Extraction and purification of the H9N2 virus nucleoprotein: A simple and practical method

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Received: 30 May 2018 Published: 21 Dec 2018

Abstract

Background: Avian Influenza disease annually entails a significant economic loss to the poultry industry around the world. Influenza virus is a polymorphic virus of the orthomyxoviridae family (single-stranded RNA genome), and nucleoprotein (NP) is the structural and internal protein of the virus. The aim of the work was to purify nucleoprotein for further investigations with a simple, low-cost, fast and practical method.

Methods: In this study, H9N2 influenza virus was isolated in specific pathogen-free embryonated chicken eggs by allantoically inoculating 10^3 to 10^5 egg-infective doses (EID50) for 9 to 11 days, purified by 10% (W/V) polyethylene glycol (PEG) 6000 with a sucrose gradient of 60% to 30%. The influenza virus proteins were collected and prepared as fractions by preparative electrophoresis. Finally, the purified NP was subjected to sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and Western blot procedures.

Results: The protein analysis with SDS-PAGE and silver nitrate staining indicated that the desired samples contained purified nucleoprotein and lacked other viral proteins. The results of the investigation of lyophilized fractions containing nucleoprotein on the SDS-PAGE revealed the absence of viral RNA in nucleoprotein and its high purity.

Conclusion: According to this study, purified nucleoprotein can be used to produce nucleoprotein vaccines, as well as to study structural, molecular and diagnostic and therapeutic materials.

Keywords: Avian influenza virus, H9N2, Nucleoprotein, Extraction, Purification

Conflicts of Interest: None declared
Funding: None

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Introduction

Avian Influenza is a highly contagious disease that causes symptoms in respiratory, digestive and nervous systems in a wide range of birds and can be transmitted to mammals, including humans (1). Dramatic changes reasortment between human and avian viruses are due to an
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Viral propagation

In this experimental study, viral propagation was performed by the method described (14). Briefly, the isolates of avian influenza A/Chicken/Iran/259/2014/H9N2 were propagated in specific-pathogen-free chicken embryonated eggs by allantochorial inoculating 10^2 to 10^3 egg-infective doses (EID₅₀) for 9 to 11 days. Eggs were incubated at 37°C with relative humidity of 55% and were candled daily for 7 days to embryo viability. Allantoic fluid from embryos that died after 24 h was collected aseptically and tested for hemagglutinating activity. If no deaths occurred after 7 days in the eggs, the embryo was opened aseptically. The allantoic fluids were pooled and inoculated into 5 more embryonated eggs. If no deaths occurred after 7 days in the eggs of the second passage, all the eggs were opened, and the allantoic fluid of each egg was tested for hemagglutinating activity. If deaths occurred in the eggs and the material had a positive HA test, samples were considered positive, and this result was confirmed with the hemagglutinating inhibition (HI) test. Allantoic fluids from embryos showing early mortality after inoculation were tested for bacterial contamination according to routine procedures, and subsequently the infected fluids which contained between 32×10⁵ EID 50 of virus per ml were harvested and pooled. Reference stocks were lyophilized and working stocks were stored in ampoules at -20°C (15). However, HA and HI tests were performed by standard methods (15).

Extraction and Purification of nucleoprotein

The mixture was incubated for 1 hour at 4°C and clarified by centrifugation at a speed of 3500 rpm and 4°C for 15 minutes. The supernatant was collected, and PEG 5% (w/v) was added to it. Then, clarified by centrifugation at a speed of 9000 rpm and 4°C for 60 minutes, precipitate in TNB buffer was created as a homogenous suspension. Finally, a homogenous suspension was prepared for dialysis to remove PEG from the suspension. The HA test was performed on the sample after dialysis. Then, the dialyzed fluid was ultra-centrifuged at a speed of 20000 rpm and 4°C for 2 h. The resulting precipitate was converted into homogenous suspension in 100 ml of TNE buffer. The HA test was also carried out at this stage. The purification of virus was performed by sucrose density-gradient centrifugation method (16). Finally, the resulting virus samples were lyophilized.

