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Arginine regulates inflammation response-induced by *Fowl Adenovirus* serotype 4 via JAK2/STAT3 pathway

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Abstract

Background: *Fowl Adenovirus* serotype 4 (FAdV-4) infection causes severe inflammatory response leading to hepatitis-hydropericardium syndrome (HHS) in poultry. As an essential functional amino acid of poultry, arginine plays a critical role in anti-inflammatory and anti-oxidative stress.

Results: In this study, the differential expression genes (DEGs) were screened by transcriptomic techniques, and the DEGs in gene networks of inflammatory response-induced by FAdV-4 in broiler's liver were analyzed by Kyoto encyclopedia of genes and genomes (KEGG) enrichment. The results showed that the cytokines pathway and JAK/STAT pathway were significantly enriched, in which the DEGs levels of IL-6, IL-1 β , IFN- α , JAK and STAT were significantly up-regulated after FAdV-4 infection. It was further verified with real-time fluorescence quantitative polymerase chain reaction (Real-time qPCR) and Western blotting (WB) in vitro and in vivo. The findings demonstrated that FAdV-4 induced inflammatory response and activated JAK2/STAT3 pathway. Furthermore, we investigated whether arginine could alleviate the liver inflammation induced by FAdV-4. After treatment with 1.92% arginine level diet to broilers or 300 μ g/mL arginine culture medium to LMH cell line with FAdV-4 infection at the same time, we found that the mRNA levels of IL-6, IL-1 β , IFN- α and the protein levels of p-JAK2, p-STAT3 were down-regulated, compared with FAdV-4 infection group. Furthermore, we confirmed that the inflammation induced by FAdV-4 was ameliorated by pre-treatment with JAK inhibitor AG490 in LMH cells, and it was further alleviated in LMH cells treatment with AG490 and ARG.

Conclusions: These above results provide new insight that arginine protects hepatocytes against inflammation induced by FAdV-4 through JAK2/STAT3 signaling pathway.

Keywords: Arginine, *Fowl adenovirus* serotype 4, Inflammation, JAK2/STAT3 pathway, Transcriptome, Broiler

Introduction

In 1987 *Fowl Adenovirus* serotype 4 (FAdV-4) was firstly reported in Pakistan [1]. Since 2015, FAdV-4 infection had large-scale epidemics in China, which caused incalculable economic losses to the poultry industry [2]. The FAdV-4 infection causes to severe

hepatitis-hydropericardium syndrome (HHS) in poultry [3, 4], which pathological features include pericardial effusion, swollen and yellowish livers with bleeding spot and bleeding or enlargement of immune organs [2, 5]. Previous studies demonstrated that the occurrence of the pathological inflammatory response caused by FAdV-4 induced excessive secretion of pro-inflammatory cytokines, resulting in loss of appetite, energy deficiency, and other physiological behaviors and metabolic abnormalities in animals [6]. Moreover, the over-expression of inflammatory factors such as interleukin-1 β (IL-1 β), interleukin-6 (IL-6) and tumor necrosis factor-alpha

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(TNF- α) induced by FAdV-4 caused hepatitis and further necrosis [7].

Janus kinase (JAK)/signal transducer and activator of transcription (STAT) signaling pathway widely participates in physiological processes such as inflammation, apoptosis, immunity, and tumor formation [8, 9]. In particular, the JAK2/STAT3 signaling pathway that is activated by inflammatory factors such as IL-6, is most closely related to inflammation [10, 11]. At the initial stage, the activation of JAK2 by inflammatory factors would phosphorylate its downstream STAT3 protein, which ultimately causes the over-expression of cytokines genes and induces cellular pathophysiology [12].

Arginine (ARG) is a conditionally essential amino acid, which participates in vital biological processes, including regulation of the normal function of the immune systems in an organism [13]. Arginine as a dietary nutrient supplement could improve immunologic function, regulate cellular growth, proliferation and immune responses [14]. Our previous studies confirmed that arginine could inhibit FAdV-4 replication in LMH cells [15]. In addition, studies verified that arginine could regulate inflammation of animal. It was indicated that the supplementation of arginine via the jugular vein during early lactation alleviated inflammation of dairy cows [16]. Furthermore, arginine could alleviate the inflammation of excessive immune response [17]. In porcine intestinal epithelial cells (IPEC-J2), L-arginine obviously suppressed the levels of IL-6, IL-8, IL-1 β , and TNF- α induced by LPS, alleviated inflammatory response and maintained intestinal integrity [18]. However, it is unknown whether arginine can alleviate the inflammation response induced by FAdV-4.

In this study, we performed transcriptomic technology to screen for DEGs in the liver (the main target organ of FAdV-4) of broilers infected with FAdV-4, and used KEGG to analyze the inflammatory pathway. Meanwhile, qPCR and Western blotting were used to validate inflammation-related factors and pathways in vivo and vitro. Furthermore, the regulation of arginine on cytokines and the JAK2/STAT3 pathway in the inflammatory state was determined.

Materials and methods

Virus

FAdV-4 strain NP (TCID₅₀ = 10^{-6.23}/0.01 mL) was proliferated in DF-1 cell line with RPMI-1640 medium (12,633,012, Gibco, Invitrogen Corp., California, U.S.A), 10% fetal bovine serum (FBS) (Serana, Washington, U.S.A), and 1% penicillin/streptomycin (10,378,016, Gibco, Invitrogen Corp., California, U.S.A) in a 37 °C incubator (Panasonic, Osaka, Japan) with 5% carbon dioxide (CO₂).

