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Development of A Direct Alhydrogel Formulation Immunoassay (DAFIA)

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Abstract

Alhydrogel® (aluminum hydroxide) is a widely used adjuvant in the US. Regulatory authorities require that vaccines be tested to determine the antigen content in the final vaccine product. The level of formulated antigen is currently determined in our laboratory by the o-Phthalaldehyde (OPA) fluorescent protein assay, and antigen identity and integrity are determined by Western blot and SDS-PAGE. However, OPA assay is non-specific and only limited to detection of total protein content, and it is often not sensitive enough to detect antigens in low dose formulations. Furthermore, antigens used in identity and integrity tests must be extracted from vaccines using an extraction procedure which is time-consuming and may not completely recover antigens for analysis or may alter the structures of antigens during extraction. The present study developed a Direct Alum Formulation Immunoassay (DAFIA) which was designed to directly (without antigen extraction), accurately, and sensitively determine the antigen content, identity and integrity on alum. The AMA1-C1/Alhydrogel formulation was used as a model vaccine in assay development and validation. The results showed that the DAFIA is highly antigen-specific, accurate $(87-100\%)$, sensitive $(0.16 \mu g/ml)$, reproducible, and simple with a linear detection range of $0.16-10 \mu g/ml$. These results demonstrate that DAFIA is an excellent assay to determine antigen content, identity and integrity of antigens bound to alum and may be used in routine vaccine quality control for testing antigens in Alhydrogel-based vaccines.

Keywords

DAFIA; Alhydrogel; vaccine; antigen content; *Plasmodium falciparum* apical membrane antigen 1

1. Introduction

Currently, the only adjuvants approved for human vaccine use in the US are aluminum containing compounds $1-4$, including aluminum hydroxide (Al(OH)₃) or Alhydrogel®, aluminum phosphate (AlPO₄), and potassium aluminum sulfate (KAl(SO_4)₂·12H₂O) or α alum^{3–5}. To ensure vaccine quality, regulatory authorities require to quantitate vaccine antigen content in the final product; for example, at least 80% for tetanus vaccine by the World Health

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Organization⁶. In particular, it is essential to determine the amount as well as the identity and integrity of the antigens bound to aluminum containing adjuvants following formulation. Alhydrogel used in our formulations has a fibrous morphology with a primary particle size of $4.5 \times 2.2 \times 10$ nm and exists as loose aggregates ranging from 1 to 10 μ m^{5, 7–10}. The presence of such aggregates in solution prevents direct quantitation of protein content in formulations using assays such as Lowry, BCA, or Bradford protein assay, not to mention that these assays are all non-specific and low in sensitivity. The o-Phthalaldehyde (OPA) fluorescent protein assay can directly and accurately determine protein content in vaccine formulations with a moderate sensitivity (20–500 µg/ml), but it only measures total proteins and lacks specificity to the protein antigens in the samples^{11–13} Furthermore, conventional ELISA is also unsuitable to directly measure protein content in formulations due to the presence of aggregates, albeit very high specificity. Presently, protein content in vaccine formulations is determined by OPA assay whose use has been limited to detection of levels of total proteins in formulations with concentrations of 20 μ g/ml or higher. Antigen identity and integrity are determined by Western blot and SDS-PAGE, but vaccine antigens used in these assays that must be extracted from Alhydrogel formulations by a laborious and time-consuming extraction procedure whose extraction efficiency and possibility to alter the structures of antigens have been in question. Taken together, the vaccine research and development fields are in need of an assay that can quickly determine the content, identity, and integrity of formulated vaccine proteins with high accuracy, sensitivity, and specificity.

The present study developed a Direct Alhydrogel Formulation Immunoassay (DAFIA) which was designed to directly (without prior antigen extraction), accurately, sensitively, and specifically determine antigen content, identity and integrity on Alhydrogel. The *Plasmodium falciparum* apical membrane antigen 1 (AMA1)-C1 (AMA1-FVO and AMA1-3D7 were mixed at a ratio of 1:1) formulated on Alhydrogel was used as a model vaccine in the assay development and validation¹⁴. The present study used 3 AMA1-C1-specific monoclonal antibodies (mAb) including mAbs 1G4, 2E3, and 1E9, recognizing the domains I/II, domain III, and an unknown epitope of the AMA-1, respectively, and a penta-His mAb which recognizes the C-terminal $His₅-tag¹⁵$. Our results indicated that this robust assay has greatly improved vaccine quality control process with high specificity, dramatically increased accuracy and sensitivity and decreased operational time needed. This assay may be widely used in vaccine research and development with a potential to monitor vaccine integrity in Alhydrogel-based formulations.

