

Aptamer-Phage Reporters for Ultrasensitive Lateral Flow Assays

Meena Adhikari,^{†,||} Ulrich Strych,^{†,||} Jinsu Kim,[‡] Heather Goux,[†] Sagar Dhamane,^{†,#} Mohan-Vivekanandan Poongavanam,^{†,||} Anna E. V. Hagström,^{‡,⊥} Katerina Kourentzi,[‡] Jacinta C. Conrad,[‡] and Richard C. Willson^{*,†,‡,§}

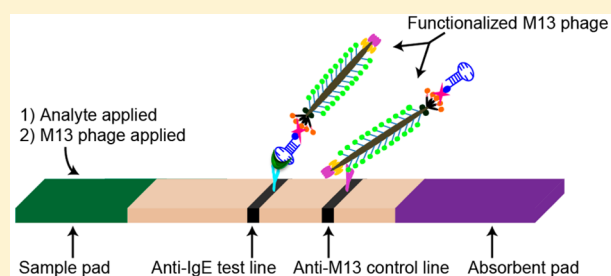
[†]Department of Biology & Biochemistry, University of Houston, Houston, Texas 77004, United States

[‡]Department of Chemical & Biomolecular Engineering, University of Houston, Houston, Texas 77204, United States

[§]Centro de Biotecnología FEMSA, Tecnológico de Monterrey, Monterrey, Nuevo León, Mexico

S Supporting Information

ABSTRACT: We introduce the modification of bacteriophage particles with aptamers for use as bioanalytical reporters, and demonstrate the use of these particles in ultrasensitive lateral flow assays. M13 phage displaying an *in vivo* biotinylatable peptide (AviTag) genetically fused to the phage tail protein pIII were used as reporter particle scaffolds, with biotinylated aptamers attached via avidin–biotin linkages, and horseradish peroxidase (HRP) reporter enzymes covalently attached to the pVIII coat protein. These modified viral nanoparticles were used in immunochromatographic sandwich assays for the direct detection of IgE and of the penicillin-binding protein from *Staphylococcus aureus* (PBP2a). We also developed an additional lateral flow assay for IgE, in which the analyte is sandwiched between immobilized anti-IgE antibodies and aptamer-bearing reporter phage modified with HRP. The limit of detection of this LFA was 0.13 ng/mL IgE, ~100 times lower than those of previously reported IgE assays.



Lateral flow assays (LFAs), in which reporter particles are transported by capillary wicking in a porous material and accumulate to form a visible line in the presence of analyte, are widely used because of their convenience, low cost, and adaptability to use at the point-of-care.^{1–3} LFAs most often employ antibodies as recognition elements, but nucleic acid recognition elements have increasingly been employed, not just in lateral flow nucleic acid hybridization assays,⁴ but also in protein detection assays where target-specific RNA or DNA aptamers serve as the molecular recognition element.^{1,5–9} Signal readout is most commonly achieved by using visible nanoparticles (gold, blue latex, or carbon). But even if the classic LFA is a brilliant approach to delivering assays in a rapid and easy-to-use format, the LFA limits of detection (LOD) lag behind more complex laboratory methods, e.g., ELISA. Thus, the development of alternative LFA readout reporters to increase assay sensitivity is a topic of great commercial and academic interest.

Bacteriophages have been explored in recent years for use as reporters in immunoassays that enable both improved recognition and enhanced signal readout. The surface of a phage, consisting of multiple copies of identical coat proteins, can be readily modified using well-established chemistries.^{10–13} M13 phage, in particular, has been modified with recognition elements (e.g., antibodies) and/or reporter elements (e.g., horseradish peroxidase (HRP), fluorophores), and has been successfully used in various protein or small molecule detection assays, e.g., enzyme-linked immunosorbent assays (ELISA),^{14–16} microarrays,¹⁷ or other protein sensors.^{18,19}

PCR amplification of the phage genetic material has led to the development of ultrasensitive immunophage PCR assays by ourselves and others.^{20–22}

We recently introduced the use of engineered bacteriophage particles as reporters in lateral flow assays.^{23–25} M13 phage particles functionalized with both target-specific antibodies and enzyme reporters were integrated into an immunochromatographic lateral flow assay (LFA) for the detection of a model virus (MS2), leading to greatly enhanced detection sensitivity (1000-fold better than a conventional gold nanoparticle LFA using the same antibodies). Our strategy often relies upon a phage-displayed “AviTag” peptide^{26–28} which can be biotinylated by biotin ligase, either *in vivo* during phage assembly in *Escherichia coli* or *in vitro* through treatment of purified phage particles with the enzyme.²⁴ This strategy also allows the convenient modification of phage reporter particles with any biotinylated molecular recognition agent, including antibodies and (as introduced here) aptamers.

Here we report the use of engineered phage particles modified with aptamers on their tail proteins and bearing reporter enzymes, in lateral flow assays with improved limits of detection, compared to a conventional gold nanoparticle assay. The conjugation of aptamers to phage constitutes a novel and promising approach to the point-of-care detection of proteins. This approach may also readily be extended to the develop-

Received: February 19, 2015

Accepted: October 11, 2015

Published: October 12, 2015

ment of higher-sensitivity lateral flow nucleic acid hybridization assays.