Protein Assay

For this purpose, Bradford method was performed to determine the protein concentration of the sample according to the previously described method with some modifications (17). In this method, duplicate volumes of bovine serum albumin (BSA) (10-100 mg/ml) was used to prepare the calibration graph. The basic Laemmli SDS-PAGE procedure was used to separate the protein of interest (18). Afterward, one of the gels was stained with Coomassie blue G-250 (Merck, Germany) and the other gel with silver nitrate. Also, native and functional biological molecules were purified by preparative electrophoresis (model 491 Prep Cell, Bio-Rad, CA), using reduced and non-reduced methods (19). Finally, to determine protein fractions the optical density was read at 280 nm using a spectrophotometer. The reduced 12% SDS-PAGE was used to analyze the integrity of the purified influenza virus proteins and the detection of fractions containing nucleoprotein. In brief, 200 μl of each protein fraction [with a high absorption] was mixed with 30 μl of 2-mercaptoethanol and incubated for 4 h at 37°C. The protein fractions [with high absorption] were dissolved in non-reducing SDS-PAGE loading buffer (without 2-mercaptoethanol) and stained with silver nitrate (20). The protein concentration was assessed with the Bradford method. Finally, in order to confirm H9N2 presence, the purified...
sample was loaded and resolved by SDS-PAGE and proteins were transported to polyvinylidene difluoride (PVDF) membranes. Membrane blocking was done using 5% bovine serum albumin (BSA), overnight. Primary specific antibody versus H9N2 was used at a 1:1000 dilution, and secondary antibody was used at a 1:10000 dilution (HRP-conjugated). Immunoreactive bands visualized with chemiluminescence HRP substrate (Abcam).

Results

Protein assay before SDS-PAGE
A protein assay was performed for influenza virus, and at this stage, BSA was used as a standard in volumes of 100, 300, 500 and 1000 mg/ml, and the results indicated that the volume of the viral solution was appropriate. According to Table 1, the highest and lowest amounts of protein were the sample F act 3 and the sample F act 5 with 387.07 mg/ml and 212.5 mg/ml, respectively.

SDS-PAGE after protein assay
Gels were prepared with concentrations of 2.5, 5 and 10 μg of each sample. According to our results (Fig. 1), the molecular weight of the nucleoprotein was estimated between 45 and 66.2 kDa.

Preparative electrophoresis results
The absorption of the fractions was obtained from the purified nucleoprotein at 280 nm (UV). Graphs 1 and 2 were plotted from the 121 fractions of the reduced and, 123 fractions of the non-reduced methods. Our results showed that 26 fractions from the reduced and 42 fractions from the non-reduced methods had a positive absorption, and these fractions contained viral NP.

SDS-PAGE and Western blotting after protein purification
SDS-PAGE was performed for the presence of NP after purification with preparative electrophoresis method (Figs. 2 and 3). According to the molecular weight of NP, fractions 2 and 3 (reduced purification) contained viral NP, but in fractions with non-reduced purification, no band was detected, and no viral protein was isolated. The NP band (H9N2) was verified again by immunoblotting assay (Fig. 4).

Discussion
The purpose of this work was to prepare and purify the influenza virus nucleoprotein in a simple, low-cost, applied and rapid method. In the current study, firstly the H9N2 virus identified and the locations of the viral proteins were determined. Subsequently the nucleoprotein was purified with adequate and accurate information about the viral proteins. The use of a live virus for propagation is necessary, but epidemiological hygiene needs to be considered, and this issue is more important in influenza viruses due to their genetic rearrangement and transmission potential to humans. However, the advantages of using the inactive virus are due to the lack of infectious potential and long-term maintenance and the absence of contamination with other microorganisms. The virus from dialysis was lyophilized to facilitate further protein purifications. By SDS-PAGE analyses, the virus protein bands were examined, and it was determined that a nucleoprotein with a molecular weight ~ 54 kDa is clearly identifiable [between bands of 45 to 65 kDa of ladder], and two bands (top and down), which are related to the HA protein, can be identified and separated. However, to ensure that the NP protein can be observed in low amounts of protein and the diluted sample, gels are stained with silver nitrate, which is about 10 to 100 times more accurate than the Coomassie blue. For purity and the presence of the desired protein, we examined the molecular weight and position of the nucleoprotein in terms of movement and detachment in the polyacrylamide gel; samples were run on a preparative electrophoresis machine. The Prep Cell 491 has a special constructional complexity and requires the

