

Response profiles of cytokines and chemokines against avian H9N2 influenza virus within the mouse lung

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Abstract The circulation of H9N2 viruses throughout the world, along with their expanded host range, poses a potential health risk to the public, but the host responses to H9N2 virus in mammals were little known. To obtain insight into the host immune responses to the avian H9N2 virus, the expressions of both cytokines and chemokines in the lungs of infected mice were examined by real-time polymerase chain reaction and enzyme-linked immunosorbent assay. We found that interferon gamma (IFN- γ) was the dominant antiviral component, and IFN- γ -induced protein 10 kDa, interleukin 6, chemokine (C–C motif) ligand 5 and macrophage inflammatory protein-1 alpha all

played a role in pro-inflammatory responses to H9N2 viruses. In conclusion, this research can make us further understand the infection characteristics of H9N2 virus in mammalian host by providing the data on mice lung immune responses to the avian H9N2 virus.

Keywords Avian influenza virus · H9N2 · Mice · Cytokines and chemokines

Introduction

H9N2 avian influenza virus (AIV) has been circulating worldwide in many avian species and resulted in great economic losses [1–3]. More importantly, human cases of avian H9N2 virus infection have been reported in Hong Kong and mainland China since the late 1990s [4–6], and seroprevalence investigations of H9N2 in poultry workers were also the solid evidence on human cases of H9N2 infection [7–9]. The above-mentioned researches have intrigued a great concern of the public.

H9N2 infection in mammals mainly depends on the ability of the virus to bind the human-like α 2,6-linked sialic acid (SA- α 2,6) receptors, and the HA receptor-binding site is critical for virus host range [10–12]. Some isolates of the H9N2 influenza viruses circulating in poultry can infect humans due to the ability to binding the SA- α 2,6 receptors [10, 13]. The expanded receptor specificity of H9N2 AIVs has raised concerns about their pathogenicity in humans.

H9N2 influenza viruses of chicken origin cause only mild symptoms in humans, but they have the pandemic potential for the high level of genetic plasticity [14–16]. For example, sequencing analyses of the novel influenza A (H7N9) virus isolated in China showed that 6 out of 8

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fragments were from H9N2 [17]. Facing the threat of H9N2 AIV, understanding the mammalian host immune responses to the virus is of importance to cope with the possible pandemic. However, up to date, little information about mammal immune responses to the H9N2 virus of chicken origin was reported.

To fill the literature gap, therefore, the expression of six cytokines [interferon beta (IFN) β , interferon gamma (IFN- γ), tumor necrosis factor (TNF) α , interleukin (IL) 1 β , IL-6, IL-10] and five chemokines [IFN- γ -induced protein 10 kDa (IP-10), chemokine (CC motif) ligand 5 (CCL-5), monocyte chemoattractant protein-1 (MCP-1), macrophage inflammatory protein-1 alpha (MIP-1 α), IL-8] were evaluated in the lungs of H9N2-infected mice with the aim to make us further understand the infection characteristics of H9N2 AIV in mammalian host.

Materials and methods

Ethics statement

All animal experiments were reviewed and approved by the Institutional Animal Care and Use Committee of Shandong Agricultural University and performed in accordance with the “Guidelines for Experimental Animals” of the Ministry of Science and Technology (Beijing, China). Animal suffering was minimized as much as possible.

Virus

The H9N2 virus, A/chicken/Shandong/w3/11/H9N2 (SDw3), belonging to the BJ94-like lineage was isolated from diseased chicken in Shandong. The virus was passaged in 10-day-old specific pathogen-free (SPF) chicken

embryos. And the 50 % tissue-culture-infective dose (TCID₅₀) were calculated by the method of Reed and Muench [18].

Mice experiments

Twenty-eight SPF female BALB/c mice (18.0–20.0 g, 6–8 weeks) from Experimental Animal Center of Shandong Province were randomly divided into the infected group and the control group. After being lightly anesthetized with CO₂, the mice were inoculated intranasally with 10⁶ TCID₅₀ of SDw3 in fifty microliters of phosphate-buffered saline (PBS) or PBS alone.