EXPERIMENTAL SECTION

Materials and Reagents. Polyclonal anti-IgE antibody was purchased from Fitzgerald Industries International (Acton, MA). Human immunoglobulin E (IgE) was obtained from Abcam (Cambridge, MA). An oligonucleotide bearing a 5' biotin and TEG (triethylene glycol) spacer with sequence matching that of a previously reported DNA aptamer binding human IgE²⁹ was purchased from Integrated DNA Technologies (Coralville, IA) (Figure 1). In addition, an aptamer

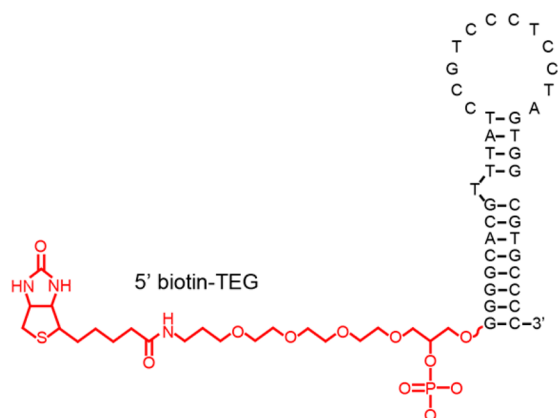


Figure 1. Biotin-TEG IgE aptamer (secondary structure adapted from ref 29).

specific for the *Staphylococcus aureus* penicillin-binding protein 2a (PBP2a) (5'-biotinylated; proprietary sequence) was purchased from Base Pair Technologies (Pearland, TX). It is a DNA aptamer that was selected from a randomized 32-mer library against MRSA (methicillin-resistant *Staphylococcus aureus*) protein. Since IgE aptamer is a DNA aptamer, a DNA aptamer (PBP2a) was used as an appropriate control for the aptamer-phage LFA. PBP2a protein was purchased from Ray Biotech (Norcross, GA).

Propagation and Modification of SAM-AviTag Phage. SAM (serine-amber)-AviTag M13 phage, a gift from Dr. Brian Kay, UIC (Chicago, IL),^{26,27,30} are derivatives of M13 phage displaying an AviTag peptide (a 15-amino-acid peptide that is a substrate for biotin ligase (BirA)) on the phage coat protein, pIII. The phage were propagated, titered, biotinylated, and conjugated with horseradish peroxidase largely as previously described.²⁴ Briefly, phage were initially grown on *E. coli* TG1 in LB medium at 37 °C, and this infected culture was then transferred to 500 mL 2xTY³¹ medium in a 2 L flask and incubated overnight at 37 °C on a shaker. After centrifugation, the phage-containing supernatant was passed through a 0.45 μ m filter, and phage were purified by precipitation with 20% polyethylene glycol (PEG) in 2.5 M NaCl. Phage titers were determined by plaque assay on X-Gal/IPTG plates.³² The pIII protein of SAM-AviTag phage was enzymatically biotinylated using *E. coli* biotin ligase (Avidity, Aurora, CO) according to the manufacturer's instructions; biotin ligase also can be prepared in-house as previously described.²⁰ The efficiency of biotinylation was determined through an ELISA on streptavidin-coated microtiter plates (StreptaWell High Bind, Roche Applied Science, Indianapolis, IN). Subsequently, phage were activated using Traut's reagent (2-iminothiolane-HCl, Thermo

Fisher Scientific, Waltham, MA). A 100 μ L portion of a 1×10^{11} phage per mL solution were suspended in 800 μ L phosphate-buffered saline (PBS), pH 7.4, 3 mM EDTA with a 20-fold molar excess of Traut's reagent. This reaction was continued for 90 min at 25 °C on a rotator. Excess Traut's reagent was removed using a 10 kDa filter (Millipore, Billerica, MA). HRP (Sigma-Aldrich, St. Louis, MO) was modified with maleimide groups using sulfo-SMCC (succinimidyl-4-(*N*-maleimidomethyl) cyclohexane-1-carboxylate, Sigma-Aldrich) by mixing HRP and sulfo-SMCC at a final concentration of 22 μ M and 1.14 mM, respectively, and incubating for 30 min. Excess sulfo-SMCC was removed using a 10 kDa filter. Traut's reagent-activated phage and maleimide-HRP were then incubated together for 90 min at 25 °C (270 000 molecules of HRP were offered per phage). Uncoupled HRP was then removed using 100 kDa filters (Millipore). Subsequently, 1 μ L of 1 mg/mL of neutravidin (Thermo Fisher Scientific) was then offered to the HRP-labeled phage and bound through the biotinylated AviTag displayed on phage protein pIII (100 molecules of Neutravidin were offered per phage). Uncoupled neutravidin was removed using 100 kDa filters (Millipore).

