Sequence-specific Nucleic Acid Detection from Binary Pore Conductance Measurement

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1 Supporting Information

ABSTRACT: We describe a platform for sequence-specific nucleic acid (NA) detection utilizing a micropipet tapered to a 2 μm diameter pore and 3 μm diameter polystyrene beads to which uncharged peptide nucleic acid (PNA) probe molecules have been conjugated. As the target NAs hybridize to the complementary PNA-beads, the beads acquire negative charge and become electrophoretically mobile. An applied electric field guides these NA-PNA-beads toward the pipet tip, which they obstruct, leading to an indefinite, electrically detectable, partial blockade of the pore. In the presence of noncomplementary NA, even to the level of single base mismatch, permanent pore blockade is not seen. We show application of this platform to detection of the anthrax lethal factor sequence.

1. INTRODUCTION

Detection of specific DNA or RNA sequences is highly desired in a range of diverse applications such as screening for genetic diseases,†1–3 pathogenic microbe identification for food safety,‡4–6 and paternity tests in forensic science.‡7,8 For many, if not most, of these applications, the paramount need is for a yes or no answer (binary response) regarding the presence or absence of the nucleic acid (NA) sequence of interest in a sample rather than a quantitative determination of concentration. The majority of conventional sequence-specific DNA or RNA detection platforms also rely on base pairing interaction between two single strands of complementary nucleic acids that can be detected by optical, mechanical, or electrochemical readout.†9 Many of these platforms require amplification by polymerase chain reaction, fluorescent or enzymatic labels, and expensive instrumentation.‡10,11

Recently, biological or solid-state nanopores have been used as sequence specific NA detection platforms.‡12–15 Nanopores are attractive for label-free molecular sensing, because the presence of single objects within the pore, down to molecular in size, can result in large modulations of pore electrical conductance. In most of this work, these conductance modulations are transient and rapid, occurring as the object passes through the pore, and therefore a fast and sensitive amplifier is required.

We have developed a binary, sequence-specific NA detection concept based on permanent attenuation of nanopore conductance by electrophoretically mobile beads. This approach provides a yes or no signal as to the presence of a target NA in a sample, but in its current embodiment, it does not give concentration data. Sequence specificity is achieved using peptide nucleic acid (PNA) probe molecules conjugated to the beads.‡16 PNA has been used previously for sequence-specific DNA detection in a number of platforms,‡17–27 including electrochemical detection of DNA hybridization with PNA-modified magnetic beads,28 and detection of DNA hybridization to PNA bound to the walls of a nanopore sensor.‡29,30 PNA offers many advantages over DNA probes including resistance to enzyme-catalyzed degradation, increased stability of complementary DNA complexes, and increased sensitivity of these complexes to single base pair mismatches.†31 Here we take particular advantage of the electrical neutrality of PNA to ensure that the beads are uncharged and electrophoretically immobile until target NA is complexed. Hybridization of complementary target NA to the PNA-beads imparts sufficient negative charge to the beads, enabling them to be driven electrophoretically to the pore. The beads then physically occlude the pore, leading to a large decrease in conductance. The coupling of target NA to the PNA-bead conjugate thereby leads to an electromechanical amplification phenomenon that gives rise to a large, easily detected, binary modulation of electrical current. The “pores” used in the proof-of-concept work described here are the capillary openings formed at prepulled glass micropipet tips. Similar glass pores have been used to detect single DNA molecules attached to a 10 nm gold nanoparticle32 and the folding state of double-stranded DNA,33 in addition to their use for electrochemical readout of NA binding events.34,35 The label-free, potentially low cost sensor with a simple binary read-out described here may have significant technological appeal.

