Single-Step Influenza and Dengue Virus Purification with Mixed-Mode CHT Ceramic Hydroxyapatite XT Media

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Virus Purification

Abstract

Past studies have shown that CHT Ceramic Hydroxyapatite Media can be used for virus purification. We purified influenza and dengue viruses on the newest member of the CHT family, CHT XT Media. We recovered over 75% of each virus in a single step of purification. In addition, >90% reduction of contaminating proteins and >60% removal of dsDNA were seen in both samples. This positions CHT XT as an additional media for simple, fast, and reproducible viral purification and overcomes the shortcomings of conventional purification methods.

Introduction

Virus preparations have been used as vaccines for over two hundred years, and more recently for gene therapy platforms. For gene delivery, conventional viral purification methods such as density gradient ultracentrifugation and precipitation can be difficult to scale and may not meet the purity standards required for a therapeutic. Consequently, chromatography has emerged as an advanced tool to meet large-scale therapeutic virus purification requirements. Viruses bind to chromatographic resins by electrostatic interactions.

We have previously demonstrated mammalian virus purification with the mixed-mode CHT Ceramic Hydroxyapatite Media (Kurosawa et al. 2014). CHT Media are a group of mixed-mode calcium affinity/cation exchange supports. The high negative surface charge on many viruses allows them to bind tightly to CHT calcium sites, permitting significant purification with high (>80%) yields. Such tight binding has also been utilized to provide robust viral clearance, which can be used to ensure viral safety during biomanufacturing (ICH Expert Working Group 1999, Moritz 2005).

CHT XT is the newest addition to the CHT family of media. It has been designed for superior physical robustness to ensure it can be used repeatedly over a large number of cycles. It maintains the unique separation properties of CHT for aggregate removal and monoclonal antibody purification. In order to test its virus purification capabilities, we performed single-step purifications of influenza and dengue viruses using CHT XT Media.

Materials and Methods

The viruses used in this study are shown in Table 1.

Table 1. Virus type and size.

| Virus | Genome | Envelope | Size, nm |
|-----------|--------|----------|----------|
| Influenza | ssRNA | + | 80–120 |
| Dengue | ssRNA | + | 50 |

Virus Production

Influenza virus: Strain A/California/07/2009 was cultured on MDCK cells in minimal essential medium (MEM; Gibco) containing 10% fetal bovine serum (FBS) and L-glutamine in 225 cm² flasks. When the cells were almost confluent, influenza virus was inoculated onto the cell monolayer at a 1:3,000 dilution in 75 ml of D-MEM/F-12 with trypsin, penicillin, and streptomycin (flu culture medium). The supernatant was harvested on days 3, 4, and 11 and clarified through 0.45 µm filters.

Dengue virus: Type 2 strain ThNH7/93 was cultured on C6/36 cells. Cells were grown in 225 cm² flasks that were precoated with poly-L-lysine at 100 µg/ml in phosphate buffered saline (PBS). Cells were cultured in Minimum Essential Medium Eagle (modified) (modified EMEM, MP Biomedicals) containing 10% FBS at 28°C for 1 week. After the cells reached confluence, dengue virus type 2 was inoculated at a 1:1,000 dilution onto the cell monolayer in 75 ml of modified EMEM containing 0.5% FBS and MEM Vitamin Solution (Invitrogen Corporation) and cultured at 28°C. The medium was changed at day 3 and culture fluid (about 75 ml) was collected at day 7. The culture fluid was filtered through a 0.45 µm filter to remove cells and large cell debris.





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Chromatography

CHT XT Media was packed into 4.6 x 35 mm columns (0.58 ml bed volume) and viral purification was performed at a flow rate of 360 cm/hr as shown in Table 2 for influenza and Table 3 for dengue. All experiments were carried out in triplicate and representative data are shown.