Aptamer Modification of Avidin Phage. Biotinylated IgE aptamer (100 μ L, 100 nM in PBS, pH 7.0, 1 mM MgCl₂) was first heat-denatured at 95 °C for 10 min (to disrupt any pre-existing higher-order structures that might interfere with phage functionalization and to allow the 5'-biotin to interact freely with its avidin binding partner³³) and then slowly renatured at room temperature. Subsequently, the biotinylated IgE aptamer was added to 10^{12} neutravidin-functionalized phage in 100 μ L PBS. The complex was incubated for 2 h at 25 °C before unconjugated IgE aptamer was removed using PEG precipitation (Figure 2). As a control, phage were function-

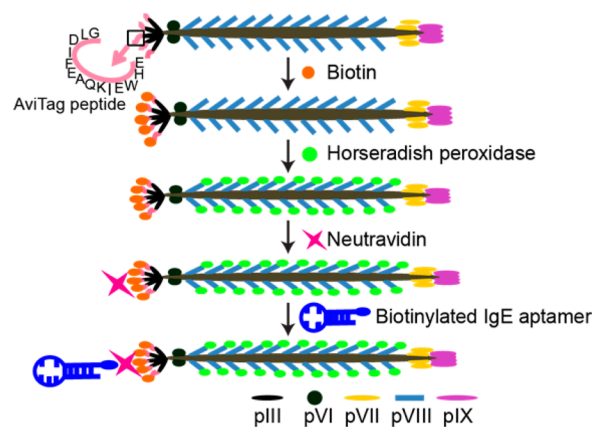


Figure 2. Functionalization of M13 phage with IgE aptamers and horseradish peroxidase (HRP). Phage displaying AviTag peptides are biotinylated using biotin ligase. Biotinylated phage are covalently modified with HRP on the major coat protein pVIII before neutravidin is bound to biotinylated AviTag on HRP-labeled phage. Neutravidin-functionalized phage are then conjugated with the biotinylated IgE aptamer.

alized with the anti-PBP2a aptamer in the same fashion. A series of 10-fold dilutions (10^5 to 10^{10} phage/mL) of unmodified phage (titer obtained by plaque counting) was used as PCR calibration standards, and the concentration of the various phage constructs was determined against that standard curve. The AviTag-targeted PCR primers were the following: forward 5'-GTTGTTTCTTTCTATTCTCACTCC-3' and

reverse 5'-CAGACGTTAGTAAATGAATTTCTG-3'. For the phage amplification, 0.1 μL of a 10 μM stock of each primer was combined with 10 μL of 2xPCR mix (Brilliant III Ultra-Fast SYBR mix, Agilent, Santa Clara, CA), 4.8 μL of ultrapure (RNase and DNase free) water, and 5 μL of each phage sample to achieve a total volume of 20 μL . The PCR conditions were the following: 10 min at 95 $^{\circ}\text{C}$, 40 cycles of 30 s at 95 $^{\circ}\text{C}$, 30 s at 62 $^{\circ}\text{C}$, and 30 s at 72 $^{\circ}\text{C}$, followed by a dissociation step to detect possible contaminations (1 min at 95 $^{\circ}\text{C}$, 30 s at 55 $^{\circ}\text{C}$, and 30 s at 95 $^{\circ}\text{C}$).

Preparation of LFA Strips. Figure 3 depicts the schematic of the aptamer-phage LFA. Polyclonal anti-human IgE rabbit

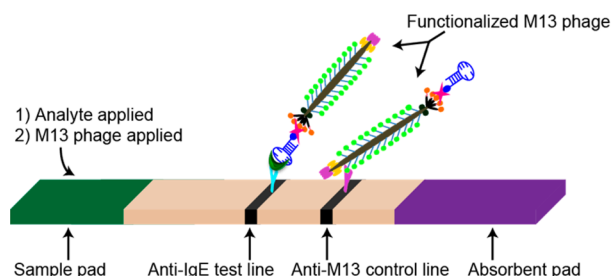


Figure 3. Aptamer-phage lateral flow assay. The M13 phage SAM-AviTag protein, pIII, was specifically biotinylated and functionalized with analyte-specific biotinylated aptamers, and the M13 coat proteins were functionalized with horseradish peroxidase. The detection line contains IgE-specific antibodies, and the control line has anti-M13 antibodies. Captured HRP-labeled phage were detected on the test and control lines using the chromogenic substrate, 3',3',5,5'-tetramethylbenzidine (TMB).

serum (Fitzgerald Industries International, Acton, MA; 5 μL , diluted 100-fold in 50 mM sodium acetate buffer (pH 3.6) based on preliminary screening experiments) and 5 μL of rabbit polyclonal anti-M13 IgG antibodies (Novus Biologicals, Littleton, CO; 0.1 $\mu\text{g}/\mu\text{L}$ in 50 mM sodium acetate buffer (pH 3.6)) were used to make test and control lines, respectively, on Fusion 5 membranes (GE Healthcare, Piscataway, NJ) using a lateral flow reagent dispenser (Claremont BioSolutions, Upland, CA) equipped with an external syringe pump (Chemxy, Stafford, TX). The Fusion 5 membrane was cut into 7 mm \times 50 mm strips using a guillotine paper cutter. CF5 membrane (GE Healthcare) was used as the absorbent pad (7 mm \times 25 mm), and Fusion 5 membrane (7 mm \times 10 mm) was used as a sample pad. The strips were allowed to dry for 1 h at 25 $^{\circ}\text{C}$ before use.

