



Structural and positional impact on DNzyme-based electrochemical sensors for metal ions

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Abstract

The rapid, accurate and convenient detection of heavy metal is very important to public health. Here, we developed a DNzyme-based electrochemical sensor for Pb²⁺. A DNzyme-including and Pb²⁺ active probe was anchored to the biosensing interface, based on the well-defined self-assembled, three-dimensional DNA nanostructure. The results indicate that the detection performance depends on the change of distances between the methylene blue and the electrode surface. The limit of detection (LOD) could reach the concentration of 0.01 μM Pb²⁺, and the signal change shows semi-logarithmic relationship with the concentration of Pb²⁺ from 0.01 μM to 100 μM. The biosensor also presents good stability and specificity to detect Pb²⁺ in tap or river water. This method not only provides promising approach for improving the performance of tetrahedra in detecting Pb²⁺, but helps deepen the understanding of tetrahedral structure design and how the position of electroactive groups affects the performance of electrochemical sensing.

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Lead ion (Pb²⁺) is an important pollutant that can cause serious physical damage, including abdominal pain, irritability, headaches, memory problems, and constipation.^{1,2} Current

methods for detecting Pb²⁺ require inductively coupled plasma/mass spectroscopy (ICP/MS),³ a rather complex laboratory technique. The selective electrode is also a widely used method for detecting lead ion, despite of its high limit of detection (LOD) and less sensitivity.⁴ Many efforts have been made for quick, easy, and sensitive detection of lead ions.^{5–10} The catalytic activity of DNzymes is metal ion cofactor-dependent, as in the case with some proteins.^{11–15} Much research has focused on the 8-17 DNzyme owing its high specificity for Pb²⁺.^{16–19} The 8-17 DNzyme has a specific sequence, and its complementary substrate chain contains only one RNA adenine. In the presence of lead ions, the 8-17 DNzyme cuts and sheds the substrate chain.^{20–22}

Xiao et al riveted the 8-17 DNzyme sequence to a gold electrode interface using a covalent modification method and, in the presence of Pb²⁺, the trans-acting catalytic strand cleaved the

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sessile phosphodiester of the substrate into two pieces at the ribonucleotide site. By electrochemically measuring the change in the current value, a detection limit of 0.3 μM was achieved.²³ However, the detection performance can be markedly affected by the single-strand surface probe jamming effect, probe intertwining, lodging, and non-specific adsorption. Yun et al used a colorimetric method to detect lead ions using a DNAzyme to cleave the substrate, and the sheared product was combined with nanogold to detect lead ions with a detection limit of 5 nM.²⁴ However, the colorimetric method involves visual observation (which is not accurate) and requires a large optical instrument. Currently, with the development of nanotechnology, a variety of approaches based on nanomaterials have been used in the sensing field,^{25–31} such as DNA tetrahedron which is a nanostructure formed by equal mixing and annealing of four single-stranded DNAs.^{32–38} Because DNA obeys specific A-T and G-C base pairing rules, it is possible to precisely control the size of tetrahedral DNA nanostructures and introduce functionalized sequences. The three-dimensional (3D) DNA tetrahedron applied in electrochemical biosensors offers a great advantage in the field of sensing.^{8,39–42} The three vertices of the tetrahedron probe are modified with a thiol group and fixed to the surface of the gold electrode. The distance between the probes can be controlled to ensure the desired orientation. The tetrahedral structure has a certain thickness (~6 nm) and provides a solution-like environment. The reaction environment improves the affinity and enables good target detection.^{40–48} Zhou et al embedded a DNAzyme into one side of a DNA tetrahedron and detected Pb^{2+} by fluorescence with a detection limit of 3.9 nM in cells.⁴⁹ However, detecting lead ions by fluorescence is cumbersome and reading the fluorescent signals requires a large instrument.

To solve these problems and explore the impact of the DNA tetrahedral structure and methylene blue electrochemical active position on the DNAzyme-based signal response to Pb^{2+} , we combined DNA tetrahedral electrochemical biosensing technology and DNAzyme-based targeted RNA-cleavage technology to establish a lead-ion detection platform. Initially, we designed a detection model (Model A) by referring to Xiao's²³ electrochemical scheme and replaced one edge of the tetrahedron with a DNAzyme. Contrary to our expectation, the detection signal decreased instead of increased upon Pb^{2+} binding, which was due to steric hindrance on the surface of the gold electrode. Thus, we designed Model B and Model C to achieve lower detection limit of lead ions. Based on these detection models, we explored the influence of the distance between the electroactive groups and the gold electrode surface on the detection performance.

