Video Article A Method to Assess Fc-mediated Effector Functions Induced by Influenza Hemagglutinin Specific Antibodies

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Abstract

Antibodies play a crucial role in coupling the innate and adaptive immune responses against viral pathogens through their antigen binding domains and Fc-regions. Here, we describe how to measure the activation of Fc effector functions by monoclonal antibodies targeting the influenza virus hemagglutinin with the use of a genetically engineered Jurkat cell line expressing an activating type 1 Fc-FcqR. Using this method, the contribution of specific Fc-FcqR interactions conferred by immunoglobulins can be determined using an *in vitro* assay.

Video Link

The video component of this article can be found at https://www.jove.com/video/56256/

Introduction

Immunity provided by the seasonal influenza vaccine has traditionally been assessed by the hemagglutination inhibition (HI) assay¹, which measures the presence of antibodies that target the receptor binding site of the hemagglutinin. These HI-active antibodies can confer sterilizing immunity, but are generally narrow in their breadth of protection, providing immunity to only a select few strains of influenza virus. The isolation and characterization of broadly reactive monoclonal antibodies that recognize the hemagglutinin (HA) suggest the development of a universal influenza vaccine is within reach^{2,3,4,5,6,7,8}. One of the major aims of a universal influenza vaccine is to induce a strong antibody response towards the conserved regions of the influenza virus^{9,10,11,12,13,14,15,16}. Broadly reactive antibodies that recognize these conserved epitopes, composed of both neutralizing and non-neutralizing antibodies^{17,18}, have been shown to require Fc-FcγR interactions for optimal protection *in vivo* and highlight the contribution of Fc-mediated immunity to the overall immune response to influenza virus^{19,20}.

Correlates of protection are crucial in assessing immunity to infection. These metrics allow scientists and clinicians to estimate the efficacy of vaccines, the significance of the data, and the course of treatment. The only established correlate of protection against influenza virus infection is the hemagglutination inhibition assay. A titer of 1:40 is associated with a 50% reduction in the risk of illness^{1,21} – and reflects the presence of antibodies that inhibit agglutination by targeting the receptor binding site located on the globular head of the HA. However, it should also be noted that T-cell responses may be a better correlate of protection against influenza virus infection in the elderly. Interestingly, a recent report suggests that neuraminidase inhibition activity may be a better predictor of immunity to influenza²². Fortunately, typical *in vitro* assays such as neutralization or neuraminidase inhibition assays can measure HA stalk- or neuraminidase-specific antibodies. Most of these conventional *in vitro* assays, however, only take into account the function of the antigen binding region of the antibody and do not measure the role of the Fc region. Additionally, the contribution of non-neutralizing antibodies that protect via Fc receptor engagement *in vivo* are not detected^{17,18}. In order to measure protection afforded by antibodies that induce Fc-mediated immunity such as antibody dependent cell mediated cytotoxicity (ADCC), a robust *in vitro* assay is needed.

The method described below assesses the ability of murine monoclonal antibodies to induce Fc-mediated functions through the use of a genetically modified Jurkat cell line expressing an activating murine type 1 FcR, FcγRIV. Antibody engagement of the FcγR transduces intracellular signaling that triggers nuclear factor of activated T-cells-mediated luciferase activity. The assay has several advantages over traditional techniques that require isolation and culture of primary effector cells and the use of flow cytometry to detect activation of Fc-mediated effector functions^{19,23,24,25,26}. First, the protocol described here can easily be adapted to incorporate different viral targets, FcγRs, and antibodies from different species (human and murine). Second, the use of a modified Jurkat cell line expressing an FcγR with a luciferase reporter gene under a nuclear factor of activated T-cell (NFAT) allows for a large format assay that can be easily analyzed using a plate reader measuring luminescence. Here, the traditional activation of primary effector cells is replaced with the induction of NFAT-luciferase upon antibody engagement of an FcγR on the surface of the Jurkat cell. This 96-well plate format assay allows for the measurement of up to four different samples in triplicates with seven dilutions – a number of samples that could be cumbersome using traditional techniques. There are additional points to consider with this assay that may affect the design of experiments and the interpretation of data. The viral target must be expressed on the cell surface in order for antibody recognition. To mitigate this, the target antigen (*e.g.* an internal viral protein) may be directly coated on

the surface of a plate. However, this has not yet been rigorously tested. Additionally, the modified Jurkat cell line expresses only one type of activating FcγR and does not express any inhibitory FcγRs, while primary cell lines express all receptors in a physiological context.