At 0.5, 2 and 6 days post-inoculation (dpi), lungs, hearts, livers, spleens, kidneys and brains of mice (3/group) in the infected group and the control were, respectively, collected for detecting SDw3 virus titers. Simultaneously, the mice lungs were used for cytokines and chemokines detection by real-time polymerase chain reaction (PCR) or enzyme-linked immunosorbent assay (ELISA) [19]. Additionally, the mouse lungs collected at 5 dpi were fixed in 10 % phosphate-buffered formalin, embedded in paraffin, then cut into 5-mm-thick sections and stained with hematoxylin and eosin (H&E) [20]. During the raising period, the weight loss and clinical signs of the mice were monitored and recorded. After the experiment, the mice were killed due to ethical reasons.

RNA extraction, cDNA preparation and real-time PCR

The RNA of the lung samples was extracted using an RNeasy Mini Kit (Qiagen, Valencia, USA). The quantity and quality of the isolated RNA were determined by UV 260/280 using a biophotometer (Eppendorf, Hamburg, Germany). A total of 500 ng RNA was used to prepare

Table 1 Primer sequences used for real-time PCR

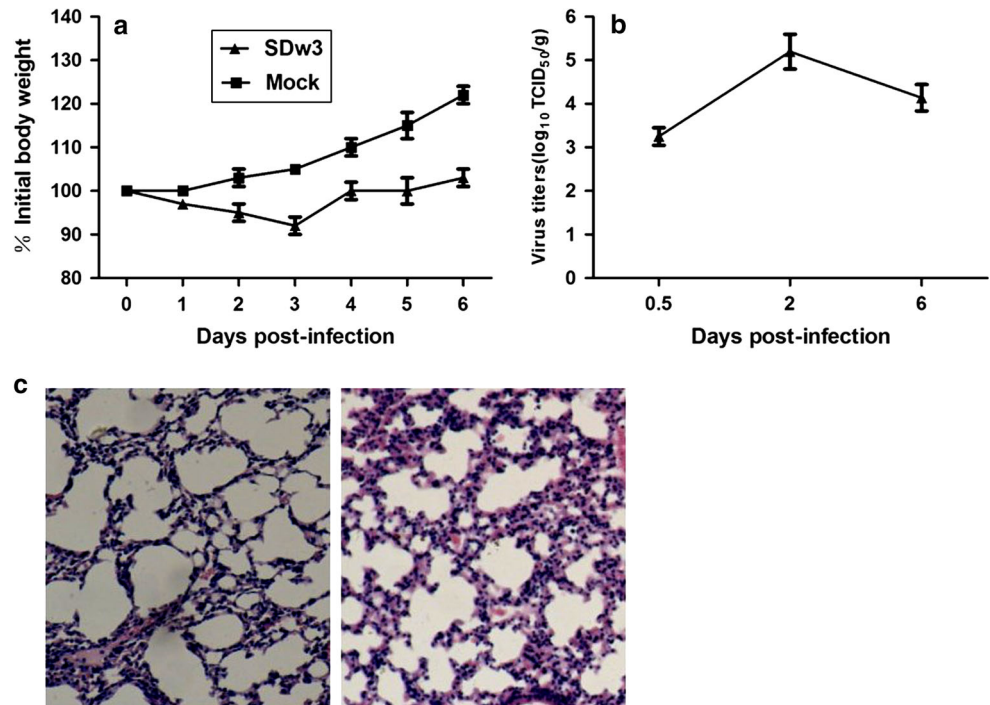
		Sequence (5' → 3')	Length(bp) ^a	Accession ^b
IFN- γ	FP	CGGCACAGTCATTGAAAGCCTA	199	NM_008337.3
	RP	GTTGCTGATGGCCTGATTGTC		
IL-6	FP	CCACTTCACAAGTCGGAGGCTTA	112	NM_031168.1
	RP	GCAAGTGCATCATCGTTGTTTCATA		
IP-10	FP	GTCCGCTGCAACTGCATCCATA	135	NM_021274.2
	RP	CTGCTCATCATTCTTTTCATCGTG		
CCL-5	FP	CTGCTGCTTTGCCTACCTCTCCC	156	NM_013653.3
	RP	TATTCTTGAACCCACTTCTTCTCTG		
IL-8	FP	CCGTCCCTGTGACACTCAAGA	178	NM_011339.2
	RP	TGGAGCATCAGGATCCAAACAA		