Methods

Chemicals and materials

Tris(2-carboxyethyl)-phosphine hydrochloride (TCEP) and 6-mercaptopentanol (MCH) were purchased from Sigma-Aldrich. $\text{Pb}(\text{OAc})_2 \cdot 3\text{H}_2\text{O}$, $\text{Zn}(\text{OAc})_2 \cdot 2\text{H}_2\text{O}$, $\text{Co}(\text{OAc})_2 \cdot 4\text{H}_2\text{O}$, $\text{Mn}(\text{OAc})_2 \cdot 4\text{H}_2\text{O}$, $\text{Ni}(\text{OAc})_2 \cdot 4\text{H}_2\text{O}$, $\text{Ca}(\text{OAc})_2 \cdot \text{H}_2\text{O}$, and $\text{Mg}(\text{OAc})_2 \cdot 4\text{H}_2\text{O}$ were purchased from Sigma-Aldrich and used without further processing. All metal acetate was dissolved in

10% CH_3COOH as a 0.1 M stock solution. Further dilution was performed with 50 mM acetic acid. The following buffers were used in the experiments: TM buffer (50 mM Tris-acetate, 1 M NaCl, pH 6.5; used to form DNA tetrahedra) and soaking buffer (50 mM Tris, 0.5 M NaCl, pH 6.5). DNA oligonucleotides with the following sequences (from 5' to 3') were synthesized and purified by Takara Biotechnology, Inc. (Shanghai, China):

Strand AZM: MB-TTCATCTCTTCTCCGAGCCGGTTCGAAATAGTGAGTACATTACAGCTTGCTACACGAGAAGAGCCGCCATAGTA.

Strand AZ: TTCATCTCTTCTCCGAGCCGGTTCGAAATAGTGAGTACATTACAGCTTGCTACACGAGAAGAGCCGCCATAGTA.

Strand B: SH-TATCACCAGGCAGTTGACAGTGTAGCAAGCTGTAATAGATGCGAGGGTCCAATAC.

Strand C: SH-TCAACTGCCTGGTGATAAAACGACACTACGTGGGAATCTACTATGGCGGCTCTTC.

Strand D: GCTTCCCACGTAGTGTCGTTTGTATTGGACCTCGCAT-SH.

Strand DS: ACTCACTATrAGGAAGAGATGGCTTCCCACGTAGTGTCGTTTGTATTGGACCTCGCAT-SH.

Strand DSM: MB-TTACTCACTATrAGGAAGAGATGGCTTCCCACGTAGTGTCGTTTGTATTGGACCTCGCAT-SH.

Strand substrate: ACTCACTATrAGGAAGAGATG.

Detection Model A involved the use of five strands: AZM, B, C, D, and the strand substrate. Detection Model B involved four strands: AZM, B, C, and DS. Detection Model C also involved four strands: AZ, B, C, and DSM. M refers to a chain with the modifications of electrochemically active group (methylene blue, MB), and rA refers to an adenine ribonucleotide).

Instruments

Electrochemical measurements were performed on a CHI 760E electrochemical workstation (CH Instruments Co., Shanghai, China). A conventional three-electrode system including a gold electrode (2 mm in diameter) as the working electrode, an Ag/AgCl electrode (3 M KCl) as the reference electrode, and a platinum wire as the auxiliary electrode was used for cyclic voltammetry (CV) and square wave voltammetry (SWV). CV was carried out from 0 to 0.7 V at a scan rate of 100 mV/s. SWV was carried out from 0.5 to 0 V at a frequency of 50 Hz with an amplitude of 50 mV.