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Previous studies of hybridized DNA-PNA 12-mers found melting after incubation, the beads were washed with 0.4× SSC buffer (60 mM NaCl, 6 mM trisodium citrate, 0.1% Triton X-100, pH 8), resuspended into 0.6 mL of coupling buffer (100 mM1-[3-(dimethylamine) propyl]-3-ethylcarbodiimide (EDC) in MES buffer). Ten nanomoles of amine-functionalized PNA probes were added to the coupling buffer with beads and incubated at 50 °C for one hour to cap any remaining unreacted carboxylic groups. After the last coupling step, the beads were washed 4 times in 0.4× SSC buffer (60 mM NaCl, 6 mM trisodium citrate, 0.1% Triton X-100, pH 8), resuspended into 0.6 mL coupling buffer (100 mM1-[3-(dimethylamine) propyl]-3-ethylcarbodiimide (EDC) in MES buffer). Ten nanomoles of amine-functionalized PNA probes were added to the coupling buffer with beads and incubated at 50 °C for one hour to cap any remaining unreacted carboxylic groups. After the last coupling step, the beads were washed 4 times in 0.4× SSC buffer (60 mM NaCl, 6 mM trisodium citrate, 0.1% Triton X-100, pH 8), resuspended into 0.6 mL of coupling buffer containing 100 mM ethanethiol, and incubated at 50 °C for one hour to cap any remaining unreacted carboxylic groups with ethanolamine. After the last coupling step, the beads were washed 4 times in 0.4× SSC buffer and were stored in PBS buffer at 4 °C.

2.3. Hybridization Assay. Prior to DNA incubation, PNA-beads were washed twice in 0.4× SSC buffer and once in hybridization buffer (750 mM NaCl, 10 mM Tris-HCl, pH 7.0), and were resuspended in 100 μL of hybridization buffer. The PNA-beads were divided between two separate 1.5 mL centrifuge tubes. To one tube was added 1 nmol of 20-mer target DNA with sequence complementary to the PNA on the beads, and to the other (control) was added 1 nmol of noncomplementary 20-mer poltDT DNA. The tubes were placed on a mechanical rotator and incubated overnight at room temperature. After incubation, the beads were washed with 0.4× SSC buffer 3 times. Previous studies of hybridized DNA-PNA 12-mers found melting temperatures of >62 °C and reductions of 15–20 °C for DNA with single base mismatch, suggesting that the hybridized PNA-DNA used here is stable at the room temperature measurement conditions.

2.4. Zeta Potential, Electrophoretic Mobility, and Size Measurements. Following resuspension of the beads in 1 mM KCl, pH 7.0 at 25 °C, a Zetasizer Nano-ZS (Malvern Instruments) was used to characterize their zeta potential and electrophoretic mobility as well as their size using dynamic light scattering. The mean and mode diameters measured for the carboxylic acid beads were 3680 and 3580 nm, respectively, and the mean and mode diameters of the amine beads were 3250 and 3090 nm, respectively (Figure S1, Supporting Information).

2.5. Sensor Apparatus and Electrical Measurements. Two identical chambers made of polydimethylsiloxane (PDMS) connected by a 1 mm diameter opening were sealed to a glass microscope slide following activation with oxygen plasma. A prepulled borosilicate micropipet with outer diameter of 1 mm and nominal inner tip (pore) diameter of 2 μm was placed in the opening between the two chambers and sealed with vacuum grease so that the micropipet is the only connection between the two chambers (Figure 1). The platform was mounted on an inverted optical microscope (Leica DMIRB).

The chambers were filled with identical volumes of buffer (1 mM KCl, 10 mM HEPES, pH 7.0); Pt electrodes were placed in each chamber, away from the pipet entrances. A potential difference of 25 V was applied between the electrodes, and the resultant current was amplified by a transimpedance amplifier and logged using acquisition hardware at 1 kHz (PCI 6052E, National Instruments) and LabVIEW software (National Instruments). Data shown in the figures was processed in MATLAB with a fifth order Butterworth 100 Hz lowpass filter. After initial set up and baseline current recording, 10 μL of the bead suspension (in 1 mM buffered KCl) were injected into the micropipet and were observed optically while the system was monitored electrically. In the absence of applied voltage, motion of the beads within the capillaries was not observed.