Cell culture and treatment

Chicken hepatocellular carcinoma cell line (LMH), a classical hepatocyte model for studying chicken pathogenicity or gene function, was used as a cell model in this experiment [19, 20]. LMH cell line (lot no. 601411-714SE, CLS, Germany) was subculture in modified RPMI-1640 medium (12,633,012, Gibco, Invitrogen Corp., California, U.S.A) with 10% fetal bovine serum (FBS) (Serana, Washington, U.S.A) and 1% penicillin/streptomycin (10,378,016, Gibco, Invitrogen Corp., California, U.S.A) in a 37 °C incubator (Panasonic, Osaka, Japan) with 5% CO₂. LMHs were treated with arginine at concentrations of 300 μ g/mL, while the cells in the control group were cultured in a normal medium for 24 h. The final culture system was 5 mL. Then, cells were collected after infection with 0.5 mL FAdV-4 ([TCID₅₀ = 10^{-6.23}/0.01 mL) for 2 h. Furthermore, AG490 (a JAK2 inhibitor) was dissolved as 50 nmol/L using dimethyl sulfoxide (DMSO) (D2650, Sigma, Burlington, U.S.A) to treat LMH cells for 2 h at 37 °C [21].

Animals and treatments

Eighteen 21-day-old normal white-feathered male broilers (Ross 308) were purchased from Fujian Shengnong Food Co. Ltd., China. All birds with negative FAdV-4 antigen and antibody were randomly divided into two groups. Chickens in the infection group were intramuscularly injected with 0.5 mL FAdV4 (TCID₅₀ = 10^{-6.23}/0.1 mL). Conversely, broilers in the mock group were injected with the same volume of sterile saline. Two days after FAdV-4 infection, liver samples were collected for transcriptome sequencing.

Additionally, 180 twenty-one-day-old broilers (Fujian Shengnong Food Co. Ltd., China) were randomly divided into 3 treatments with 5 replicates per group and 12 broilers per replicate. The FAdV-4 and ARG + FAdV-4 groups were treated by randomly selecting 6 sample broilers of similar weight from the control feeding group and the 1.92% level of arginine feeding group for intramuscular injection of 0.5 mL FAdV-4, respectively. In contrast, 6 broilers from the control feeding group were injected with sterile saline as the mock group. After 2 d, the chickens were deeply anaesthetized with CO₂ and euthanized for liver collection.

All broilers were fed with corn-soybean meal diet. The diets were formulated to meet the nutritional requirements of broilers as defined by the National Research Council of China (NRC) [22, 23]. The composition and nutritional components of broiler diets are shown in Table 1. Our previous results showed that the 1.92% arginine nutrient level had the most significant effect on resistance to FAdV-4 infection in broilers.

Table 1 Composition and nutrient levels of white-feathered broiler diets (air-dry basis)

	Arginine levels in diet (%)	
	1.20%	1.92%
Ingredients		
Corn	62.78	64.67
Corn gluten meal	16.62	14.42
Soybean meal	9.36	10.47
Puffed soybean	3.00	3.00
Limestone	1.25	1.25
CaHPO ₄	2.20	2.20
NaCl	0.30	0.30
L-Lys	0.65	0.65
DL-Met	0.19	0.19
L-Thr	0.09	0.09
L-Trp	0.04	0.04
Zeolite powder	2.28	0.76
Premix ^a	1.00	1.00
ARG	0.24	0.96
Total	100	100
Nutrient levels		
ME(KJ/kg)	12.54	12.54
CP	21.50	21.50
Ca	1.00	1.00
P	0.45	0.45
Lys	1.15	1.15
Met + Cys	0.91	0.91
Thr	0.81	0.81
Trp	0.21	0.21
ARG	1.20	1.92

^a The premix provides following for per kg diet: vitamin A, 9,800 U; vitamin D₃, 2,000 U; vitamin E, 21 mg; vitamin K₃, 2.24 mg; vitamin B₁, 2.4 mg; vitamin B₂, 6.1 mg; vitamin B₃, 36.5 mg; vitamin B₅, 10.1 mg; vitamin B₆, 9.8 mg; vitamin B₁₂, 0.02 mg; biotin, 0.2 mg; folic acid, 1.12 mg; choline, 1,300 mg; Cu, 8 mg; Fe, 100 mg; Mn, 120 mg; Zn, 100 mg; Se, 0.3 mg; I, 0.7 mg

Thus, the ARG treatment was defined as 1.92% of the total nutrient level [24].

KEGG enrichment analysis

Total RNA was extracted from the livers of broilers in FAdV-4 infection group and control group. The

appropriate fragments were selected for PCR amplification by agarose gel electrophoresis. Sequencing libraries were constructed and sequenced using Illumina[®] HiSeq TM 2000 (Illumina, Shenzhen, China). And the quality and saturation of sequenced reads were evaluated. Then, low-quality reads and impurities from the raw reads were removed to obtain clean reads. The RPKM (reads per kilobase per million mapped reads) method was used to screen and analyze DEGs. The *P*-value was controlled by the false discovery rate, while the false discovery rate for DEGs was ≤ 0.001 , and its difference ratio was less than twice [2].

The KEGG functional enrichment analysis was used to screen significant pathways. A pathway with $Q \leq 0.05$ was characterized as enriched. Significant pathway enrichment enabled the identification of biochemical metabolic pathways and signaling pathways associated with DEGs [2].

Polymerase chain reaction (PCR) detection

The specific primers for FAdV-4 Fiber 1 (Gene ID: 10,399,487) were designed (Table 2) via Primer 5 (Molecular Biology Insights, Cascade, Co). The virus nucleic acid was extracted with the virus DNA extraction kit ((TransGen Biotech, Beijing, China) and amplified by PCR [2]. The PCR reaction system (25 μ L) comprised 1 μ L of DNA template, 12.5 μ L of PCR Nucleotide Mix (10 mmol/L), 9.5 μ L of nuclease-free water, and 1 μ L of each upstream and downstream primers. The reaction conditions were as follows: denaturation at 95 °C for 30 s, annealing at 55 °C for 30 s, and extension at 72 °C for 30 s, 30 cycles. PCR products were analyzed by 1% agarose gel electrophoresis.