Methods

In the beginning, we designed Model A, which is an incomplete tetrahedron formed by the annealing of five strands including strand AZM, strand B, strand C, strand D, and the substrate strand. The 5' end of the A chain was modified with MB, and the substrate strands were later hybridized with it. The edge of the tetrahedron containing DNAzyme was free to move in solution. After the addition of lead ions, the phosphodiester bond at the rA of the substrate chain was cleaved into two pieces. Since the free energy was reduced, both pieces of the substrate were latterly shed from the tetrahedron, and the A chain was opened from the neck ring structure to a single-stranded state. Then, the distance between the MB to the electrode surface

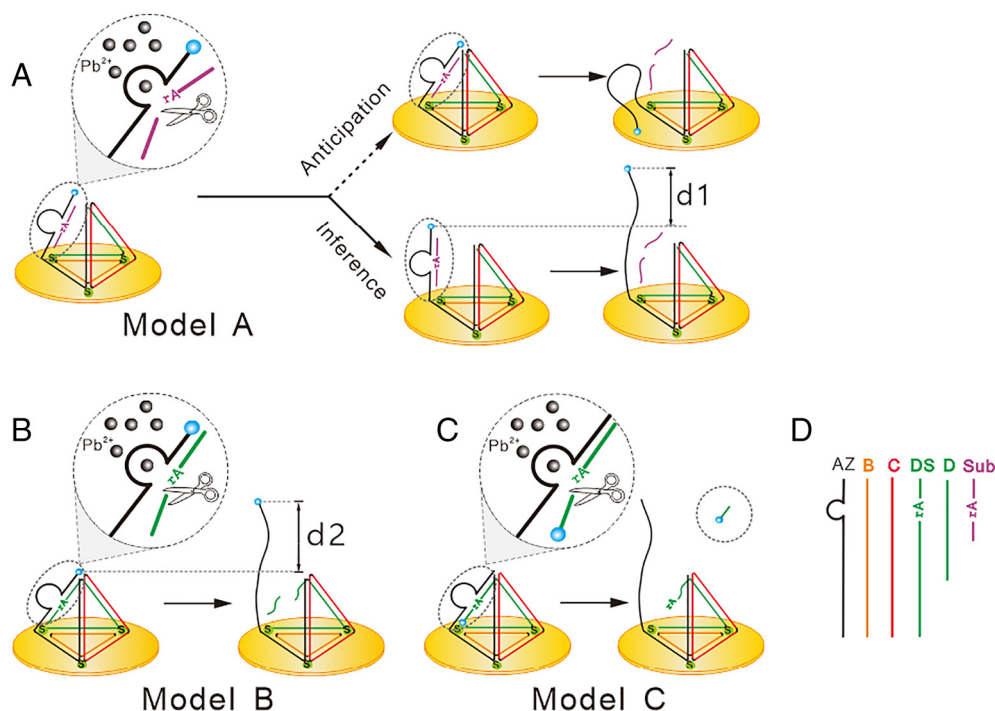


Figure 1. Schematic illustration of tetrahedral DNAzyme-based enzymatic cleavage. (A) Two hypothetical types of structural changes occurring in the presence of lead ions in Model A. (B) Structural change occurring in the presence of lead ions in Model B. (C) Structural change occurring in the presence of lead ions in Model C. (D) DNA strands making up the DNA tetrahedron.

increased. In anticipation, we initially considered that the chain modified with MB would be free in solution and have a certain degree of flexibility. The electroactive group collides freely with the surface of the electrode, and an increased electrochemical signal could be displayed as also described by Xiao Yi in his single-stranded electrochemical detection procedure. Interestingly, the signal decreased after adding lead ions. This phenomenon may be due to steric hindrance of the tetrahedron, as shown by inference in Figure 1, A. To investigate the effect of the position of modification of the electroactive group and the tetrahedron structure on the detection performance, we designed Model B (Figure 1, B) and Model C (Figure 1, C). Model B involves a complete tetrahedron formed by the annealing of strands AZM, B, C, and DS. The substrate chain is integrated into the tetrahedral strand D as DS. MB is also modified at the 5' end of the A chain. In the presence of lead ions, the DNAzyme cleaves the substrate. Thus, the A chain is free in solution and the distance of the MB label from the electrode surface is increased. Model C involves a complete tetrahedron formed by the annealing of strands AZ, B, C, and DSM, where the 5' end of the D chain is modified with MB (Figure 1, C). In the presence of lead ions, the DNAzyme cleaves the substrate strand into two pieces at the ribonucleotide site, and the substrate chain with the MB is detached from the tetrahedron. We assembled the synthesized tetrahedron onto the surface of the gold electrode and recorded the absolute current values of the three models every two minutes using SWV. Wait until the current value stabilizes (as shown in Figure S1 in the Supporting Information)

to calculate the percentage change in current value before and after the reaction.