FP forward primer, RP reverse primer

^a Amplicon length in base pairs

^b GenBank accession number of cDNA and corresponding gene, available at <http://www.ncbi.nlm.nih.gov>

Fig. 1 Virulence of SDw3 in mice. Mice were inoculated intranasally with 10^6 TCID₅₀ of SDw3 or PBS. **a** Body weight were monitored and recorded daily. **b** Virus titers in lungs of SDw3-infected mice. The lungs of SDw3-infected mice ($n = 3$ per time point) were collected at 0.5, 2 and 6 dpi for virus titration. Virus titers were given in units of log₁₀ TCID₅₀ per g. Bars represent mean \pm standard deviation (SD) of three mice. **c** Histopathology of mock (*left*) and SDw3-infected (*right*) mice. Lungs were collected at 5 dpi and fixed in 10 % formalin, embedded in paraffin and sectioned (magnification $\times 100$)



cDNA by the reverse transcription reaction with a Prime-Script RT reagent Kit (TaKaRa, Dalian, China).

Real-time PCR was performed using SYBR Premix Ex Taq kit (TaKaRa, Dalian, China) and a standard 7500 Real-Time PCR System (Applied Biosystems, Foster City, USA) to obtain the relative expression quantity of message RNAs (mRNAs) of six cytokines and five chemokines (β -action was used as housekeeping genes) [21, 22].

Except that the primers (IFN- β , TNF- α , IL-1 β , IL-10, MCP-1, MIP-1 α and β -action) were cited according to the published references [23, 24], the other primers (Table 1) designed using Premier 5.0 software (Applied Biosystems, Foster City, USA).

Enzyme-linked immunosorbent assay

The protein levels of cytokines and chemokines in lung homogenates were measured using the mouse Quantikine Kits (RayBiotech Inc., Norcross, GA, USA; R&D Systems, Minneapolis, MN, USA). Briefly, the entire lung of each mouse was homogenized individually in 500 μ l Cell/tissue lysis buffer (RayBiotech Inc., Norcross, GA). After centrifugation at 10,000g for 15 min, the supernatants were collected for analyzing cytokines and chemokines simultaneously according to the kit instructions. The plates were read on a spectrophotometer at wavelength 450 nm by a microplate reader (Bio-Rad, Richmond, CA, USA). Finally, the cytokine or chemokine levels were recorded as ng/ml homogenate.

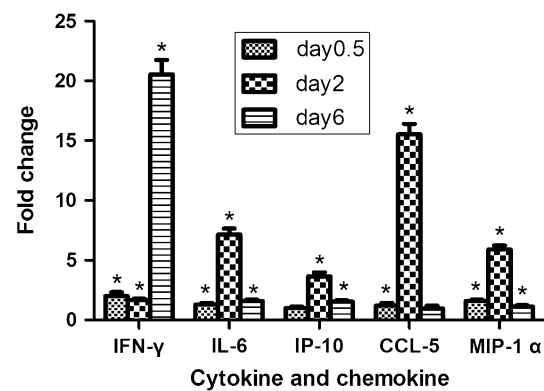


Fig. 2 Cytokine and chemokine mRNA response of mouse lung to H9N2. Cytokine and chemokine mRNA expressions in lungs of mice ($n = 3$ per time point) were measured by real-time PCR and were presented as the mean fold change (\pm SD) compared with values in mock mice at the days after inoculation. Statistical analysis was performed by comparing with the data of mock mice. Asterisk $p < 0.05$ between mock and SDw3-infected mice

Statistical analysis

Statistical analysis was performed using the Statistical Product and Services Solutions, version 10.0 (SPSS, Cary, NC, USA). ANOVA analysis was performed by comparing infected data to uninfected data. A p value < 0.05 was considered statistically significant.