Lateral Flow Assay with Aptamer-Phage. A 10-fold dilution series of human IgE was prepared in PBS, pH 7.0 with 0.5% bovine serum albumin (BSA). Each IgE sample (100 μL) was dispensed onto the sample application area of a prepared LFA strip. The strips were then washed with 200 μL of LFA buffer [(50- μL aliquots; 4 \times) PBS, pH 7.0; 1% Tween-20; 1% Triton; 0.1% PEG-3350] followed by 10 μL of the aptamer-phage reporter construct ($\sim 10^9$ phage), and finally with 500 μL LFA buffer (50- μL aliquots; 10 \times). Signals were visualized by spotting 25 μL of TMB liquid substrate system for membranes (Sigma-Aldrich, St. Louis, MO) on each line. The signals were allowed to develop for 10 min, and then the strips were scanned on a Perfection V600 flatbed color scanner (Epson, Long Beach, CA). The scanned images were analyzed using ImageJ's Gel Analysis Tool and by plotting the line intensity profile. The intensity of each line was given by the area under

each peak that was numerically integrated using ImageJ's Gel Analysis Toolbox. The ratio of the intensity of the test line to that of the control line for each strip (T/C) was used as an indicator of signal strength over the background.

RESULTS AND DISCUSSION

Evaluation of Aptamer-Phage for Direct Detection of Spotted IgE and PBP2a in LFA. Initial LFA experiments tested the binding of the anti-IgE and anti-PBP2a aptamer-phage directly to IgE and PBP2a proteins, respectively, spotted on a Fusion 5 membrane. For the IgE experiments, 5 μL of IgE (0.2 mg/mL) and 5 μL of anti-M13 antibodies (0.1 mg/mL) in 50 mM sodium acetate buffer (pH 3.6) were spotted on the test and control lines, respectively, of a Fusion 5 membrane. For the anti-PBP2 aptamer-phage, 5 μL PBP2a protein (0.2 mg/mL) was spotted on the test line and anti-M13 antibodies on the control line. The strips were dried for 1 h at room temperature. Aptamer-phage dilutions (10 μL) ranging from 10^7 to 10^9 phage were dispensed onto the sample application area of the LFA strip. The strips were washed with 500 μL LFA buffer, and signals were obtained by spotting TMB liquid substrate on each line. As shown in Figure 4, a clear signal was obtained at the

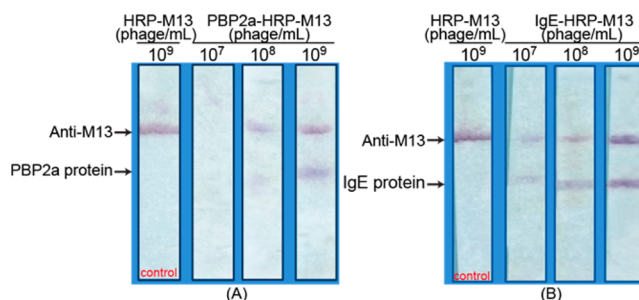


Figure 4. Direct detection of spotted target analytes [PBP2a (A) or IgE (B)] with aptamer-phage. (A) PBP2a protein and anti-M13 antibodies were spotted on the test and control lines, respectively. (B) IgE protein and anti-M13 antibodies were spotted on the test and control lines, respectively. Different concentrations of aptamer-phage constructs were then passed through the membranes. Phage bearing HRP reporters but no aptamers (HRP-M13) served as a specificity control.

anti-M13 control line for all samples containing $>10^7$ aptamer-phage. This result confirms that the aptamer-phage moved successfully through the membrane. The signals obtained at the test lines using 10^8 and 10^7 aptamer-phage were weak, but with 10^9 phage the signals for both IgE and PBP2a were clear and distinguishable from other aptamer-phage concentrations (Figure 4). Tests with phage modified with HRP but no aptamer gave no visible test line signal with either protein, thus confirming that M13 phage without aptamers are not retained nonspecifically by IgE or PBP2a protein at the test line but still yield a specific signal at the anti-M13 control line.

Lateral Flow Sandwich Immuno-Aptamer Assay for IgE Detection. We next developed a sandwich IgE LFA based on the IgE-dependent capture of phage modified with anti-IgE aptamers and HRP on anti-IgE polyclonal antibodies at a test line on Fusion 5 strips. Serial dilutions of the IgE test analyte ranging from 0.013 to 130 ng/mL were prepared in PBS at pH 7.0 with 0.5% BSA. Each IgE sample (100 μL) was dispensed onto the sample application area of the prepared LFA strips. A clear signal on the anti-M13 antibody control line for all samples confirmed that the phage reporters had successfully

moved through the membrane. A signal clearly distinguishable from the no-IgE control was obtained at the test line of the strip with 0.13 ng/mL IgE (0.68 pM) (Figure 5, Table S1). The