Synthesis and characterization of DNA tetrahedra

For each model (Model A: strands AZM, B, C, D, and the substrate strand; Model B: strands AZM, B, C, and DS; Model C: strands AZ, B, C, and DSM), the strands were dissolved in TM buffer, and the mixtures were heated at 95 °C for 10 min and quickly cooled down to 4 °C within 1 min, yielding a final DNA tetrahedra concentration of 1 μM. Each set of tetrahedral constituent chains is mixed and annealed to form a tetrahedral nanostructure, and then verified by polypropylene gel electrophoresis. The details of the polyacrylamide gel electrophoresis method are given in Table S1 in the Supporting information.

Preparation of self-assembling DNA tetrahedra on gold electrodes

Before preparing self-assembling DNA tetrahedra, gold electrodes were first electrochemically cleaned in 0.5 M NaOH. Then, the gold electrodes were polished for 2 min using a microcloth with 0.3 and 0.05 μm γ-alumina particles. Then, the polished electrodes were sonicated in ethanol and Milli-Q water (3 min for each sonication step). Next, the electrodes were electrochemically cleaned in 0.5 M H₂SO₄ by scanning the potential between the oxidation and reduction of gold (0.35 and 1.5 V, respectively), and finally the electrodes were cycled in 0.5 M H₂SO₄/0.01 M KCl. To determine the

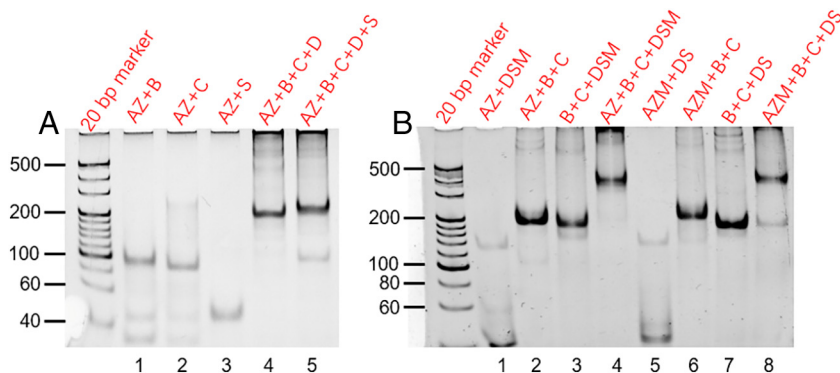


Figure 2. PAGE (8%) analysis of the self-assembled DNA strands using all possible combinations of the four strands. The bands in panel **A** (lanes 1-3) and panel **B** (lanes 1-3, 5-7) feature a common tailing phenomenon, suggesting that the hybridization between two or three strands was incomplete. The concentration of all strands was 1 μ M. **(A)** Lane 4 represents AZM, B, C, D, four chains consisting of non-intact tetrahedra. Lane 5 represents AZM, B, C, D, substrate, five chains form non-integral tetrahedra. **(B)** Lane 4 and lane 8 show complete tetrahedral PAGE bands for the Model B and Model C designs, respectively.

effective active surface area, the electrodes were scanned in 0.05 M H_2SO_4 . After cleaning the electrodes, 3 μL of 1 μM DNA tetrahedron, which was annealed in TM buffer with 3 mM TCEP, was incubated overnight on the freshly cleaned gold electrodes at room temperature to form a self-assembled monolayer.

Procedures for electrochemical measurement

DNA tetrahedra were used to form a self-assembled monolayer on the electrode. Then, the modified electrodes were exposed to a 2 mM MCH solution in phosphate-buffered saline (pH 6.5; 50 mM Tris, 1 M NaCl) at room temperature for 1 h to avoid interference from oxygen, which shows a similar reduction potential to that of MB. After the blockage, the electrodes were transferred to soaking buffer for 1 h. Next, the current signal response was interrogated by CV and SWV. For enzymatic cleavage reactions, the electrode was reacted with various concentrations of target Pb^{2+} in buffered Tris (50 mM Tris-acetate 0.5 M NaCl, pH 6.5) for 60 min at room temperature to achieve maximum substrate cleavage on the gold surface, after which the electrodes were transferred (within 1 h) to buffered Tris (pH 6.5) and thoroughly washed to remove the substrate chain. Control electrodes were prepared in an identical fashion.