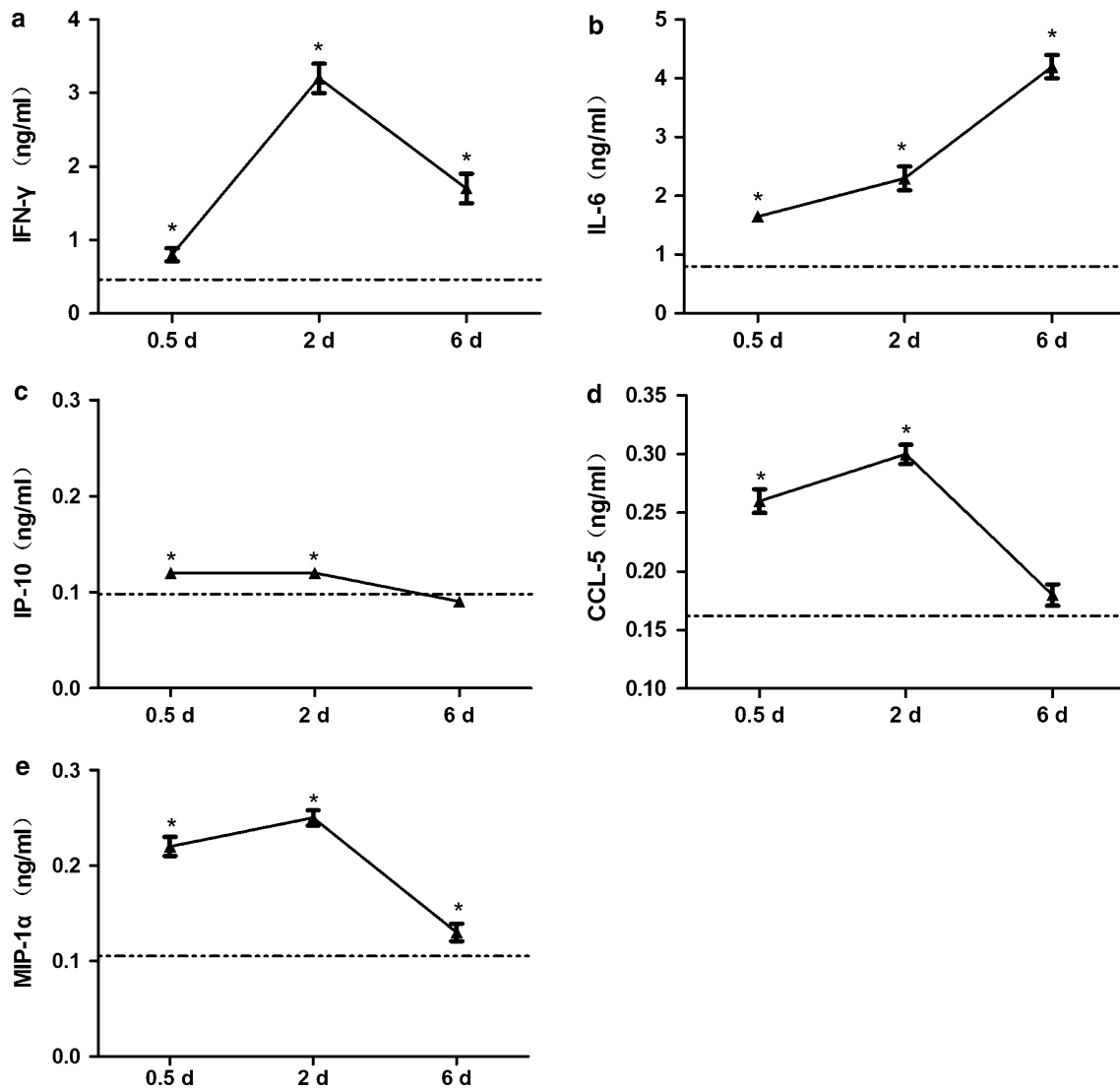


Fig. 3 H9N2 stimulates cytokine and chemokine release in mouse lung. Mice were inoculated intranasally with 10^6 TCID₅₀ of SDw3 or PBS. The lungs of SDw3-infected mice ($n = 3$ per time point) were harvested and homogenized in 500 μ l of Lysis Buffer. The protein of cytokines and chemokines were measured by ELISA. Baseline

protein levels of cytokines and chemokines from PBS-inoculated mice ($n = 5$) were shown as a dashed line in each graph. The data was presented as mean \pm SD for three independent experiments. Asterisk $p < 0.05$ between mock and SDw3-infected mice