2. Material and Methods

2.1 Monoclonal antibodies

Anti-AMA1 monoclonal antibodies were generated by A&G Pharmaceutical, Inc. (Columbia, MD) with the specificity to recognize AMA1 domains I/II (1G4), domain III (2E3), or an unknown epitope of the AMA1 (1E9). Penta-His™ antibody was purchased from Qiagen (Hilden, Germany). mAb 1G4, 2E3 and 1E9 were used at 1:200 dilutions and penta-His mAb used at 1:100 dilutions in 3% skim milk (Difco™, Becton, Dickinson and Company, Franklin Lakes, NJ) in 1xTBS.

2.2 Preparation of AMA1-C1/Alhydrogel formulations

The clinical grade *P. falciparum* apical membrane antigen 1 (AMA1)-FVO lot WRAIR0932 and AMA1-3D7 lot WRAIR0941 were purified and characterized by Malaria Vaccine Development Branch (MVDB), National Institute of Allergy and Infectious Disease, National Institutes of Health, with the methods developed in $MVDB¹⁶$. Protein concentration was measured by UV spectrum at 280nm and calculated using extinction coefficient of 1.206 or 1.205 for AMA1-FVO or AMA1-3D7, respectively. Purified AMA1-FVO and AMA1-3D7

were mixed at 1:1 ratio, each concentration (0.02, 0.04, 0.08, 0.16, 0.31, 0.63, 1.25, 2.5, 5, 10, 20, or 40 μ g/ml) of the mixture (assay standards) was prepared individually in Alhydrogel $(1,600 \,\mu\text{g/ml}$ in saline, Brenntag Biosector, Denmark), aliquoted and stored at 4° C until used. The mixture was then rotated in a rotary spinner (Appropriate Technical Resources, Laurel, MD) at 16–24 rpm for 1 h at room temperature. Test samples (AMA1-C1/Alhydrogel or BSA/ Alhydrogel) at 10, 40 or 160 μ g/ml were prepared similarly. The standards or test samples were aliquoted and kept at 4°C until use. Test samples were further diluted to a final concentration of 1 μ g/ml with 1,600 μ g/ml Alhydrogel (placebo) prior to analysis by DAFIA. Samples for OPA assay were used without dilutions.

2.3 Western blot

AMA1-FVO or AMA1-3D7 (1 µg per lane) were run on pre-cast 4–20% gradient Tris-glycine SDS-PAGE gels under reducing and non-reducing conditions at a constant current of 30 mA per gel for approximately 40 min using an XCell SureLock elecrophoresis Mini-Cell apparatus (Invitrogen Corp., Carlsbad, CA), and then transferred to nitrocellulose membranes (Invitrogen, Carlsbad, CA) at 30 V constant for 60 min. After transfer, the membranes were stained with Ponceau S solution (Sigma-Aldrich Co., St. Louis, MO) to visualize the protein bands. The lanes of each membrane were then cut into individual strips, rehydrated in 1x Tris buffered saline (TBS), and blocked with blocking buffer (3% skim milk in TBS) for 30 min, followed by incubation for 1 h at room temperature with individual mAbs. After extensive washing in TBS/0.05% Tween 20 (TBS/T, Sigma-Aldrich Co.), the strips were incubated with goat anti-mouse immunoglobulin G (IgG)-alkaline phosphatase conjugate (KPL Labs, Gaithersburg, MD) for 1 h at room temperature. Following further washes, the blots were developed by incubation with 5-bromo-4-chloro-3-indolyl-phosphate/nitroblue tetrazolium (BCIP/NBT substrate, Kirkegaard and Perry Labs, Gaithersburg, MD) according to the manufacturer's instructions. The reaction was stopped when additional bands ceased to develop.