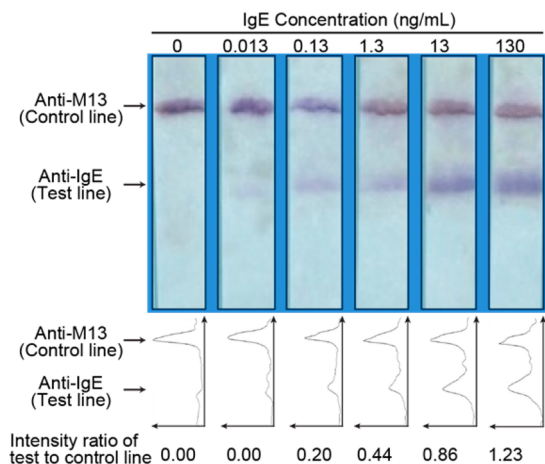


Figure 5. Lateral flow assay with aptamer-phage for IgE detection. Various concentrations of IgE (volume: 10 μ L) were detected using anti-IgE antibodies at the test line and IgE aptamer-HRP-phage reporters. Control line consisted of anti-M13 antibodies. Line intensity profiles as evaluated by ImageJ density analysis of the LFA strips are shown below each strip. The area under each peak was numerically integrated using the ImageJ Gel Analysis Toolbox to give the intensity for that line. The intensity of the test line divided by the intensity of the control line for each strip is shown below each line intensity profile.

relative intensities of the test lines were divided by the control line intensities (T/C) to adjust for any background, and this ratio was used as the reported LFA signal. The limit of detection (LOD) was determined using Student's t test from five independent replicate LFAs. We report the LOD for IgE detection using aptamer-phage to be at least 0.13 ng/mL, defined as the lowest tested concentration that was significantly different from the negative control with no IgE at a significance level of $\alpha = 0.05$. Table S2 in the Supporting Information shows that the average T/C differs between the negative and the positive control at a significant level of $\alpha = 0.05$, and we thus reject the null hypothesis for all the IgE concentrations except for 0.013 ng/mL. Therefore, the LOD of our assay is concluded to be at or lower than 0.13 ng/mL, in that the positive control signal from 0.13 ng/mL of the analyte concentration is statistically distinguishable from the negative control.

Compared to previously reported assays using the same anti-IgE aptamer, our phage-based LFA achieves a greater sensitivity in detecting IgE with a reduction in the LOD by ~ 230 times compared to the assay reported by Tran et al.³⁴ (30 ng/mL), who used an aptamer-based impedimetric biosensor using a nanocrystalline diamond (NCD) film as a working electrode to detect IgE and by ~ 8 times compared to the assay reported by Wei et al.³⁵ (10 ng/mL) who used silver nanoparticles modified with the IgE aptamers to detect captured IgE on microarrays with metal-enhanced fluorescence. It is noteworthy that Tran et al.³⁴ reported similar LODs for the detection of IgE in buffer and in human serum using this aptamer. Initial testing with human IgE spiked into human serum confirms that these samples are compatible with phage transport through the LFA strip, preservation of LOD, and specific capture of the phage constructs on the test line (preliminary data not shown).

Currently, the most popular FDA-cleared assay for IgE detection in serum is the ELISA-based ImmunoCAP system, which requires considerable investment in instrumentation and a moderately high cost per assay with a detection range for total IgE 0–100 kU/I (0–240 ng/mL; 1 kU/I = 2.4 ng/mL). ImmunoCAP Rapid is a related commercially available qualitative lateral flow test that measures specific IgE in whole blood, and its sensitivity ranges between 0.72 and 1.68 ng/mL. The ALFA (allergy lateral flow assay) Total IgE test from Dr. Fooke Laboratories (GmbH) reports a detection sensitivity of 12 ng/mL in serum or whole blood.

Aptamer-Phage Specificity. The specificity of the aptamer-phage LFA was tested by spotting the unrelated murine IgG1 anti-lysozyme antibody, HyHEL-5,³⁶ onto the test line. Neither the no-IgE samples, nor 13 ng/mL IgE, gave any signals at the HyHEL-5 test line, implying that neither IgE nor aptamer-phage bind nonspecifically to this unrelated antibody (Figure 6, strips 1 and 2). Furthermore, no signal was observed when HyHEL-5 was passed as a mock-analyte on an LFA membrane with an anti-IgE test line followed by anti-IgE aptamer-phage (Figure 6, strip 7). The strips where IgE was added confirmed that IgE binds specifically to the anti-IgE test line (Figure 6, strips 3, 4, and 5) and aptamer-phage are