We have previously used this assay to demonstrate that epitope specificity in a polyclonal context plays a crucial role in regulating Fc-mediated effector functions and that optimal activation of antibody-dependent cell-mediated response requires two points of contact^{27,28}. Here, we describe a method that assesses the ability of stalk-specific monoclonal antibodies to engage and activate the murine activating FcγRIV. Although there are exceptions^{29,30}, a large number of broadly reactive antibodies target the antigenically conserved stalk region of the hemagglutinin and thus play an important role in the development of a universal influenza vaccine.

Protocol

The experiments preformed in this manuscript were done following Icahn School of Medicine's institutional code of ethics and regulations.

1. Expression of Influenza Virus Hemagglutinin via Transfection or Infection

Transfection:

- 1. Plate Human Embryonic Kidney (HEK 293T) cells at a density of 2 x 10⁴ cells/well in a white tissue culture treated 96-well plate and let it sit for 4 h in a 37 °C incubator (with 5% CO₂).
- 2. Transfect cells with 100 ng of DNA coding for viral hemagglutinin and 0.2 µL of transfection reagent per well in a total volume of 50 µL.
- NOTE: Both plasmid and transfectant reagent are diluted in 1x Opti-Minimum Essential Media (MEM) Reduced Serum Medium.
- 3. After incubating plates in a 37 °C incubator with 5% CO₂ for 18 h, remove transfection media.

2. Infection

- 1. Plate A549 or Madin Darby canine kidney cells at a density of 2 x 10⁴ cells/well in a white tissue culture treated 96-well plate and grow for at least 18 h in a 37 °C incubator with 5% CO₂.
- Dilute virus in 1x MEM and infect cells at a multiplicity of infection of 3, 1, or 0.33 for 1 h in a 37 °C incubator with 5% CO₂. Total volume should equal 100 μL.
- NOTE: 10x MEM is diluted with water to make a 1x MEM working stock for virus dilution mentioned above.
- 3. After infection, remove the inoculum and add 100 µL of 1x MEM into the plate in the absence of trypsin.
- 4. Grow infected cells for 18 to 24 h in a 37 °C incubator with 5% CO_2 .

3. Assessing the FcRg/NFAT-mediated Activation of Luciferase Activity by Monoclonal Antibodies or Sera

- 1. First, replace growth media of transfected or infected cells with 25 μL of assay media (1x RPMI 1640 supplemented with 4% (vol/vol) low IgG fetal bovine sera).
- In a separate 96-well plate, perform four-fold dilutions of the antibodies of interest starting at 30 µg/mL using the assay media. Perform dilutions in triplicates and an example plate layout is illustrated in Figure 1A. Make sure to include a control that excludes antibody (no antibody control) as well as a background control (assay buffer alone).
- NOTE: For both the 'no antibody' and background controls, add 25 µL of assay buffer instead of antibody.
- 3. Transfer 25 µL of the serially diluted antibodies from Step 3.2 to corresponding wells on plate of transfected cells or infected cells.
- 4. Incubate plate in a 37 °C incubator (with 5% CO₂) for 15 min.
- 5. Add 25 µL of the appropriate modified effector Jurkat cell expressing murine FcgRIV or human FcgRIIIa (75,000 of cells/well). Total volume for each well should be 75 µL and the starting final concentration of the antibody is at 10 µg/mL (Figure 1B). NOTE: For the background control, add 25 µL of assay buffer instead of effector cells. To examine antibody-dependent cell-mediated phagocytosis, a modified Jurkat cell expressing the human FcgRIIa can be used.
- 6. Incubate plate in a 37 °C incubator with 5% CO₂ for 6 h.
- Thaw and warm luciferase substrate buffer in a 37 °C water for about 1 h before use. NOTE: To make the luciferase substrate buffer, reconstitute lyophilized luciferase assay substrate in assay buffer provided. The reconstituted substrate buffer can be stored at -30 °C to -10 °C for up to six weeks.
- 8. Add 75 µL/well of luciferase substrate buffer to all wells except for the background control.
- 9. Read on a luminometer.
- 10. Calculate fold induction with the following equation: Fold induction = (Induced value-Background value) / (No mAb value-Background value).

Representative Results

We demonstrated that a stalk-specific mAb, 6F12, but not head-specific a head-specific mAb, PY102, was able to activate primary natural killer cells by upregulating CD107a and interferon γ^{19} . To model the ability of a stalk-specific antibody to activate primary human NK cells, we show that a stalk-specific mAb, 6F12, can robustly activate the modified Jurkat cell expressing the murine FcγRIV by ~14-fold. On the other hand, the head-specific mAb, PY102 and the control IgG do not (**Figure 2**)²⁸.