2.4 OPA assay

OPA reagent was purchased from Pierce (Rockford, IL) and the assay was performed in a 96 well black flat-bottom plate (Thermo Fisher Scientific, Waltham, MA) following manufacturer's instructions.

2.5 DAFIA Procedure

The DAFIA procedure is depicted in Fig. 1. Briefly, standards or test samples (200 µl/well) were added to black, opaque, 96-well U-bottom plates (Coring Inc. NY) and washed 3 times with phosphate-buffered saline (PBS, pH 7.4) by centrifugation at 1,000 g for 4 min at room temperature. The plates were then blocked with 200µl 3% BSA/PBS at room temperature for 1.5 h, followed by 3 washes with PBS. Mouse anti-AMA1-C1 monoclonal antibodies as primary reagent in 100 µl were added and incubated for 1 h at room temperature, followed by 3 washes with PBS. Goat anti-mouse IgG (H+L) conjugated to fluorescein (1:100, Pierce, Rockford, IL) as secondary reagent in 100 μ l was added to plates and incubated for 1 h at room temperature and washed 3 times with PBS. Following the final wash, pellets were resuspended in 100 µl PBS and read by a fluorometer (PerkinElmer, Waltham, MA) at 485nm/535 nm. Controls included Alhydrogel alone (1,600 µg/ml, placebo) with first and secondary Abs, Alhydrogel with secondary Ab alone, and AMA1-C1/Alhydrogel formulation at 1.25 µg/ml with secondary antibody alone.

2.6 Data analysis

The standard curves were generated with four-parameter nonlinear regression and the amount of standards or test samples were calculated from the equation of $Y = ((b/(a-log(X)))^A(d)-1)/$

c, where a, b, c or d are value of the parameters, Y is the amount determined and X is fluorescence readings. The back calculation was performed by converting the observed fluorescence readings at 485/535 nm of the standards to concentrations of the antigens using the four-parameter nonlinear regression equations. The percent accuracy was calculated by the following formula: percent accuracy $=$ (|calculated concentration – nominal concentration | / nominal concentration) \times 100. Inter-assay variation was calculated for the same samples (from different aliquot) that were tested in different assays and reported as coefficient of variation (CV) which is the standard deviation divided by the mean.

3. Results

3.1 Western blot analysis of AMA1-FVO and AMA1-3D7 by mAbs

The specificity of the monoclonal antibodies to AMA1-FVO and AMA1-3D7 was determined by Western blot (Fig. 2). All mAb used in this study recognized both AMA1-FVO and AMA1-3D7 under reducing and non-reducing conditions (Fig. 2).

3.2 Establishment and validation of DAFIA

Standard samples were tested between the range of 0.02 and 40 μ g/ml by DAFIA using 4 mAbs and standard curves were generated between the concentrations of 0.16 and $40 \mu g/ml$ with four-parameter nonlinear regression analysis, with a correlation coefficient (R^2) greater than 0.99 (Fig. 3). The lower doses $(0.02 - 0.08 \,\mu\text{g/ml})$ were tested, but the data points were eliminated due to poor correlations to the nominal concentrations (data not shown).

Back calculation was used to confirm the reliability of the standard curves and determine the detection range of the assay. For each data point of standard curve, the protein concentration of AMA1-C1/Alhydrogel was back calculated using the fluorescent readings and the four parameter nonlinear regression equation (Table 1). The protein concentrations determined by back calculation agreed well with the nominal amounts of AMA1-C1(with a percent accuracy of 84% or higher) between a detection range of 0.16 and 10 μ g/ml, indicating the standard curves within the detection range generated by four parameter nonlinear regression were useful to extrapolate AMA1-C1/Alhydrogel concentrations in vaccines. All anti-AMA1-C1 mAbs had a similar detection limit of 0.16 and 10 μ g/ml with some variations in the lower detection range, with an exception for penta-His mAb which had a detection limit of 0.31 and 10 μ g/ml (Fig. 2; Table 1).