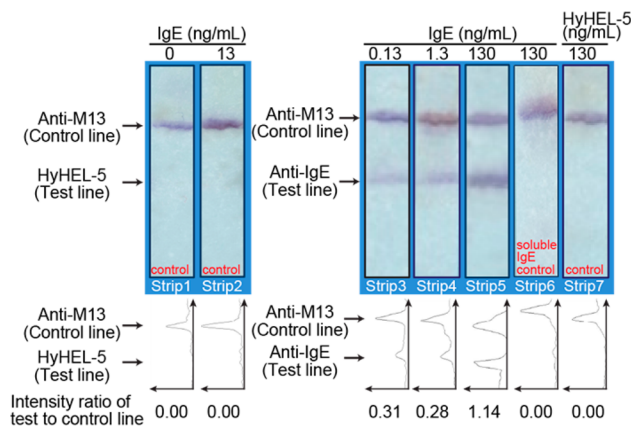


Figure 6. Control for nonspecific binding of IgE and aptamer-phage to an unrelated protein (the murine antilysozyme IgG antibody HyHEL-5). Strips 1 and 2: HyHEL-5 was spotted on the test line and anti-M13 antibodies on the control line. IgE protein (100 μ L in LFA buffer) was passed through the membrane and aptamer-phage were added as reporters. No signal was observed on the HyHEL-5 test line. Strips 3, 4, and 5: To confirm specificity and sensitivity of the assay, varied concentrations of IgE were passed through the membrane. Strip 6: Competition assay where free IgE aptamer was passed on the strip prior to offering the aptamer-phage construct; 130 ng/mL IgE in 100 μ L of LFA buffer were passed, followed by soluble IgE aptamer (10 μ M aptamer in 100 μ L of LFA buffer), and then the aptamer-phage construct was offered. No signal was observed on the test line confirming that soluble anti-IgE aptamer competes with aptamer-phage for the available binding sites on the IgE protein. Strip 7: 100 μ L of HyHEL-5 in LFA buffer was passed through the membrane with anti-IgE and anti-M13 lines and detected using the IgE aptamer-phage reporter. No signal was observed on the HyHEL-5 test line, confirming specificity of the assay. Line intensity profiles as evaluated by ImageJ density analysis of the LFA strips. The area under each peak was numerically integrated using the ImageJ Gel Analysis Toolbox to give the intensity for that line. The intensity of the test line divided by the intensity of the control line for each strip is shown below each line intensity profile.

subsequently retained, hence demonstrating the specific performance of the aptamer-phage LFA for IgE detection.

The specificity of the IgE LFA was also demonstrated by a competition assay (Figure 6, strip 6). Here, IgE (130 ng/mL) was passed through the membrane (with anti-IgE test line and anti-M13 control line) followed by an excess amount of soluble anti-IgE aptamer (10 μ M aptamer in 100 μ L of LFA buffer) to saturate the binding sites on the captured IgE protein and subsequently adding the anti-IgE aptamer-phage reporters. We previously confirmed that IgE was captured in the anti-IgE test line, as shown in Figure 6, strips 3, 4, and 6. However, the excess of competing soluble IgE aptamer occupies the sites on the captured IgE, so anti-IgE phage cannot bind, and hence no signal was observed, indicating that phage do not bind nonspecifically by interactions other than those mediated by aptamer-IgE recognition.

PBP2a Aptamer-Phage as Specificity Control for IgE-Aptamer-Phage Reporters. As a control for the specificity of the IgE aptamer-phage construct, we used an aptamer-phage functionalized with a proprietary aptamer recognizing the *Staphylococcus aureus* penicillin-binding protein, PBP2a. This PBP2a-M13 phage construct was prepared in a fashion analogous to that used for the anti-IgE aptamer-phage construct.

IgE (13 ng in 100 μ L of LFA buffer) was passed through the strip, followed by the PBP2a-M13 phage construct. As expected, the anti-PBP2a aptamer on the phage did not recognize the IgE protein captured on the test line, and hence there was no signal (Figure 7, strip 6). To confirm the performance of the PBP2a-M13 phage construct, strips were made with PBP2a protein and anti-M13 antibodies on the test and control lines, respectively (Figure 7, strip 2). A clear signal was obtained on the strip where the PBP2a-M13 phage construct was applied, indicating detection of the target PBP2a protein. As a further control, the strip with HRP-phage reporters bearing no PBP2a aptamers did not show any signal, indicating that phage do not bind nonspecifically to the PBP2a protein (Figure 7, strip 1). The strips where IgE was added reveal that IgE binds specifically to the anti-IgE test line (Figure 7, strips 4 and 5) and functionalized aptamer-phage are subsequently retained there, hence demonstrating the efficient performance of the aptamer-phage LFA for IgE detection. The test and control lines were quantified by ImageJ density's analysis function.

CONCLUSIONS

We have demonstrated a novel, specific, and highly sensitive lateral flow assay for the detection of proteins using viral nanoparticles as scaffolds bearing aptamers as biorecognition elements. With further refinement, the use of aptamer-phage as reporters in lateral flow assays promises a convenient, cost-effective, and specific framework for the detection of analytes. In a comparison to antibodies, aptamers have lower production cost, and can easily and rapidly be selected and synthesized. They are chemically more stable, and can be selected to have high affinity toward their targets. The 2700 copies of the M13 coat protein provide multiple sites for enzyme conjugation, and thus an enhanced signal per individual affinity agent is generated, resulting in increased sensitivities. Phage nanoparticles represent versatile scaffolds with high colloidal stability and very low nonspecific binding that can accommodate a variety of recognition and reporter elements, thus making them broadly applicable and ultrasensitive biodetection reporters.