Results

Characteristic of DNA tetrahedra

The assembly of the DNA tetrahedra was characterized by polyacrylamide gel electrophoresis (PAGE), and cross-hybridizations between two or three strands during the thermal annealing process were also evaluated. Inspection of Figure 2 reveals that the bands in Figure 2, A (lanes 1-3) and Figure 2, B (lanes 1-3, 5-7) featured a common tailing phenomenon, suggesting that incomplete hybridization occurred between the two or three strands. In contrast, when all strands were combined in equimolar quantities, a clearly independent and non-trailing band with much lower electrophoretic mobility was observed (lanes 4 and 5 in Figure 2, A, lanes 4 and 8 in Figure 2, B).

Characterization of the enzyme cleavage reaction

To analyze the enzymatic cleavage reaction, we used PAGE to characterize the DNzyme-based tetrahedron cleavage products. Figure 3, A and B shows enzymatic cleavage results in polyacrylamide gels with concentrations of 8% and 15%, respectively, using Model A. The results obtained with the 8% polyacrylamide gel showed that the bands for tetrahedra that hybridized to the substrate chains (Figure 3, A, AZ + B + C + D) had a lower electrophoretic mobility than those that did not hybridize to the substrate chains (Figure 3, A, AZ + B + C + D + S). After the addition of lead ions, the tetrahedron band (Figure 3, A, AZ + B + C + D + Pb^{2+}) showed a higher electrophoretic mobility than the tetrahedron band that hybridized to the substrate chain (Figure 3, A, AZ + B + C + D + S). The migration distance of the band (Figure 3, A, AZ + B + C + D + Pb^{2+}) was the same as the tetrahedral bands that did not hybridize with the substrate (Figure 3, A, AZ + B + C + D). This phenomenon indicates that the substrate strand was capable of hybridizing to the enzyme strand and then could be successfully cleaved in the presence of lead ions. After the addition of lead ions, a new band appeared as shown by the black arrow in Figure 3. This new band was the substrate-cleavage product, and the band provides strong evidence that the enzymatic cleavage reaction occurred.

Figure 3, C and D shows the polyacrylamide gels for Model B and Model C at concentrations of 8% and 15%, respectively. The 8% polyacrylamide gel shows that, after the addition of lead ions, the bands for the tetrahedra showed a higher electrophoretic mobility (versus the absence of Pb^{2+}), which indicates that a cleavage reaction occurred in the presence of Pb^{2+} and that the substrate chain detached from the tetrahedron after cleavage. The 15% polyacrylamide gel shows that, after adding lead ions, a new band was formed, i.e., the substrate-cleavage product (shown by the arrow). These two phenomena indicate that the enzymatic cleavage reaction occurred.

Electrochemical signal responses of three detection models

As shown in Figure 4, A, the DNzyme-based tetrahedron sensor has a very strong response to the presence of Pb^{2+} . In the absence of Pb^{2+} , we observe only small Faradaic current change.