Inter-assay variation analysis showed that coefficient of variation (CV) of the test samples were from 0 % to 16.0 % at 1.0 µg/ml (Table 2). In general, the acceptable CV between the assays should be less than 15%. For all 12 experiments performed, only one experiment (1E9 at 40 μ g/ml dose) gave a higher than expected CV (16%), which indicate that the DADIA was repeatable within the defined detection range.

Alhydrogel alone (1,600 µg/ml, placebo) with first and secondary Ab, placebo with secondary Ab alone or AMA1-C1/Alhydrogel formulation at 1.25 µg/ml with secondary antibody alone were used as controls. BSA/Alhydrogel was formulated at 10, 40 or 160 µg/ml and used as irrelevant antigen control for the specificity of Penta-His™ antibodies. These controls had readings similar to those of the background (data not shown).

3.3 Comparative analysis of formulation samples by DAFIA and OPA

Freshly prepared Alhydrogel formulations at 10, 40 or 160 µg/ml were analyzed by DAFIA using 4 monoclonal antibodies. All samples were pre-diluted to a final concentration of $1 \mu g$ / ml by placebo prior to analysis. The results showed that a % accuracy of 87% or higher for all samples tested was achieved, with an exception that the % accuracy (76.6 \pm 9.8) for sample

formulation at 10 µg/ml analyzed with 2E3 was slightly lower than those of other mAbs (Table 2).

AMA1-C1/Alhydrogel formulation at 40 or 160 μ g/ml doses was also tested by OPA assay to confirm the protein content. The $10 \mu g/ml$ dose was excluded due to insufficient assay sensitivity with OPA assay. The results of OPA assay were comparable to the results of DAFIA for 40 and 160 µg/ml doses (Table 2).

4. Discussion

There is no generic method available for the determination of protein content for the alumbased vaccines. Katz reported the use of ELISA to quantitate antigens formulated on Alhydrogel¹⁷. This report utilized a 2-step procedure for analysis, involving elution of antigens from Alhydrogel and followed by ELISA. However, the elution procedure was very tedious using 1.2 M potassium phosphate, and the efficacy of elution and the chance to alter the structure of antigens are in question. Our laboratory adapted OPA assay for the direct protein content determination of intact vaccines, but the sensitivity of detection of this assay is limited and unable to determine individual antigens in the multivalent vaccines due to lack of specificity.

The results from current study clearly showed that the DAFIA was highly accurate (87–100%), sensitive $(0.16 \mu g/ml$ with 3 out of 4 mAbs), and simple to perform with a detection range of $0.16-10 \mu$ g/ml. In theory, DAFIA is able to detect protein content of vaccine formulations with concentrations ranging from $0.16 \mu g/ml$ to the highest concentrations formulated, because diluting seem not to compromise the accuracy of this test as shown with formulations between 10 and 160 µg/ml. At the least, this assay should be able to overcome the disadvantage of OPA assay and accurate detection of protein content in low dose formulations (10 µg/ml or lower) in our quality control process.

In addition to its sensitivity and accuracy for protein content determination, DAFIA has several important potential applications while protein content is simultaneously measured using single or multiple specific mAbs. First of all, since this is an antibody-based assay, DAFIA can specifically determine the protein identity on Alhydrogel without protein extraction. Secondly, it is highly feasible that DAFIA may be used to determine protein integrity while the intact formulation state is maintained. Our results clearly demonstrated that the epitopes recognized by all antibodies were present on AMA1-C1/Alhydrogel, in part depicting the intact state of the vaccine. Additional work is warranted using molecularly engineered truncated vaccines and multiple specific antibodies recognizing structures critical to the immunogenicity of the vaccine of interest. Thirdly, the use of DAFIA eliminates the laborious protein extraction process, and therefore prevents protein loss due to experimental and human errors and structure alterations including loss of critical epitopes by inevitable exposure to the extraction reagents and temperature variations. Finally, DAFIA can shorten the operational time and simplify the operation procedures. Moreover, it may be reasonably to speculate that DAFIA is particularly useful in determining protein content, identity, integrity and/or structural alterations of multivalent vaccines provided that all the specific antibodies to each of the vaccine components are available.

In summary, DAFIA represents a novel assay which may have broad applications in quality control of vaccine research and development or other similar research areas.