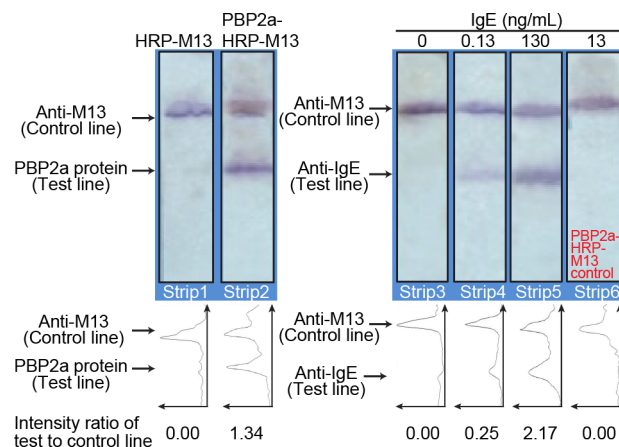


Figure 7. PBP2a aptamer-phage as a specificity control for IgE-aptamer-phage reporters. Strips 1 and 2: Strips with PBP2a protein and anti-M13 antibodies on the test and control lines, respectively, were made to evaluate the binding of PBP2a-M13 to PBP2a protein. A clear signal on the test line of strip 2 indicates binding of the PBP2a-phage to the PBP2a protein. Strip 1 with HRP-phage devoid of any signal confirms that phage do not bind nonspecifically to the PBP2a protein. Strips 3, 4, and 5 show the various concentrations of IgE protein (100 μ L in LFA buffer) that were used. IgE binds on the anti-IgE test line. Signals on the control lines indicate the proper functioning of the assay. Strip 6: A PBP2a-M13 phage construct was passed through a strip with anti-IgE and anti-M13 lines. The anti-PBP2a aptamer on the phage did not bind to the IgE protein on the test line as indicated by the absence of any signal on the test line. Line intensity profiles as plotted by ImageJ density analyses of the LFA strips are shown below each figure. The area under each peak was numerically integrated using the ImageJ Gel Analysis Toolbox, and the average intensity of the test line divided by the intensity of the control line for each strip is also shown below each line intensity profile.

While the aim of the present study was primarily to demonstrate the feasibility and performance of aptamer-phage LFAs, the success of this approach suggests that phage bearing nucleic acids also could serve as the basis of high-sensitivity lateral-flow nucleic acid assays. It is also worth noting that the detection of IgE, used as a model here, has clinical value in itself. IgE is known to play an important role in the allergic response and is widely reported as a marker of atopic diseases such as asthma, dermatitis, and pollenosis.³⁷ Testing of specific IgE, particularly at low levels of IgE, can be helpful in evaluating a patient's sensitivity profile and risk factors for other severe conditions such as anaphylaxis.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.analchem.5b00702.

Details on test line intensity profiles and Student *t* tests (PDF)

AUTHOR INFORMATION

Corresponding Author

*Phone: +1 713-743-4308. Fax: +1 713-743-4323. E-mail: willson@uh.edu.

Present Addresses

[†]Department of Pediatrics, Baylor College of Medicine, Houston, Texas 77030, United States.

[†]MedImmune, Purification Process Sciences, Gaithersburg, Maryland 20878, United States.

[#]Patheon, Downstream Process Development, St. Louis, Missouri 63134, United States.

Notes

The contents of the paper are solely the responsibility of the authors and do not necessarily represent the official views of the funding agencies.

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

This work was supported in part by the Welch Foundation [E-1264], NIH [U54 AI057156 and 1R21AI111120-01A1], and National Science Foundation [CBET-1511789]. Postdoctoral scholarships for AEV Hagström from the Olle Engkvist Byggmästare Foundation and Carl Trygger Foundation are gratefully acknowledged. We thank Bill Jackson for helpful discussions.