The mRNA levels were determined by qPCR

Total RNA was extracted from chicken livers and LMH cells using TransZol Up RNA Kit (ER501-01, Beijing TranGen Biotech Co. Ltd., China). The concentration and purity of RNA were identified by measuring OD₂₆₀/OD₂₈₀ values with the NanoDrop 2000 spectrophotometer (Thermo Scientific, Waltham, MA), then they were frozen at -80 °C. The reverse transcription system was as follows: 1 μ g of total RNA, 2 μ L RT Master Mix (5 \times), and RNase-free water to fill a total 10 μ L. The products

Table 2 Primer sequences for PCR

Genes	Sequence (5'to3')	Product size, bp	GenBank No
FAdV-4 Fiber 1	F: CCCTCGAGATGTCGGCCCTAATCGCCTCC R: CGGAATTCGGGGCCCGAGCATTGT	1315	APA_19531
β -actin	F: CCCACACCCTGTGATGAAA R: TAGAACTTTGGGGCGTTTCG	148	NM_205518

(cDNA) were stored at -20°C for subsequent real-time qPCR [25].

We found the gene sequences of chicken β -actin, IL-6, IL-1 β , and IFN- α in GenBank with accession numbers of NM_205518, NM_204628, NM_204524, and NM_205427, respectively. Specific primers (Table 3) for qPCR were designed using Primer 15 software. Real-time qPCR was performed to detect the expression levels of IL-6, IL-1 β , and IFN- α genes in different cDNA samples using TransStart[®] Green qPCR SuperMix (AQ101-01, Beijing TranGen Biotech Co. Ltd., China). Refer to Wang [26], the reaction mixtures were incubated in a LightCycler 480 II real-time PCR system (Basel, Switzerland). The relative mRNA expression levels were calculated using the $2^{-\Delta\Delta\text{Ct}}$ method, and the data were calibrated as the relative value to the control group. All samples were analyzed in triplicate.

Western blotting examines protein levels

Radio immunoprecipitation assay (RIPA) buffer (P0013B, Beyotime, Shanghai, China) encompassing 1% protease (ST506, Beyotime, Shanghai, China) and 1% phosphatase inhibitors (P1045, Beyotime, Shanghai, China) were added to obtain the total protein supernatant. Proteins were detached using 10% sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) (P0012A, Beyotime, Shanghai, China) and then were transferred to polyvinylidene fluoride (PVDF) membranes (BSP0161, Pall, New York, U.S.A). After blocking the membranes with 5% skimmed milk (P0216, Beyotime, Shanghai, China), the membranes were transferred to anti-JAK2 (1:1,000; 3230S, Cell Signaling Technology, Boston, U.S.A), anti-STAT3 (1:1,000; 12640S, Cell Signaling Technology, Boston, U.S.A), anti-phospho-JAK2 (1:1,000; 3771S, Cell Signaling Technology, Boston, U.S.A), anti-phospho-STAT3 (1:1,000; 9145S, Cell Signaling Technology, Boston, U.S.A) and anti- β -actin (1:2000; 4970S, Cell Signaling Technology, Boston, U.S.A) at 4°C overnight [26, 27]. Membranes were washed 3 times with $1 \times$ tris

buffered saline tween (TBST) (T1081, Solarbio, Beijing, China), then were incubated with secondary antibody (1:3000; 7074S, Cell Signaling Technology, Boston, U.S.A) that was conjugated with horseradish peroxidase (HRP). Immunoblots were developed using enhanced chemiluminescence (ECL) reaction (KF001, Affinity Biosciences, OH, U.S.A) and were imaged on a ChemiDoc XRS System (BioRad, California, U.S.A). The value of the target proteins were quantitatively calculated using Image J software.

Statistical analysis

The data were presented as the mean ($n=6$) \pm standard difference (SD) unless noted otherwise. Data were analyzed by independent-samples ANOVA using the SPSS statistical software (Ver.16.0 for windows, SPSS Inc., Chicago, U.S.A). $P < 0.05$ was considered statistically significant. Data handling and statistical processing were performed using GraphPad Prism 7.0 (GraphPad Software, San Diego, CA, U.S.A).

Results

FAdV-4 promoted inflammatory cytokines and JAK2/STAT3 signaling pathway in LMH cells

Results showed that FAdV-4 could significantly induce the mRNA expression levels of IL-6, IL-1 β and IFN- α in LMH cells (Fig. 1 A-D) ($P < 0.05$). Furthermore, the imprinting degree of p-JAK2 and p-STAT3 protein bands in LMH cells infected with FAdV-4 was higher than that in the control group (Fig. 1 E–F). Meanwhile, by analyzing the gray value of the protein bands and transforming them into histograms, we could intuitively observe that the phosphorylation of JAK2 and STAT3 proteins was aggravated by FAdV-4 infection (Fig. 1 G), which indicated that FAdV-4 infection promoted the activation of JAK2 and STAT3.

Arginine relieved inflammatory factors and JAK2/STAT3 signaling pathway in LMH cells-infected by FAdV-4

In the present study, the mRNA levels of IL-6, IL-1 β and IFN- α in LMH cells -infected with FAdV-4 co-treatment with 300 $\mu\text{g}/\text{mL}$ arginine medium were significantly relieved (Fig. 2 A-D) ($P < 0.05$), compared with FAdV-4 infection cells. Meanwhile, cells-treated with 300 $\mu\text{g}/\text{mL}$ arginine medium reduced the phosphorylation and gray values of p-JAK2 and p-STAT3 after infection with FAdV-4, compared with FAdV-4 infection cells (Fig. 2 E–G) ($P < 0.05$). These results demonstrated that arginine treatment also could alleviate the inflammatory response of LMH cells infected with FAdV-4.

