In vivo infection with an avian leukemia virus causing fowl glioma: viral distribution and pathogenesis. Avian Pathol 32:617–624
9. Toyoda T, Ochiai K, Ohashi K, Tomioka Y, Kimura T, Umemura T (2005) Multiple perineuromas in chicken (Gallus gallus domesticus). Vet Pathol 42:176–183
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