REFERENCES

- (1) Gubala, V.; Harris, L. F.; Ricco, A. J.; Tan, M. X.; Williams, D. E. *Anal. Chem.* **2012**, *84*, 487–515.
- (2) Ngom, B.; Guo, Y.; Wang, X.; Bi, D. *Anal. Bioanal. Chem.* **2010**, *397*, 1113–1135.
- (3) Anfossi, L.; Baggiani, C.; Giovannoli, C.; D'Arco, G.; Giraudi, G. *Anal. Bioanal. Chem.* **2013**, *405*, 467–480.
- (4) Carter, D. J.; Cary, R. B. *Nucleic Acids Res.* **2007**, *35*, e74.
- (5) Liu, G.; Mao, X.; Phillips, J. A.; Xu, H.; Tan, W.; Zeng, L. *Anal. Chem.* **2009**, *81*, 10013–10018.
- (6) Fang, Z.; Wu, W.; Lu, X.; Zeng, L. *Biosens. Bioelectron.* **2014**, *56*, 192–197.
- (7) Wong, R. C.; Tse, H. Y. *Lateral Flow Immunoassay*; Springer: New York, 2009.
- (8) Bruno, J. G. *Pathogens* **2014**, *3*, 341–355.
- (9) Chen, A.; Yang, S. *Biosens. Bioelectron.* **2015**, *71*, 230–242.
- (10) Li, K.; Chen, Y.; Li, S.; Nguyen, H. G.; Niu, Z.; You, S.; Mello, C. M.; Lu, X.; Wang, Q. *Bioconjugate Chem.* **2010**, *21*, 1369–1377.
- (11) Sapsford, K. E.; Soto, C. M.; Blum, A. S.; Chatterji, A.; Lin, T.; Johnson, J. E.; Ligler, F. S.; Ratna, B. R. *Biosens. Bioelectron.* **2006**, *21*, 1668–1673.
- (12) Strable, E.; Finn, M. G. *Curr. Top. Microbiol. Immunol.* **2009**, *327*, 1–21.
- (13) Soto, C. M.; Ratna, B. R. *Curr. Opin. Biotechnol.* **2010**, *21*, 426–438.
- (14) Kim, H. J.; Ahn, K. C.; Gonzalez-Techera, A.; Gonzalez-Sapienza, G. G.; Gee, S. J.; Hammock, B. D. *Anal. Biochem.* **2009**, *386*, 45–52.
- (15) Kim, H. J.; Rossotti, M. A.; Ahn, K. C.; Gonzalez-Sapienza, G. G.; Gee, S. J.; Musker, R.; Hammock, B. D. *Anal. Biochem.* **2010**, *401*, 38–46.
- (16) Brasino, M.; Lee, J. H.; Cha, J. N. *Anal. Biochem.* **2015**, *470*, 7–13.
- (17) Domaille, D. W.; Lee, J. H.; Cha, J. N. *Chem. Commun. (Cambridge, U. K.)* **2013**, *49*, 1759–1761.
- (18) Park, J. S.; Cho, M. K.; Lee, E. J.; Ahn, K. Y.; Lee, K. E.; Jung, J. H.; Cho, Y.; Han, S. S.; Kim, Y. K.; Lee, J. *Nat. Nanotechnol.* **2009**, *4*, 259–264.
- (19) Lee, J.-W.; Song, J.; Hwang, M. P.; Lee, K. H. *Int. J. Nanomed.* **2013**, *8*, 3917–3925.
- (20) Litvinov, J.; Hagstrom, A. E.; Lopez, Y.; Adhikari, M.; Kourentzi, K.; Strych, U.; Monzon, F. A.; Foster, W.; Cagle, P. T.; Willson, R. C. *Biotechnol. Lett.* **2014**, *36*, 1863–1868.
- (21) Kim, H. J.; McCoy, M.; Gee, S. J.; Gonzalez-Sapienza, G. G.; Hammock, B. D. *Anal. Chem.* **2011**, *83*, 246–253.
- (22) Zhang, H.; Xu, Y.; Huang, Q.; Yi, C.; Xiao, T.; Li, Q. *Chem. Commun. (Cambridge, U. K.)* **2013**, *49*, 3778–3780.
- (23) Kim, J.; Adhikari, M.; Dhamane, S.; Hagstrom, A. E.; Kourentzi, K.; Strych, U.; Willson, R. C.; Conrad, J. C. *ACS Appl. Mater. Interfaces* **2015**, *7*, 2891–2898.
- (24) Adhikari, M.; Dhamane, S.; Hagstrom, A. E.; Garvey, G.; Chen, W. H.; Kourentzi, K.; Strych, U.; Willson, R. C. *Analyst* **2013**, *138*, 5584–5587.
- (25) Hagstrom, A. E.; Garvey, G.; Paterson, A. S.; Dhamane, S.; Adhikari, M.; Estes, M. K.; Strych, U.; Kourentzi, K.; Atmar, R. L.; Willson, R. C. *PLoS One* **2015**, *10*, e0126571.
- (26) Scholle, M. D.; Collart, F. R.; Kay, B. K. *Protein Expression Purif.* **2004**, *37*, 243–252.
- (27) Scholle, M. D.; Kriplani, U.; Pabon, A.; Sishtla, K.; Glucksman, M. J.; Kay, B. K. *ChemBioChem* **2006**, *7*, 834–838.
- (28) Kehoe, J. W.; Kay, B. K. *Chem. Rev.* **2005**, *105*, 4056–4072.
- (29) Wiegand, T. W.; Williams, P. B.; Dreskin, S. C.; Jouvin, M. H.; Kinet, J. P.; Tasset, D. *J. Immunol.* **1996**, *157*, 221–230.
- (30) Scholle, M. D.; Kehoe, J. W.; Kay, B. K. *Comb. Chem. High Throughput Screening* **2005**, *8*, 545–551.
- (31) Maniatis, T.; Fritsch, E. F.; Sambrook, J. *Molecular Cloning: A Laboratory Manual*; Cold Spring Harbor Laboratory: New York, 1982.
- (32) Lee, C. M.; Iorno, N.; Sierro, F.; Christ, D. *Nat. Protoc.* **2007**, *2*, 3001–3008.
- (33) Tennico, Y. H.; Hutanu, D.; Koesdjojo, M. T.; Bartel, C. M.; Remcho, V. T. *Anal. Chem.* **2010**, *82*, 5591–5597.
- (34) Tran, D. T.; Vermeeren, V.; Grieten, L.; Wenmackers, S.; Wagner, P.; Pollet, J.; Janssen, K. P.; Michiels, L.; Lammertyn, J. *Biosens. Bioelectron.* **2011**, *26*, 2987–2993.
- (35) Wei, X.; Li, H.; Li, Z.; Vuki, M.; Fan, Y.; Zhong, W.; Xu, D. *Anal. Bioanal. Chem.* **2012**, *402*, 1057–1063.
- (36) Xavier, K. A.; Willson, R. C. *Biophys. J.* **1998**, *74*, 2036–2045.
- (37) Platts-Mills, T. A. *Am. J. Respir. Crit. Care Med.* **2001**, *164*, S1–S5.