Table 3 Primer sequences for real-time qPCR

Genes	Sequence (5'to3')	Product size, bp	GenBank No
IL-6	F: AAATCCCTCCTCGCCAATCT	133	NM_204628
	R: CCCTCACGGTCTTCTCCATAAA		
IL-1 β	F: CTGCTGCAGAAGAAGCCT	164	NM_204524
	R: TGTCAGCAAAGTCCCTGCTC		
IFN- α	F: CAACGACACCATCTGGACA	147	NM_205427
	R: ATCCGGTTGAGGAGGCTTTG		
β -actin	F: CTGGCACCTAGCACAAATGAA	90	NM_205518
	R: CTGCTTGCTGATCCACATCT		

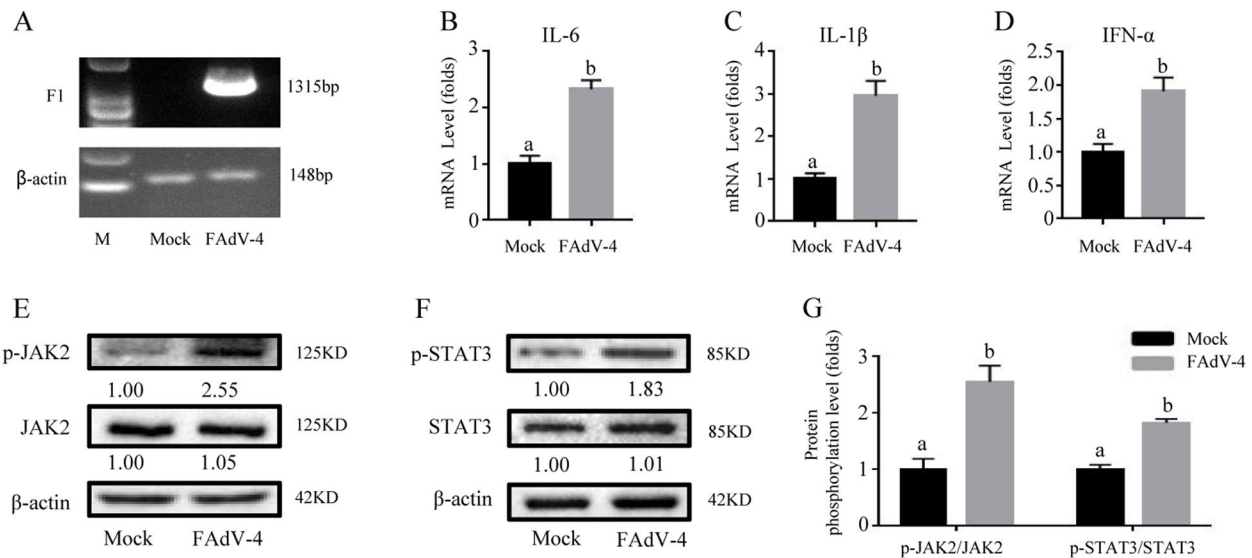


Fig. 1 FAdV-4 promoted inflammatory cytokines and the JAK2/STAT3 pathway in LMH cells. The inflammatory indexes of cells infected with or without FAdV-4 were evaluated. After infection with FAdV-4 24 h (A), the virus load of FAdV-4 was determined by PCR with the primers of F1 gene, and the β-actin was used as a conference gene. The mRNA levels of cytokines (B) *IL-6*, (C) *IL-1β*, and (D) *IFN-α* was determined by qPCR, while the phosphorylation of crucial proteins (p-JAK2 and p-STAT3) in the JAK2/STAT3 pathway was analyzed by WB (E-F). (G) The Western blotting was analyzed by gray scale and described by histogram, histogram is the result of the ratio of the phosphorylated protein to the total protein, and is the digital embodiment of the protein phosphorylation level of JAK2 and STAT3. Significance between the treatments was determined by T-test analysis using SPSS software (Version 20.0). Means with different alphabets (a, b) denotes significance at $p < 0.05$. β-actin was used as a control for protein loading, and the following were consistent. All values are expressed as Means ± SD (n = 6). The a, b, c, d bars in each panel without a common superscript letter were significantly different ($P < 0.05$). All remain consistent below unless otherwise stated