Table 2. Influenza purification protocol.

| 1 M NaOH | | 5 |
|--|--|---|
| 600 mM sodium phosphate | 7.2 | 10 |
| 10 mM sodium phosphate | 7.2 | 10 |
| 24% of 600 mM sodium phosphate | 7.2 | 10 |
| 24% of 600 mM sodium phosphate | 7.2 | 24 |
| Gradient elution from 150–600 mM (24% to 100%) sodium phosphate | 7.2 | 11 |
| 600 mM sodium phosphate | 7.2 | 5 |
| | 600 mM sodium phosphate 10 mM sodium phosphate 24% of 600 mM sodium phosphate 24% of 600 mM sodium phosphate Gradient elution from 150–600 mM (24% to 100%) sodium phosphate 600 mM sodium phosphate | 600 mM sodium phosphate7.210 mM sodium phosphate7.224% of 600 mM sodium phosphate7.224% of 600 mM sodium phosphate7.2Gradient elution from 150–600 mM7.2(24% to 100%) sodium phosphate7.2600 mM sodium phosphate7.2 |

Table 3. Dengue purification protocol.

| Step | Mobile Phase | pН | Volume, ml |
|-------------------------------|--|-----|------------|
| Pre-wash | 1 M NaOH | | 5 |
| Wash | 600 mM sodium phosphate | 7.2 | 10 |
| Equilibration | 10 mM sodium phosphate | 7.2 | 20 |
| Sample loading and washing | 10 mM sodium phosphate | 7.2 | 20 |
| Elution | Gradient elution from 10–600 mM sodium phosphate | 7.2 | 15 |
| Column wash | 600 mM sodium phosphate | 7.2 | 5 |

Viral Activity Assay

Influenza viral activity in the recovered fractions was determined via hemagglutination analysis (HA test) with chicken red blood cells. Activity of the dengue virus in the recovered fractions was determined via HA test with goose red blood cells as described previously (Kurosawa et al. 2012).

Estimating Double-Stranded DNA (dsDNA)

The concentrations of dsDNA in fractions were determined using the Quant-iT PicoGreen dsDNA Assay Kit (Invitrogen Corporation) according to the manufacturer's instructions.

Protein Analysis in Virus Fractions

Viral fractions were analyzed for total protein concentrations with the Micro BCA Protein Assay Kit. Host cell protein (HCP) analysis for the influenza viral fractions was performed with the MDCK HCP ELISA Kit (Cygnus Technologies).

TCID₅₀ Influenza Virus Infectivity Assay

A confluent monolayer of MDCK cells in 96-well microplates was used for this test. MDCK cells (2 x 10⁴ cells/well) were cultured in flu culture medium for 1 day prior to the testing. Each virus fraction underwent tenfold serial dilution with flu culture medium. Diluted viral fractions (50 μ l) were added to each well (n = 3) and cultured for one week. Cytopathic effects were studied under a microscope. Titers were calculated using the Reed-Muench method.

Viral Purity Analysis

Viral purity was assessed by electrophoresis on a 12.5% SDS-PAGE gel and by immunoblotting. The protocols used for these analyses were as described previously (Kurosawa et al. 2012). Cell culture fluid was concentrated tenfold and all other samples were concentrated 30-fold by ultrafiltration with a 10 kD MWCO filter before use. Bio-Rad's Gel Doc EZ Imaging System was used to quantitate band density. For immunoblotting, primary antibody anti-HA mAb (Abnova) was used at a 1:200 dilution with the influenza virus samples and mouse anti-D2 serum (made in house by HOYA Technosurgical Corporation for these experiments) was used at a 1:500 dilution with the dengue virus samples. Secondary antibody anti-mouse IgG (Fc)-POD (American Qualex) was used at a 1:1,000 dilution. The bands were visualized with EzWestBlue (ATTO Corporation).

Results Influenza Virus Viral Activity

The chromatographic results from the influenza virus purification are shown in Figure 1. Influenza virus elutes at approximately 250 mM phosphate, with good separation from the bulk of UV-absorbing material and partial resolution from dsDNA. Hemagglutination activity (red bars on the chromatograms) measures the titer of virus that can bind to molecules present on the surface of red blood cells and cause agglutination. As seen in Figures 1A and 1B, HA activity is predominantly seen between retention volumes of 26 and 31 ml. The total viral activity recovered from these fractions was calculated to be about 75% of the starting activity. This activity was well separated from the bulk of the DNA and A280-absorbing impurities. The samples showing bulk viral activity were pooled for further analytical assays.



Fig. 1. Replicate elution of influenza virus from CHT XT. HA titer (—); DNA (—); A₂₈₀ (—); A₂₈₀ (—); conductivity (—).