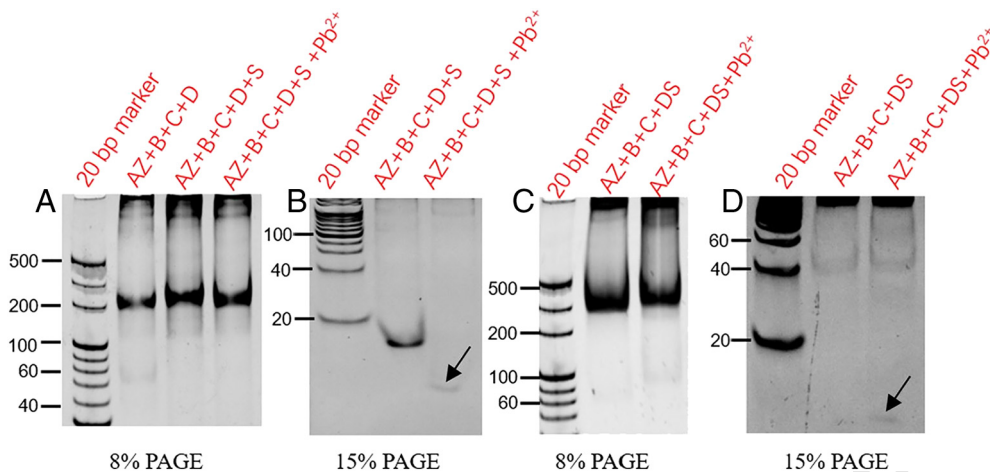


Figure 3. PAGE (8% and 15%) analysis of the enzymatic substrate-cleavage reaction with the tetrahedron. The concentration of all strands was 1 μM . (A and B) Electrophoretic results for the Model A enzymatic cleavage reaction with polyacrylamide concentrations of 8% and 15%, respectively. (C and D) Electrophoretic results of polyacrylamide gels of Model B and Model C at concentrations of 8% and 15%, respectively. The arrows in B and C indicate the new band products.

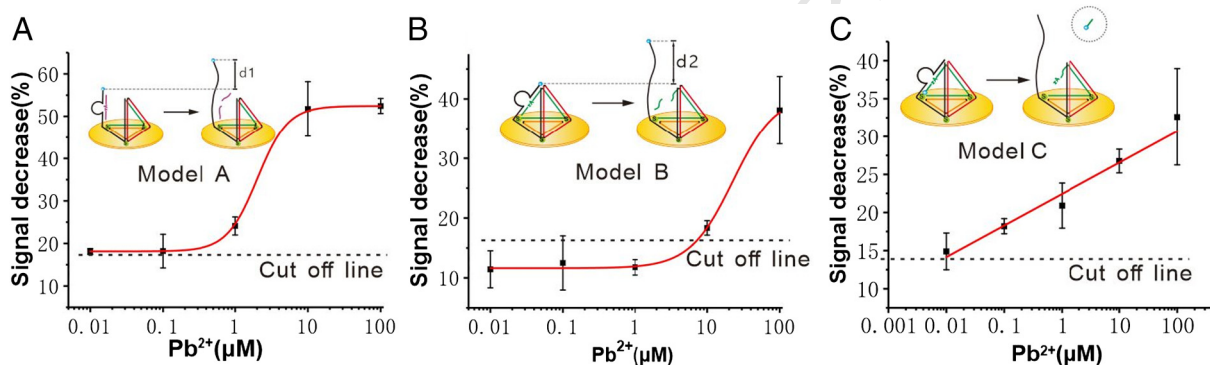


Figure 4. Simultaneous detection of various concentrations of Pb^{2+} (from 10.0 nM to 100.0 μM) with the three different types of DNAzyme tetrahedron nanoprobe. The figure is a schematic representation of the different structure designs. The error bars reflect the standard deviation of three independent measurements. The dashed line represents the percent decrease in the electrochemical signal of the control group.

Upon the increase of the Pb^{2+} concentration, we observed a large Faradaic current change. Model A is sensitive to detect as low as 0.1 μM Pb^{2+} . The percentage change of the measured current towards a target of 0.1 μM (18.41%) is still larger than the mean blank value (13.94%) plus 3 standard deviations (1.47%). These results confirmed that the nanoprobe could achieve sensitive detection of the target metal ions. As shown in Figure 4, B, the resulting change percentage of Faradaic currents exhibited semi-logarithmic linear relationships with the concentrations of the target metal ions in the range of 10–100 μM . Model B is sensitive to detect as low as 10 μM Pb^{2+} . The change percentage of the measured current from a target of 10 μM (18.39%) is still larger than the mean blank value (11.66%) plus 3 standard deviations (1.56%). As shown in Figure 4, C, the change percentage of Faradaic currents also presents semi-logarithmic relationship along with the concentrations of the target Pb^{2+} from 0.01 μM to 100 μM . Model C is sensitive to detect as low as 0.01 μM Pb^{2+} . The percentage change of the measured current from a target of 0.01 μM (15.44%) is still larger than the mean blank value (11.79%) plus 3 standard deviations (1.00%). There is no

significant difference in current change between the blank groups of the three models. The error bars shown here reflected the standard deviation of three independent measurements, and the cutoff line (dashed line in Figure 4) is the average current change percentage of the blank group plus 3 times the standard deviation of the current change. Table S2, Table S3 and Table S4 in the Supporting Information show the current signal change of the biosensor before and after the addition of Pb^{2+} for the three detection models. The CV and SWV curves for the three models are shown in Figure S2, Figure S3 and Figure S4 in Supporting Information.