Table 1. Results of Protein assay

<table>
<thead>
<tr>
<th>Sample name</th>
<th>OD 595nm</th>
<th>Protein mg/ml</th>
</tr>
</thead>
<tbody>
<tr>
<td>St 100</td>
<td>0.08</td>
<td>100</td>
</tr>
<tr>
<td>St 300</td>
<td>0.31</td>
<td>300</td>
</tr>
<tr>
<td>St 500</td>
<td>0.53</td>
<td>500</td>
</tr>
<tr>
<td>St 1000</td>
<td>0.97</td>
<td>1000</td>
</tr>
<tr>
<td>F act 1</td>
<td>0.22</td>
<td>212.9</td>
</tr>
<tr>
<td>F act 2</td>
<td>0.32</td>
<td>309.67</td>
</tr>
<tr>
<td>F act 3</td>
<td>0.40</td>
<td>387.07</td>
</tr>
<tr>
<td>F act 4</td>
<td>0.27</td>
<td>261.29</td>
</tr>
<tr>
<td>F act 5</td>
<td>0.17</td>
<td>212.5</td>
</tr>
</tbody>
</table>

Fig. 1. SDS PAGE gel for absorption (reduction method) protein fractions stained silver nitrate
1. F act 4 (10mg), 2. F act 4 (5mg), 3. F act 4 (2.5mg), 4. F act 5 (10mg), 5. F act 5 (5mg), 6. F act 5 (2.5mg)

Fig. 2. SDS PAGE gel for absorption of (reduction method) protein fractions
1. Prestaining Marker, 2. Virus sample before purification, 3. Absorption fraction containing nucleoprotein, 4. Absorption fraction containing nucleoprotein...
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Graph 1. Absorption of 121 fractions from (reduced method) at 280 nm

Graph 2. Absorption of 123 fractions from (non-reduced method) at 280 nm

To check the accuracy of the results, the influenza virus before and after lyophilization, and a purified nucleoprotein after lyophilization were tested. The results of electrophoresis indicated no genomic band in the nucleoprotein sample. Thus, during testing, it has been proven that with the help of the Prep Cell 491, we will be able to completely purify the H9N2 avian influenza virus and save time and cost. Purified nucleoprotein can be used to provide nucleoprotein anti-sera against influenza viruses in future researches. Significant points in this study are the high speed and precision of the test results, which can be expressed as a superior advantage over the success of the research. Studies have been conducted on nucleoprotein and different methods used for purification of nucleoprotein. Harmon et al., investigated an immunoassay for serological diagnosis of influenza type using a recombinant DNA producing nucleoprotein antigen and monoclonal human antibody IgG (21). They used immunoaffinity chromatography method for NP purification. This method is time-consuming and
costly, and although the appropriate amount of nucleoprotein was purified, supplementary experiments and a large number of viral samples are required. In the present study, with the least number of the viral samples, nucleoprotein can be obtained. Li et al., (2014), isolated and purified the NP by SDS-PAGE for NP purification and then mass spectrometry was performed (22). In the SDS-PAGE electrophoresis method, viral proteins are isolated but not completely purified, and when cutting from the gel, there is a possibility of error. On the other hand, in this method, the accuracy and skill of the researcher are of great importance, and the cost of mass spectrometry is significant for confirmation of the protein.

Conclusion
Considering the results of our study, the purified nucleoprotein was obtained very dynamically, with high speed, precision, and purity. Purified nucleoprotein can be used to produce nucleoprotein vaccines, as well as to study structural, molecular and diagnostic and therapeutic materials.

Acknowledgments
We thank our colleagues from Department of Botany, Biotechnology and Bioinformatics, Payame Noor University who provided insight and expertise that greatly assisted the research.

Conflict of Interests
The authors declare that they have no competing interests.

References