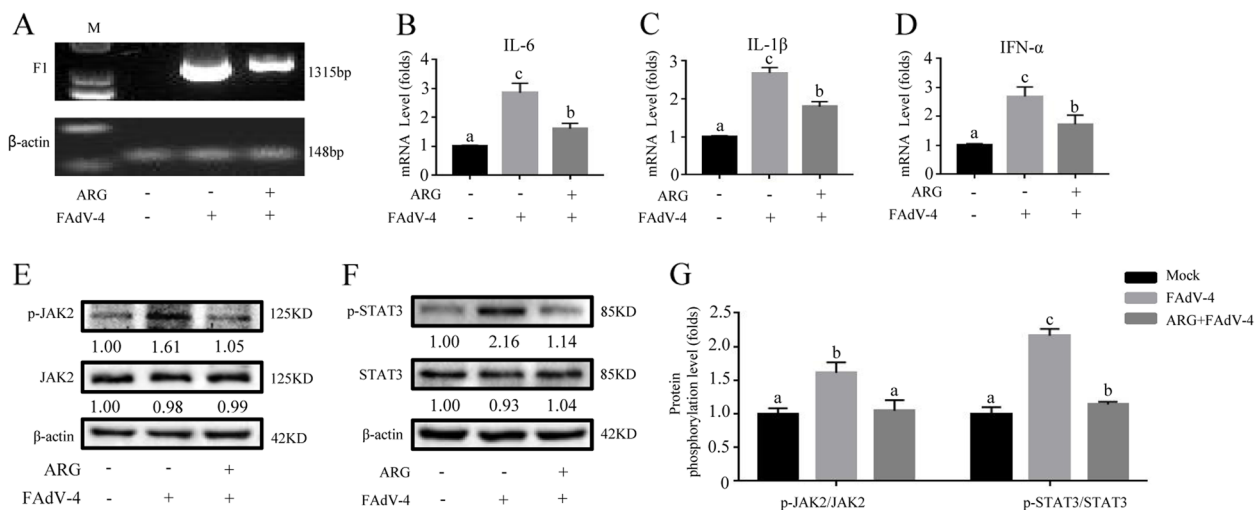
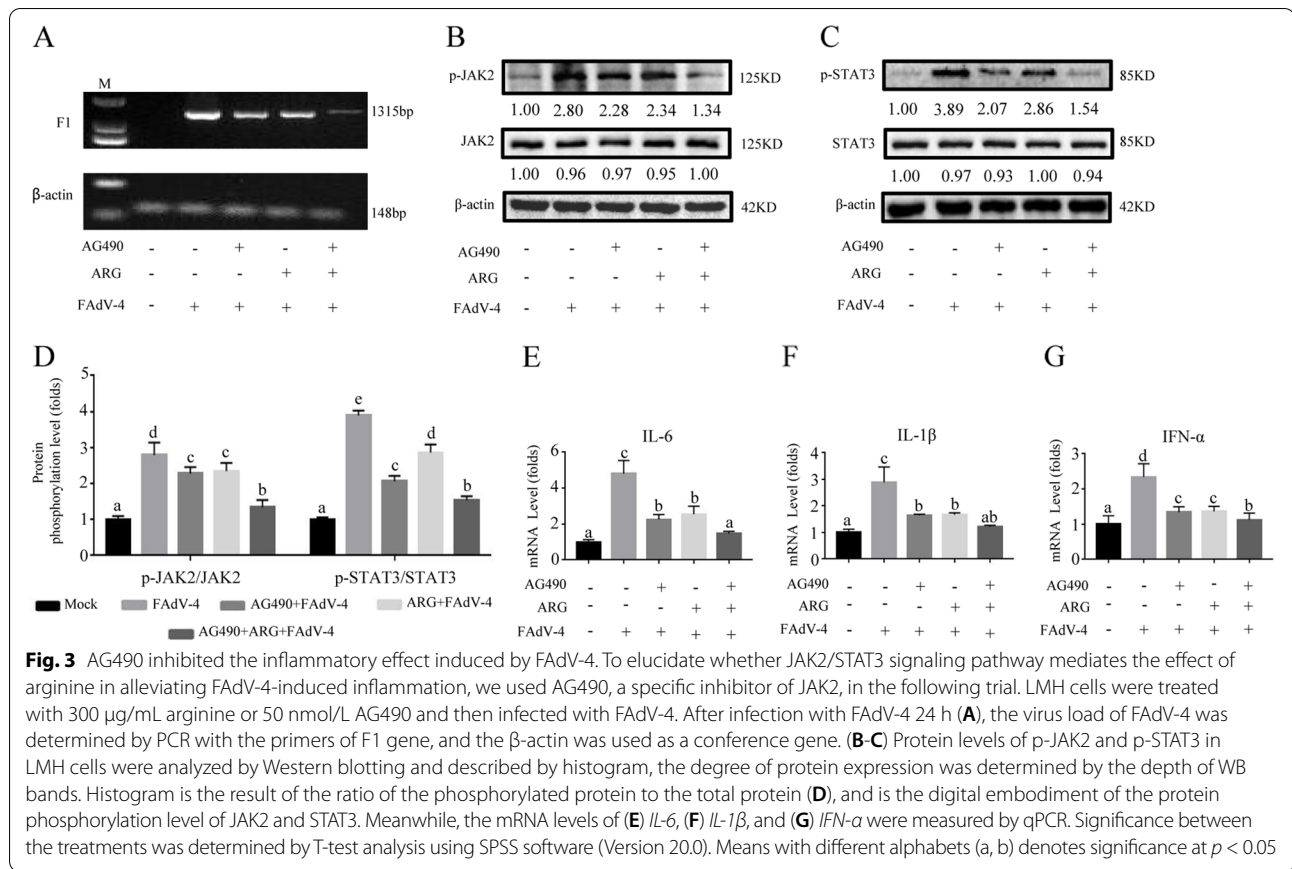


Fig. 2 Arginine treatment down-regulated inflammatory response and JAK2/STAT3 pathway-activated by FAdV-4 in LMH cells. The cells were divided into three groups: mock and FAdV-4 groups were cultured with a basal medium containing 10% FBS; ARG + FAdV-4 group was cultured with 300 μg/mL arginine medium. In addition, the FAdV-4 and the ARG + FAdV-4 groups were treated with the FAdV-4 strain NP for 2 h. On the contrary, the mock group was still cultured with a basic medium. After infection with FAdV-4 24 h (A), the virus load of FAdV-4 was determined by PCR with the primers of F1 gene, and the β-actin was used as a conference gene. The hepatic mRNA levels of (B) *IL-6*, (C) *IL-1β*, and (D) *IFN-α* were assessed. Similarly, the effect of arginine on the JAK2/STAT3 signaling pathway was determined by evaluating whether arginine interferes with the phosphorylated expression of JAK2 and STAT3 (E-F). (G) The degree of protein expression was determined by the depth of WB bands. Histogram is the result of the ratio of the phosphorylated protein to the total protein, and is the digital embodiment of the protein phosphorylation level of JAK2 and STAT3. Significance between the treatments was determined by T-test analysis using SPSS software (Version 20.0). Means with different alphabets (a, b) denotes significance at $p < 0.05$