To investigate if the multiplicity of infection (MOI) can affect the induction of Jurkat cells expressing the murine FcγRIV, we infected MDCK cells with X31 (a reassortant influenza A virus expressing the hemagglutinin and neuraminidase of A/Hong Kong/1/68 (H3N2) virus with the internal

segments of A/Puerto Rico/8/34 (H1N1)) with an MOI of 3, 1, or 0.33. Twenty four hours later, Jurkat cells expressing murine FcγRIV and mAb 9H10 were added unto the infected MDCK cells. In **Figure 3**, we show that cells infected with an MOI of 3 can induce the luminescence around 3-fold higher than MDCK cells infected with an MOI of 0.33.



Figure 1: A schematic of a sample plate layout and composition of the experimental groups. (A) The starting concentration for each antibody sample is at 10 μ g/mL and serially diluted across the plate (column 1 to column 12) 4-fold. Each sample is done in triplicates. Row G is reserved for the no mAb control and Row H is background. (B) Final volume for each experimental group before addition of luciferase substrate buffer is 75 μ L. Of note, the sample and 'no mAb groups' have 25 μ L of effector cells added, while the background control groups have 75 μ L of assay media. Please click here to view a larger version of this figure.



Figure 2: Induction of modified Jurkat cells expressing FcgR by a stalk monoclonal antibody. A549 cells were infected with A/Puerto Rico/8/34 (H1N1) with an MOI of 3. At 24 hours post infection, murine FcγRIV-expressing Jurkat cells in combination with PY102, 6F12 or a IgG control were added. Induction of Jurkat cells were assessed by luminescence signal generated by firefly luciferase under the NFAT response element. Error bars = standard deviations. Please click here to view a larger version of this figure.



Figure 3: The multiplicity of infection affects induction of a stalk-specific antibody. MDCK cells were infected with X31 (H3N2) with an MOI of 3, 1, or 0.33. At 24 h post infection, FcgR-expressing Jurkat cells in combination with 9H10 (starting dilution of 10 μ g/mL) added. Induction of Jurkat cells were assessed by luminescence signal generated by firefly luciferase under the NFAT response element. Error bars = standard deviations. Please click here to view a larger version of this figure.

Discussion

The method described here allows the user to measure the ability of an HA-specific monoclonal antibody to engage the murine FcyRIV. The target antigen, influenza virus hemagglutinin, is expressed on the surface of cells after infection by virus or transfection of plasmid DNA. The

assay is further amenable to using different surface-expressed viral proteins in combination with other FcqRs (humans or murine). Additionally, we have used sera samples to assess the ability of antibodies in a polyclonal context to induce Fc effector functions (data not shown), which may provide more relevant information when studying the adaptive immune response induced by infection or vaccination. We additionally speculate that the assay can be modified to measure the induction of antibodies that recognize internal viral proteins by directly coating the plates with soluble protein. However, this methodology has not been rigorously tested.

Induction of antibody-dependent effector functions can be affected by several factors including but not limited to the effector to target ratio (number of effectors cells added per well), the concentration of antibody used, the length of incubation between effector and target cells, and the concentration of low IgG serum used in the assay buffer.

Historically, the standard assay to measure the antibody-dependent cellular cytotoxicity activity of antibodies or polyclonal sera involves primary natural killer (NK) cells harvested from donors. Engagement of FcvR and activation of the intracellular pathway is typically determined by the expression of activation markers CD107a and/or IFN- $\gamma^{19,23,24,31}$. Although the classical assay uses more relevant effector cells as they have not been genetically engineered, donor variation due to allelic differences in the FcvR may affect NK cell activation and result in batch to batch variation. Alternative assays used as surrogates for ADCC activity include an FcvR dimer-based ELISA or a chromium release killing assay²⁵. Though the use of an FcvR dimer-based ELISA is high throughput, the assay may not be physiologically relevant as this assay does not use effector cells³². While the chromium release assay measures direct killing, it is not high throughput and requires primary NK cells which again are donor dependent³³. One recent method describe the use of murine T-cell hybridomas stably expressing chimeric FcvR-CD3 ζ chain molecules and measures activation through the secretion of IL-2. The assay described is similar to the described method here, which makes use of a modified T-cell as a surrogate for an effector cell expressing a singular activating FcvR³⁴. Alternatively, a cell line (NK92.05-CD16) expressing the FcvRIIII can be used as the effector cell and upregulation of the degranulation antigen CD107a is used as an activation marker. However, the use of virions and viral peptides on ELISA plates on the latter protocol should be noted as proteins coated onto hydrophobic surfaces may not always recapitulate the physiological structure of antigens³⁵.

The method described here is a 96-well format assay, makes use of physiologically relevant and natively expressed viral targets and can be modified for murine or human antibodies. The genetically modified Jurkat cells can be maintained in culture and cryopreserved³⁶. Though currently the effector cells express only one activating $Fc\gamma R$, the many advantages of using this protocol described in this study should be taken into consideration when measuring antibody-dependent cell-mediated effector function.

Disclosures

Authors declare no conflicts of interest.

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