3. RESULTS AND DISCUSSION

Initial experiments in this proof of concept demonstration relied on pH to modulate the charge of carboxylic acid- or amine-terminated polystyrene beads thereby manipulating their electrophoretic mobility and ability to effect pore blockage. At an pH of 7.0, the carboxylic acid beads carried substantial negative surface charge (zeta potential = −87 mV) due to the deprotonation of carboxylic acid groups (pKₐ ≈ 4.5) thereby making the beads responsive to an electric field. With the capillary tip at high electric potential (positively charged...
When repeated at pH 11.5, above the bead pK\(_a\) (data summarized in Table T2 in the Supporting Information). and reversible reduction in the measured current (Figure 2) above was applied, the amine beads were observed to move opposite to that used in the carboxylic acid bead experiments (potential = +69 mV at pH 7.0). When a potential of sign long as desired. When the same experiments were conducted at pH 2.5 (below the carboxylic acid bead pK\(_a\), measured zeta potential = +1.79 mV), the beads were observed to be immobile and no pore blockade could be achieved. Bead blockades were observed for applied potentials between 5 and 25 V; all measurements described below were conducted with an applied potential of 25 V.

Similar experiments were conducted with amine-terminated beads that are positively charged at pH 7.0 (pK\(_a\) ∼ 9.5, zeta potential = +69 mV at pH 7.0). When a potential of sign opposite to that used in the carboxylic acid bead experiments above was applied, the amine beads were observed to move toward the pore and block it, also producing stable, indefinite, and reversible reduction in the measured current (Figure 2) (data summarized in Table T2 in the Supporting Information). When repeated at pH 11.5, above the bead pK\(_a\), the deprotonated and neutral amine beads (zeta potential = +6.3 mV) still moved in the same direction, but more slowly and with insufficient driving force to block the pore. This most likely resulted from electroosmotic flow caused by the deprotonated silanol (Si–OH) groups (pK\(_a\) ≈ 4) on the capillary surface. To confirm this, we microscopically examined the same beads and solution above the planar surface of a borosilicate glass Petri dish, and observed that the beads moved only when close to the glass surface, where the electroosmotic flow is largest. This is also consistent with the complete immobility of the carboxylic acid beads at acidic pH, since both the silanol groups on the capillary surface and the carboxylic acid groups on the beads are protonated and neutral. We presume that, in the experiments with the carboxylic acid beads at neutral pH, the force on the beads due to the electric field acting on the charged beads is greater than the opposing force due to electrosmotic flow, thereby enabling the beads to be driven to the pore.

Figure 2 shows a larger fraction of blocked current obtained with the amine beads compared to the carboxylic beads (81% vs 24%). In general, from measurements of the carboxylic acid beads in all capillaries and amine beads in three capillaries, the amine beads resulted in higher blockages (48–91%) than the carboxylic acid beads (4.7–66%) (Tables T1 and T2 and Figure S4 in Supporting Information), with some exception (Capillary 4 for carboxylic acid beads and Capillary 3 for amine beads). Optical observations indicated that the amine beads were typically immobilized closer to the capillary tip than the carboxylic acid beads, and it was common to observe some of the amine beads passing completely through the capillary tip. Since the amine beads were on average smaller than the carboxylic acid beads, this and additional observations suggest that the magnitude of the blockade is highly dependent on the relative sizes of the beads and the capillary tip.

In further support of this, a subsequent experiment with the carboxylic acid beads in a different capillary (with same nominal pore diameter of 2 μm) resulted in microscopic observation of some of the beads passing through the pore and, for those that blocked the pore, a larger reduction of current (63% average current blockade, Capillary 4, Table T1 in Supporting Information). Since the carboxylic acid beads in this experiment were drawn from the same batch as the carboxylic acid beads in previous experiments, complete passage of the beads through the pore suggests that the tip diameter of the micropipet was larger than the ones used in previous experiments. Additionally, measurement of the carboxylic acid beads in another capillary (Capillary 2 in Table T1 in Supporting Information) showed a reproducible bimodal blockade current (average blockade percentage of 24% and 6.9%) and two reproducible immobilization locations (with the 24% block occurring closer to the capillary tip (Figure S2, Supporting Information)) as the voltage was reversed and the experiment repeated. Since this measurement was obtained with one capillary, this suggests that beads of two different sizes or shapes were separately participating in the blockade.