JAK2 inhibitor AG490 further alleviates inflammation in FAdV-4-infected LMH cells co-treatment with arginine

To further investigate whether the JAK2/STAT3 pathway mediates the protective effects of ARG, a widely accepted JAK2 inhibitor AG490 was employed. ARG treatment or AG490 treatment both significantly inhibited the phosphorylation level of p-JAK2 and its downstream p-STAT3 induced by FAdV-4 (Fig. 3 A-D) ($P < 0.05$). And after co-treatment with ARG and AG490, the phosphorylation level of p-JAK2 and p-STAT3 was significantly down-regulated in LMH cells infected with FAdV-4, compared with only ARG treatment or AG490.

In addition, the result showed that the inflammatory cytokines IL-6, IL-1β, and IFN-α that were induced by FAdV-4 were down-regulated in the LMH cell line after treatment with AG490, compared with cells that was infected with FAdV-4 ($P < 0.05$). After ARG and AG490 co-treatment, the mRNA levels of IL-6, IL-1β and IFN-α were significantly decreased in LMH cells infected with FAdV-4 ($P < 0.05$), compared with only ARG or AG490 treatment cells (Fig. 3 E-G).

KEGG functional analysis on the livers infected with FAdV-4

In this study, we concentrated on the pathway about the inflammatory response by the KEGG functional analysis

(Ref: 220,192). Table 4 showed that the cytokines pathway and the JAK/STAT pathway were enriched in the liver at post-infection with FAdV-4, compared within mock livers. Furthermore, DEGs including IL-6, IL-1β, IFN-α, JAK and STAT were up-regulated. These results revealed that FAdV-4 infection caused the inflammatory response in livers of broilers, which was regulated by JAK/STAT pathway.

FAdV-4 up-regulated the expression of inflammatory cytokines, p-JAK2 and p-STAT3 in broiler liver

Furthermore, we determined the mRNA levels of IL-6, IL-1β, and IFN-α in the liver of FAdV-4-challenged chickens by qPCR. After FAdV-4 infection, the mRNA levels of IL-6, IL-1β, and IFN-α were significantly increased ($P < 0.05$) (Fig. 4 A-D). Concurrently, as shown in Fig. 4 E-F, the phosphorylation blotting of p-JAK2 and p-STAT3 were significantly improved in livers of FAdV-4 infection broilers ($P < 0.05$). Histogram results of gray

Table 4 KEGG analysis on the broiler livers infected with FAdV-4

Pathway	DEGs	Express trend
Cytokines	IL-6、IL-1β、IFN-α	up-regulated
JAK-STAT	JAK、STAT	up-regulated

values further confirmed the promotion of FAdV-4 on JAK2 and STAT3 phosphorylation in broilers (Fig. 4 G). These results further supported the KEGG data in vivo.

Arginine down-regulated the expression of inflammatory cytokines and inhibits JAK2/STAT3 signaling pathway in broiler liver-infected with FAdV-4

In our present study, 1.92% level of dietary arginine treatment could significantly down-regulate the expression of pro-inflammatory factors IL-6, IL-1β, and IFN-α in broiler liver-infected with FAdV-4 ($P < 0.05$) (Fig. 5 A-D). Moreover, the protein levels of p-JAK2 and p-STAT3 were significantly reduced in 1.92% level of dietary arginine treatment group-infected with FAdV-4 ($P < 0.05$) (Fig. 5 E-G). These results demonstrated that arginine could down-regulate inflammation response and JAK2/STAT3 pathway in the broiler liver-infected with FAdV-4.

Therefore, these results revealed that arginine alleviated inflammation- induced by FAdV-4 through regulating the JAK2/STAT3 signaling pathway.

Discussion

The FAdV-4 infection caused severe economic lose to the global poultry industry [28]. At present, the pathogenic mechanism of FAdV-4 infection still has not been

fully understood. It was well known that IBH and HHS were typical fatal infectious diseases caused by FAdV-4 [29]. Commonly, the pathological feature of FAdV-4-induced HHS disease were severe hydropericardium and liver inflammation [30]. The major findings were gross lesions in the hydropericardium with the accumulation of yellow gelatinous fluid in the pericardium [31]. At histopathological analysis, the most consistent findings were multifocal areas of necrosis and mononuclear cell infiltration in the liver, including basophilic intranuclear inclusion bodies in hepatocytes [32]. Poultry-infected with FAdV-4 showed obvious multifocal areas of necrosis in the liver and abundant inflammatory cells in liver tissue [33]. Previous studies have shown that the JAK/STAT signaling pathway was a major mechanism of broad cytokine signaling that regulates cell proliferation, growth, apoptosis, and inflammatory responses [34, 35]. Studies on severe acute pancreatitis (SAP) in rats have found that JAK2 and STAT3 protein expression levels were significantly increased following induction of SAP [36]. In our study, results revealed that FAdV-4 infection can activate JAK2/STAT3 signal transduction and induce the expression of inflammatory factors, leading to liver inflammation and hepatic necrosis.