Viral Infectivity

In order to determine if the purified influenza virus maintained its infectivity, a TCID_{50} test was performed (Figure 2) to assay for cytopathic effects (CPE). This test determines the quantity of active, infectious particles. A mean recovery of ~68% infectivity was calculated for the replicates. As seen in Figures 2A and 2B, the bulk of the infectivity is seen at retention volumes that also showed the highest HA activity. This confirmed that fractions 26 to 32 contained active, purified virus.



Fig. 2. Infectivity of the influenza virus purified on CHT XT. Infectivity (—); HA titer (—); A_{260} (—); A_{260} (—); conductivity (—).

Analytical Assays

Three analytical assays were performed on the pooled samples from the influenza virus run on CHT XT Media. The results of the analytical assays are shown in Table 4. Single-step influenza virus purification on CHT XT led to significant (>99%) clearance of total and host cell proteins as well as removal of over 60% of dsDNA.

| Table 4. Results from the analytical assays on the pooled influenza | |
|---|--|
| samples from the HA assay. | |

| Log Reduction | Log Reduction | Log Reduction |
|---------------|------------------|----------------|
| in dsDNA | in Total Protein | in HCP Content |
| 0.42 | 2.82 | 2.54 |

Viral Purity

SDS-PAGE analysis of the active pooled fractions from Figure 1 confirmed the significant purification of species migrating at the expected molecular weight for the two forms of the HA protein (Figure 3A). Immunoblotting of the same samples confirmed that the concentrated bands are HA (Figure 3B).

A

в



Fig. 3. SDS-PAGE (A) and immunoblotting analysis (B) of fractions. Protein bands on the gels were visualized by silver staining and the HA proteins in active samples were detected using an HA-specific monoclonal antibody. Lane 1: 3 μ l protein MW marker; lane 2: 10 μ l cell culture fluid; lane 3: 10 μ l flow-through fractions; lane 4: 10 μ l pooled fractions from HA assay.

Dengue Virus Viral Activity

The chromatographic results from the dengue virus purification are shown in Figure 4. Dengue virus elutes at approximately 400 mM phosphate, with good separation from the bulk of UV-absorbing material and dsDNA. HA activity was predominantly seen between retention volumes of 28 to 38 ml; fractions 30–35 were pooled for further analysis. The total viral activity recovered from these fractions was calculated to be about 87%. Viral activity was not detected in the flow-through fractions in any of the replicates.



Fig. 4. Replicate elution of dengue virus from CHT XT. HA titer (—); DNA (—); A₂₈₀ (—); A₂₆₀ (—); conductivity (—).

Analytical Assays

Two analytical assays were performed on the pooled samples from the dengue virus run on CHT XT. The results of the analytical assays are shown in Table 5. Single-step dengue virus purification on CHT XT led to significant clearance of protein (>93%) and dsDNA (>91%) contaminants.

Table 5. Results from the analytical assays on the pooled dengue samples from the HA assay.

| Log Reduction in dsDNA | Log Reduction in Total Protein |
|------------------------|--------------------------------|
| 1.07 | 1.15 |

Viral Purity

SDS-PAGE analysis of the pooled active retention fractions from Figure 4 showed that the pool contains proteins that migrate at the expected position for dengue E and C proteins as shown on the immunoblot (Figure 5). Note that a contaminant can be seen migrating immediately above the E protein on the SDS-PAGE gel.

Α

в



Fig. 5. SDS-PAGE (A) and immunoblotting analysis (B) of fractions. Protein bands on the gels were visualized by silver staining and the viral proteins in active samples were detected by antiserum against dengue virus type 2. Lane 1: 3 µl protein MW marker; lane 2: 5 µl dengue positive control; lane 3: cell culture fluid; lane 4: 10 µl flow-through fractions; lane 5: 5 µl UV-rich fractions; lane 6: 10 µl pooled fractions from HA assay.

Conclusion

Chromatography media are continuously evolving to meet the increasing demands of the bioprocessing industry. We developed the new CHT XT Media to meet the demand for a robust media with excellent life time and single-step purification functionality for healthy process economics. With both high purity and high yield for both influenza and dengue viruses, we show that CHT XT can be used for single-step impurity clearance. This positions CHT XT as a media for simple, fast, scalable, and reproducible viral purification.

References

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