The current measured for carboxylic acid beads passing completely through the tip of Capillary 4 (Figure S3, Supporting Information) displayed similar characteristics to previously reported measurements of beads traversing conical capillaries, specifically with respect to the rapid decrease in blockade current as the bead passes through the tip to the
The model predicts increasing blockade resistance with increasing ratio of particle radius to capillary radius \( r_p/r_c \). For a spherical particle fully contacting the interior of a conical capillary with circular cross-section, \( r_p/r_c \) is close to unity and the model predicts complete block of the current. None of our measurements showed complete blockades, although one capillary measured with the amine beads showed 90% average block (Capillary 2, Table T2, Supporting Information). These measurements of incomplete block indicate that the particles did not fully contact the capillary interior, possibly due to noncircular cross sections of the particle or capillary, presence of asperities on the bead or capillary surfaces, or adherence of the particle to the capillary wall before full contact. Evidence from the previously discussed carboxylic acid beads with a conical capillary as a simple cone, we estimated the electric field in the capillary with circular cross-section, \( \mathbf{E} \), by modeling the electric force on a 20-mer ssDNA on the bead surface (Supporting Information). On the basis of the average current measured for the target DNA sequence in three capillaries (average blockades: 38%, 39%, and 37%; Table T3 in the Supporting Information), the blockade of one of the capillaries could not be reversed after five measurements.

We hypothesize that incubation of the beads with ssDNA results in a significant amount of nonspecific binding for both complementary and noncomplementary sequences. In the control experiments, the DNA binding to the bead is entirely nonspecific and presumably less strongly bound than the complementary DNA. Still, the beads with nonspecifically bound DNA are negatively charged and electrophoretically mobile, allowing them to be driven to the pore. In the pore, the electric field is sufficiently strong to remove the nonspecifically bound DNA from the bead, which causes a reduction in bead charge and electrophoretic mobility, enabling the electroosmotic flow to exceed the electrophoretic force and carry the bead away from the pore. For complementary DNA sequences, the bead likely carries specific and nonspecifically bound DNA but the strong electric field is insufficient to remove the hybridized DNA from the bead (Figure 3b). This transient blockade was observed infrequently, with most of the control beads unable to block the pore.

The control and target experiments were repeated at least three times; measured zeta potentials and electrophoretic mobilities are listed in Table 1 with the qualitative results of the electrical measurements (Data summarized in Tables T3 and T4 in the Supporting Information). Quantitatively, the magnitude of the blockades measured for the target DNA was consistent between the three capillaries (average blockades: 22.0%, 25.6%, and 25.6%; Table T3 in the Supporting Information). The blockade of one of the capillaries could not be reversed after five measurements.

We modeled the bead electrophoretic force by equating it to the drag force on the bead when it is moving with constant speed. The measured mobility is the proportionality constant between the speed and the electric field. By modeling the capillary as a simple cone, we estimated the electric field in the capillary as a function of position and found forces between 1.36 and 5.44 nN as the capillary radius tapered from two bead radii to one bead radius (Supporting Information).

Next, nucleic acid detection was measured using 20-mer polyA ssDNA as a simple target sequence and PNA-beads conjugated with 12-mer polyT PNA as the complementary probe. To assess the PNA conjugation to the carboxylic acid beads, the beads’ zeta potential before PNA conjugation was measured to be −87 mV, after ethanolamine capping +5.75 mV, and after three washes with 0.4x SSC buffer −4.39 mV. After capping and washing, the beads were observed to aggregate. Without incubation with DNA, the PNA-beads in the micropipet were seen to follow the electroosmotic flow away from the pipet tip, indicating that the PNA-beads alone were unable to block the pore. Incubation of the PNA-beads with polyA target ssDNA resulted in well-dispersed beads with a measured zeta potential of −71.1 mV, and motion toward the pipet tip in the same applied voltage, ultimately blocking it (Figure 3d). This transient blockade was observed infrequently, with most of the control beads unable to block the pore.