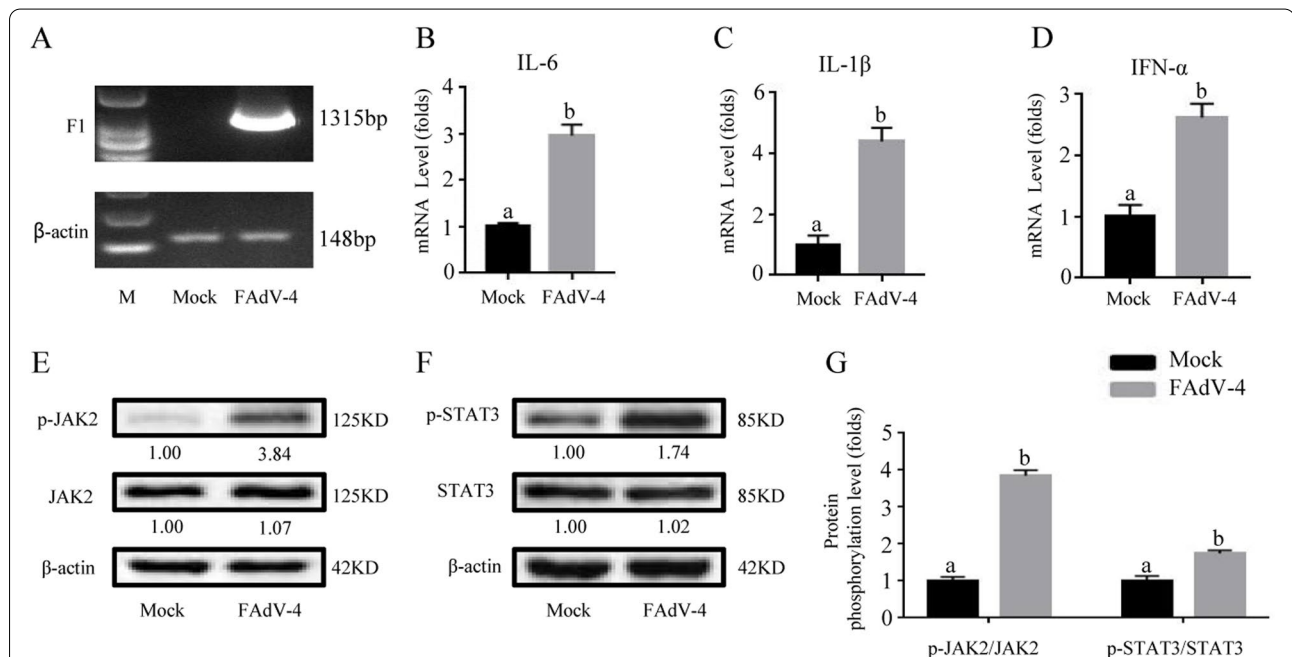
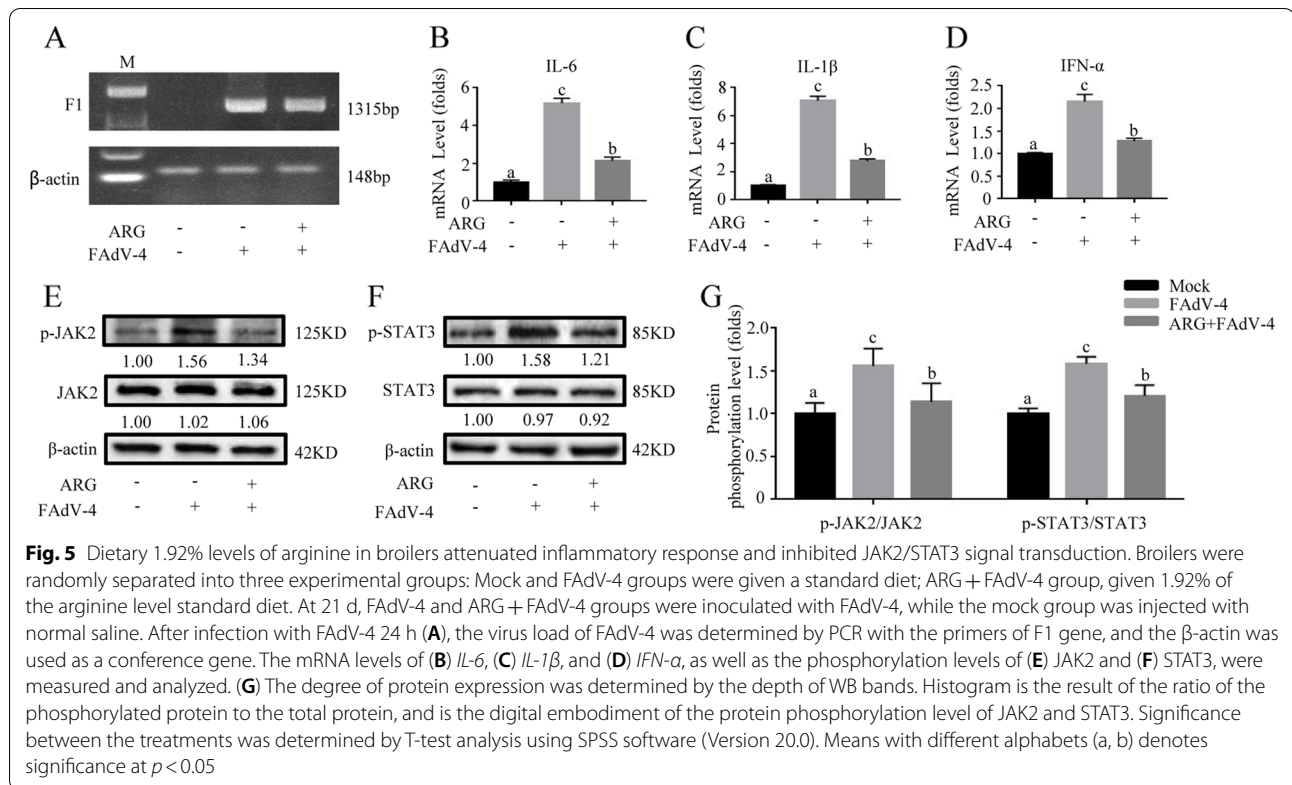