Abbreviations

DAFIA, Direct Alhydrogel Formulation Immunoassay; OPA, o-Phthalaldehyde; AMA1, *Plasmodium falciparum* apical membrane antigen 1; AMA1-C1, AMA1-FVO and AMA1-3D7 were mixed at a ratio of 1:1; STDEV, standard deviation..

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Fig. 1.

A schematic representation of the DAFIA procedure. The assay consists of 4 steps: Step 1, addition of Alhydrogel formulation and wash (3x), followed by blocking with 3% BSA/PBS and wash $(3x)$; Step 2, addition of and incubation with primary antibody and wash $(3x)$; Step 3, addition of and incubation with secondary antibody-flourescein and wash (3x); and Step 4, determination of fluorescent intensity by a fluorometer at 485nm/535 nm.

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Fig. 2.

The specificities of monoclonal antibodies to AMA1-FVO and AMA1-3D7 were determined by Western blot. A. 1G4, B. 1E9, C. 2E3, and D. anti-his. Lane 1, AMA1-3D7 under nonreducing condition; Lane 2, AMA1-3D7 under reducing condition; Lane 3, AMA1-FVO under non-reducing condition; Lane 4, AMA1-FVO under reducing condition. Antigen was loaded at 1 µg per lane and mAbs were used at 1:500 dilution.

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AMA1-C1/Alhydrogel Concentration (µg/ml)

Fig. 3.

DAFIA standard curves for AMA1-C1/Alhydrogel formulations using 4 monoclonal antibodies specific to AMA1-C1. A. Penta-his₅ mAb; B. mAb 1G4; C. mAb 2E3; and D. mAb 1E9. Results showed the detection range of 0.16 – 10 µg/ml using all 4 monoclonal antibodies. All standard curves were fitted by 4-parameter nonlinear regression with R^2 greater than 0.99.

Back calculation and percent accuracy of standard curves *a*

back cacuuauon was periorineu by convering ine ooserven reaungs at 400min 2001 on the standards to concentrations or antigers using the rour parameter nominear regression equations, percent accuracy is percent similarity b accuracy is percent similarity between the amount of AMA1-C1 calculation (detected by DAFIA) and the known amount of AMA1-C1. Percent accuracy = (|calculated concentration ²Back calculation was performed by converting the observed readings at 485nm/535 nm of the standards to concentrations of antigens using the four parameter nonlinear regression equations; percent - nominal concentration | / nominal concentration) × 100. Results are means ± standard deviation, representing $2-4$ independent experiments. – nominal concentration| / nominal concentration) × 100. Results are means ± standard deviation, representing 2–4 independent experiments.

 b Known amount of AMA1-C1 (AMA1-FVO and AMA1-3D7 at 1:1 ratio) was fresh formulated on Alhydrogel. *b*Known amount of AMA1-C1 (AMA1-FVO and AMA1-3D7 at 1:1 ratio) was fresh formulated on Alhydrogel.

Concentrations detected by DAFIA were obtained by back calculation using the four parameter nonlinear regression equations. *c*Concentrations detected by DAFIA were obtained by back calculation using the four parameter nonlinear regression equations.

 $d_{\mbox{ND, not detectable.}}$ d _{ND}, not detectable.

*b*Known amount of AMA1-C1 (AMA1-FVO and AMA1-3D7 at 1:1 ratio) was fresh formulated on Alhydrogel.

 b Known amount of AMA1-C1 (AMA1-FVO and AMA1-3D7 at 1:1 ratio) was fresh formulated on Alhydrogel.

*c*CV, coefficient of variation between assays. CV is calculated as standard deviation divided by mean.

 $^{\circ}$ CV, coefficient of variation between assays. CV is calculated as standard deviation divided by mean.

*d*Test sample AMA1-C1-Alhydrogel formulations were prepared fresh and diluted to a final concentration of 1.0 µg/ml with placebo (1,600 µg/ml Alhydrogel).

 d_{Test} sample AMA1-C1-Alhydrogel formulations were prepared fresh and diluted to a final concentration of 1.0 µg/ml with placebo (1,600 µg/ml Alhydrogel).

ND, not detectable.

ND, not detectable.