The control and target experiments were repeated at least three times; measured zeta potentials and electrophoretic mobilities are listed in Table 1 with the qualitative results of the electrical measurements (Data summarized in Tables T3 and T4 in the Supporting Information). Quantitatively, the magnitude of the blockades measured for the target DNA was consistent between the three capillaries (average blockades: 22.0%, 25.6%, and 25.6%; Table T3 in the Supporting Information). The blockade of one of the capillaries could not be reversed after five measurements.
capillaries (Table T3, Supporting Information), a force of 13.9 pN is obtained from the model for the 20-mer ssDNA on the bead surface. This force is less than the 57 pN rupture forces measured for 8 bp DNA-PNA with optical tweezers.42

To further investigate the selectivity of the sensor, we detected ssDNA with a nonrepeating sequence, a 12-mer portion of a gene encoding the anthrax lethal factor.43 To allow direct comparison with the previous experiment detecting 20-mer ssDNA, we added an 8-mer polyA tail to the 12-mer anthrax sequence. To enhance the binding of complementary ssDNA and minimize nonspecific binding, a PEG spacer was added to the amine-functionalized 12-mer complementary PNA capture probe.44 After PNA conjugation, capping with ethanolamine, and washing, the measured zeta potential of this PNA-bead preparation was −2.75 mV. The PNA-beads were divided into two volumes, one incubated with 20-mer target anthrax ssDNA and the other with control 20-mer polyT ssDNA. Measured zeta potentials after incubation were −56.7 mV for target beads and −39.0 mV for control beads. As with the previous experiments described above, the presence of complementary DNA led to permanent blockades, whereas its absence led to transient or no blockade (Figure 4) (Videos, Figure 3. Schematic of motion of PNA-bead under an applied potential and resultant measured current. (a) (i) PNA-beads (green) with nonspecifically bound ssDNA and specifically bound ssDNA hybridized to PNA on the bead are negatively charged and electrophoretically mobile. (ii) In the strong electric field at the pore tip, the nonspecifically bound DNA is removed from the bead, but the hybridized DNA is not, leaving the bead with sufficient negative charge to remain blocking the pore indefinitely. (b) Measured permanent current blockade for PNA-beads incubated with complementary polyA DNA, corresponding to (a). (c) For PNA-beads incubated with only noncomplementary DNA, any DNA bound to the bead is nonspecific; the strong electric field at the pore tip removes the nonspecifically bound DNA, reducing the bead’s charge sufficiently so that the electroosmotic flow (red arrows) is able to remove the bead from the pore tip (ii). (d) Transient current blockade measured for PNA-beads incubated with noncomplementary polyT DNA, corresponding to (c).

Table 1. Summary of Experimental Results for Target and Control Samples

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<th>control PolyT</th>
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“Zeta potential and electrophoretic mobility measured after ssDNA incubation and results of micropipet electrical measurements.

Figure 4. (a) Permanent ionic current drop caused by beads incubated with target anthrax ssDNA. The blockade was reversible and repeatable, as seen by reversals of the applied voltage (dashed lines). (b) Transient ionic current drops were seen occasionally with beads incubated with the noncomplementary, control ssDNA.
With either or both of these mechanisms contributing, there DNA from the bead as described above in Table 2 and Figure 4.

capillaries 4

recorded only in the presence of target anthrax ssDNA. Since the melting temperature for the single base mismatch DNA is 7.39 mV. As described above, the PNA-beads were divided into two volumes, one incubated with the anthrax LF ssDNA and one with the single base mismatch ssDNA (Table T5 (Capillaries 4–6) and Table T7, Supporting Information). In each of the capillaries measured, following blockade, the voltage was reversed to remove the bead from the capillary tip to attempt further blockades. In one of the six capillaries, after five detection events, the bead was not able to be removed from the blockade site with reversal of applied voltage and the experiment was terminated.

To investigate the sequence specificity of the sensor, we created a 20-mer ssDNA with the same sequence as the 20-mer ssDNA for the anthrax LF experiment described above but with a single base mismatch. The measured zeta potential after PNA conjugation, capping with ethanolamine, and washing was −7.39 mV. As described above, the PNA-beads were divided into two volumes, one incubated with the anthrax LF ssDNA and one with the single base mismatch ssDNA (Table T5 (Capillaries 4–6) and Table T7, Supporting Information). Table T3 summarizes the results of three separate experiments, which are consistent with our previous results described above. Occasional transient blockades were observed in the presence of the mismatch DNA sample; permanent blockades were recorded only in the presence of target anthrax ssDNA.