Results

Pathogenicity of SDw3 in mice

Upon being injected with SDw3, the mice showed mild signs of illness, but they recovered naturally at 6 dpi. The body weight of infected mice dropped lightly during the first 3 days, but they almost regained the original body weight at 6 dpi (Fig. 1a). The virus titer was detectable only in the mice lungs, with a peak at 2 dpi (Fig. 1b). H&E staining of the infected mice lungs (5 dpi) showed that SDw3 caused mild and limited interstitial pneumonia; the interstitial tissue was

lightly thickened and filled with immune cells and inflammatory compared with the mock lungs (Fig. 1c).

Cytokine/chemokine mRNA expression

After infection, there was a two- to 20-fold mRNA induction of IFN- γ , IL-6, IP-10, CCL-5 and MIP-1 α ; the concentrations of CCL-5 and IFN- γ were, respectively, increased up to over 15-fold at 2 dpi and 20-fold at 6 dpi (Fig. 2). The data showed the peak induction for most of the mRNAs (IL-6, IP-10, CCL-5 and MIP-1 α) occurred at 2 dpi, except for IFN- γ which was most increased at 6 dpi.

By contrast, the other mRNAs were not up-regulated significantly (data not shown).

Cytokine/chemokine protein profiles

Protein expressions of the five corresponding cytokines and chemokines (IFN- γ , IL-6, IP-10, CCL-5, and MIP-1 α) were confirmed by ELISA, and their protein levels were all above the control at 0.5 dpi (Fig. 3). Protein expressions of IFN- γ , CCL-5 and MIP-1 α peaked at 2 dpi, whereas IL-6 at 6 dpi.

Discussion

SDw3, the dominant viral isolate in China, belongs to the BJ94-like lineage [25, 26]. The pathogenicity result of this study showed that the virus was replicable and pathogenic in mice without prior adaptation, which is consistent with the previous experiments [27, 28]. This was the main reason for us to further understand the immune responses of mice challenged by the virus. During the viral infection, body weight and clinical signs of mice changed lightly mainly due to the mild pathogenicity of the SDw3.

Cytokines play an important role in clearance of virus. For example, IFN- γ mediates the production of nitric oxide, subsequently resulting in the recruitment of more neutrophils and macrophages [29, 30]; IL-6 is a multi-functional cytokine that not only regulates immune and inflammatory responses involved in the activation, growth and differentiation of T-cells, but also contributes to T cell-mediated inflammatory reactions [31]. The analyses of cytokine in this study showed that IFN- γ and IL-6 were both up-regulated in H9N2-infected mice lungs. Compared with cytokine changes in H9N2-infected chicken lungs, TNF- α , IFN- α , - β and IFN- γ were all up-regulated [32]. The differences of the breed and the immune system may be the major contributors. During the influenza virus infection, the production of chemokines, including MCP-1, MIP-1, MIP-1 α , CCL-5, IP-10 and CCL-5, was all or partly up-regulated [33, 34]. This response is crucial in pro-inflammatory responses to influenza viruses. Similarly, MIP-1 α , CCL5 and IP-10 were all up-regulated in this study, which indicated that protective responses occurred in H9N2-infected mice lungs.

Time point selection of detecting cytokines and chemokines mRNA/protein expressions and observing histopathology changes was based on previous references [19, 20]. Although this selection was not perfect, it can truly reflect main change trend of cytokines and chemokines in mice lungs challenged by H9N2 influenza virus. Additionally, during the experiment, production changes of mRNA and protein at the same time point were not

consistent, which can largely be attributable to the time difference between transcription and translation.

In conclusion, after infection of H9N2, IFN- γ was the dominant antiviral component, and IP-10, IL-6, CCL-5 and MIP-1 α all played an important role in pro-inflammatory responses to H9N2 viruses. It is useful for us to understand the infection characteristics of H9N2 virus in mammalian host.

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Conflict of interest The authors declare that they have no conflict of interest.

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