Fig. 4 FAdV-4 induced the expression of inflammatory factors and activated the JAK2/STAT3 pathway in the broiler liver. After infection with FAdV-4 24 h (A), the virus load of FAdV-4 was determined by PCR with the primers of F1 gene, and the β-actin was used as a conference gene. The mRNA levels of (B) IL-6, (C) IL-1β, and (D) IFN-α, as well as the phosphorylation levels of (E) JAK2 and (F) STAT3, were determined by real-time qPCR and Western blot in the liver of broilers. (G) The degree of protein expression was determined by the depth of WB bands. Histogram is the result of the ratio of the phosphorylated protein to the total protein, and is the digital embodiment of the protein phosphorylation level of JAK2 and STAT3. Significance between the treatments was determined by T-test analysis using SPSS software (Version 20.0). Means with different alphabets (a, b) denotes significance at $p < 0.05$



The prevention and control of FAdV-4 infection, especially in response to inflammation, was still not very successful [37]. Recent studies found that arginine regulated the immune system, specifically reducing inflammation [38]. Jiang found that when arginine was added both to the diet of Jian carp or a primary enterocyte culture media, the inflammation induced by lipopolysaccharide was significantly suppressed [39]. Our previous study revealed that FAdV-4 infection disturbed arginine metabolism to benefit its replication [17], and the addition of dietary arginine inhibited FAdV-4 replication [26]. In this present study, we further revealed that arginine treatment significantly inhibited the mRNA levels of IL-6, IL-1β and IFN-α in the broiler livers and LMH cells induced by FAdV-4, respectively.

The JAK/STAT was a classic familial signaling pathway in which each molecule has many subtypes. At the same time, each subtype has different modes of action and phosphorylation sites, which could regulate the different biological properties of cells [40]. In particular, the JAK2/STAT3 pathway has been extensively studied for its inflammatory regulation [12]. Specifically, JAK2 was important for cytokine receptor signaling. Upon activation, JAK2 kinase phosphorylates STAT3, causing inflammation [40, 41]. Evidence suggested that blocking

the JAK2/STAT3 signaling transduction could inhibit inflammation [42, 43]. By targeting JAK2/STAT3 signal transduction, it interfered with the phosphorylation of JAK2, a key protein in the pathway, thus blocking the phosphorylation and activity of downstream STAT3 protein, and provided strong protection against cell damage induced by small-molecule substances such as cytokines [42]. Intervention with AG490 (a highly selective JAK2 inhibitor) in a rat model of inflammatory bowel disease (IBD) significantly decreased the levels of IL-6 and IL-17A and increased the levels of IL-10, suggesting a protective effect on IBD [23]. Our present results indicated that arginine could inhibit JAK2/STAT3 signal transduction in inflammatory conditions induced by FAdV-4.

To explore whether JAK2/STAT3 signaling pathway was the leading way for arginine to alleviate FAdV-4-induced inflammation by comparing the inhibition of the JAK2/STAT3 pathway and the expression of inflammatory factors between arginine and AG490 (specific inhibition of JAK2 protein). The results revealed that both arginine and AG490 treatments could reduced inflammation response-induced with FAdV-4. Furthermore, the interesting discovery was that co-treatment with arginine and AG490 further promoted the inhibition of inflammation induced by FAdV-4.

Conclusion

In summary, arginine down-regulates inflammatory response induced by FAdV-4 via JAK2/STAT3 pathway in vivo and in vitro, which will provide new insight into the prevention against FAdV-4 infection.

Abbreviations

ARG: Arginine; CO₂: Carbon dioxide; DEGs: Differential expression genes; DMSO: Dimethyl sulfoxide; ECL: Enhanced chemiluminescence; FAdV4: Fowl adenovirus serotype 4; FBS: Fetal bovine serum; HHS: Hepatitis-hydropericardium syndrome; HRP: Horse radish peroxidase; IFN- α : Interferon- α ; IL-6: Interleukin-6; IL-1 β : Interleukin-1 β ; JAK: Janus kinase; KEGG: Kyoto encyclopedia of genes and genomes; LMH: Chicken hepatocellular carcinoma cell line; PVDF: Polyvinylidene fluoride; RIPA: Radio immunoprecipitation assay; PBS: Phosphate-buffered saline; PCR: Polymerase chain reaction; qPCR: Real-time fluorescence quantitative polymerase chain reaction; RIPA: Radio-immunoprecipitation assay; SDS-PAGE: Sodium dodecyl sulfate–polyacrylamide gel-electrophoresis; STAT: Signal transducers and activators of transcription; TNF- α : Tumor necrosis factor- α ; TBST: Tris buffered saline tween; WB: Western blotting.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12917-022-03282-9>.

Additional file 1: Supplementary Material 1. Original data of WB.

Additional file 2: Supplementary Material 2. Original data of PCR.

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Not applicable.

Authors' contributions

Quanxi and Changkang conceived and designed all the experiments, did a critical reading and revising of this present manuscript. Silin and Ruiling carried out the experiments. Both Silin and Qing analyzed the data in the current study. Then Lihui performed the statistics by SPSS software. Silin wrote and revised the manuscript. All authors have reviewed and approved the final manuscript.

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Availability of data and materials

All data generated or analysed during this study are included in this published article [and its supplementary information files].

Declarations

Ethics approval and consent to participate

All animal experiments were conducted following the guidelines of the Research Ethics Committee of the College of Animal Science, Fujian Agriculture and Forestry University. The animal protocol used in this study was approved by the Research Ethics Committee of the College of Animal Science, Fujian Agriculture and Forestry University, China (permit number: PZCAS-FAFU2016004). The experiments were also carried out in compliance with the ARRIVE guidelines.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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