Since the melting temperature for the single base mismatched DNA is above room temperature, the mismatched DNA-PNA is stably hybridized following incubation of the DNA with the PNA-beads. The measurement of the transient block could result from the mismatched hybrid having a sufficiently low force of dissociation that the DNA strands are removed in the strong electric field of the pore, or the number of mismatched strands hybridized to the bead could be less and the electric field of the pore is removing nonspecifically bound DNA from the bead as described above in Table 2 and Figure 4. With either or both of these mechanisms contributing, there was no permanent blockade resulting from the single base mismatched DNA.

Comparison of beads incubated with complementary target ssDNA in Table 1 (target: poly A) and Tables 2 and 3 (target: Anthrax-LF) shows that the magnitudes of the zeta potentials and mobilities in Table 1 were larger than those in Tables 2 and 3. A possible explanation for these results is the lack of registration required for the hybridization of polyA ssDNA, compared to the exact registration required for hybridization of Anthrax ssDNA. Longer ssDNA targets may improve the electrophoretic mobility of the hybridized beads, while longer strands of nonspecifically bound ssDNA would still be expected to detach from the bead in the strong electric field at the sensing zone to result in only transient ionic current blockades.

The limits of detection were probed by serially diluting the 20-mer target anthrax ssDNA in hybridization buffer and repeating the incubation with PNA beads and nanopore measurement as described above (Table T8, Supporting Information). Pore blockade was observed down to a concentration of 10 pM. At this concentration, we observed some beads only transiently blocking the pore before permanent block was achieved, indicating the presence of both nonspecific and complementary ssDNA bound to the beads, as well as a smaller amount of bound complementary DNA. We are currently working to reduce the concentration limit of detection of our micropipet-based system through the use of longer target ssDNA oligomers.

Overall, our system performed as expected for detection of specific DNA sequences. Using the conditions described, polyA or anthrax DNA were successfully detected in every capillary tried (nine capillaries total), with no false positives (no permanent blockade) observed in any capillary (nine capillaries total), including ssDNA with only a single base mismatch. The lowest DNA concentration successfully detected with our unoptimized system was ∼10 pM, an unimpressive detection level compared to other published approaches, including a PNA sandwich-hybridization assay for anthrax with a DNA detection limit of as low as 10 zmol. Yet, this binary detection system could exhibit a very low detection limit if the system were scaled down such that a submicrometer PNA-bead conjugate would assume sufficient charge for electrophoretic mobility and pore blocking upon binding one or a few target DNA molecules. Of course this low detection limit comes at the expense of the capability to determine target DNA concen-
tration. Nevertheless, as discussed in the Introduction, there are a number of important applications where a binary (yes/no) signal for the presence/absence of the DNA target is sufficient. Further, the large and sustained reduction in current resulting from the blockade of the pore by a PNA-bead conjugate with bound target DNA provides an easily detectable signal for the presence of the DNA target that can be displayed with simple electronics. Using a simple inverting operational amplifier and light emitting diodes, we constructed a binary indicator of the target DNA detection, demonstrating its potential application in a simple, potentially low-cost device. In the future, we will work to scale down using silicon micromachining technology and to optimize this system in order to achieve meaningful detection limits and to apply it to the sensing of NA in biological samples.

ASSOCIATED CONTENT

3 Supporting Information
Supporting Information contains the following: DLS diameter measurements of carboxylic acid and amine beads; a list of unblocked and blocked currents measured for all capillaries; microscopic images of beads immobilized at a capillary tip; measurements of current during passage of two carboxylic acid beads completely through a capillary tip; measurements of zeta potential and occurrence of blockade as a function of Anthrax beads completely through a capillary tip; measurements of zeta potential and occurrence of blockade as a function of Anthrax DNA PNA-bead in a capillary. This material is available free of charge via the Internet at http://pubs.acs.org.

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Notes
The authors declare no competing financial interest.

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