Electrochemical signal analysis of tetrahedra with three different probe designs

The three models were found to have different detection limits and linear ranges of detection. Comparing Model A with Model B, we found that the detection limit of Model A (0.1 μM) was 100-fold lower than that of Model B (10 μM). Comparing Model B with Model C, we found that Model C had a lower

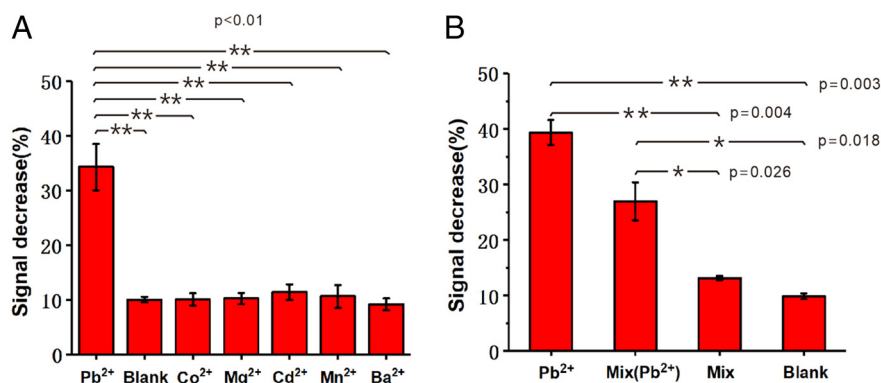


Figure 5. The Model C sensor was both specific (capable of rejecting similar ions) and selective (uninhibited by complex samples). (A) Little signal change was observed when the sensor was challenged with various divalent metal ions other than Pb²⁺ (all at 100 μM). The signal change for positive group is significantly different from that for the blank control ($P < 0.01$). (B) The sensor was selective enough to employ with mixed ions (all at 100 μM).

detection limit of 0.01 μM, which was 1000-fold better than Model B. The linear detection range of Model A was 1 μM to 10 μM, that for Model B was 10 μM to 100 μM, and that for Model C was 0.01 μM to 100 μM.

According to Xiao's work, we designed a DNA tetrahedron with one side consisting of a DNAzyme (Model A). Our initial analysis revealed that, in the presence of Pb²⁺, a catalytic strand cleaves the edge of the DNA tetrahedron. These fragments presumably dissociate from the complex, allowing the electroactive group at the 5' end of the A chain to collide freely with the electrode surface, in turn allowing the MB group to transfer electrons to the electrode and produce an increased signal (Figure 1, anticipation). However, our results showed a signal decrease, so we investigated the underlying cause of this unexpected result using Xiao's DNAzyme, a method of chemically detecting lead ions wherein a single strand of DNAzyme is assembled on the surface of a gold electrode. The substrate chain was later hybridized. After the cleavage reaction, the substrate chain is cut into two pieces, and the single strands with enzymatic properties are flexible to some extent. This flexibility causes the MB modified at the 5' end of the DNAzyme chain to freely collide with the surface of the gold electrode, so that an increase in the current value can be detected. However, Model A involves a tetrahedron-based sensor, and the tetrahedron is constrained by steric hindrance on the surface of the gold electrode. Thus, the 5' end of the AZM chain is not free to collide with the surface of the gold electrode, but stands upright in solution, which led to the decline of the Model A electrochemical signal. In addition, the Model A structure is not completely tetrahedral. The 5' end of the A chain has a flexible structure. Thus, we designed Model B and Model C to further study the effect of the position of the electrochemically active group (MB) and the integrity of the tetrahedron on the performance of the detection platform. Our analysis was as follows.

The percentage of the signal decrease after the digestion of Pb²⁺ towards DNAzyme substrate should depend on the change of the effective distance between the MB and the electrode in Model A, B and C. For Model C, the MB is totally detached from the tetrahedral structure after digestion, and thus MB at this time has the largest distance from the electrode surface, so the

detection performance is best. For Model A and Model B, the distance between MB and electrode surface is equal after digestion, so we need to compare the distance between the MB and the electrode surface before digestion. The difference is shown in the dotted circle in Figure 1. For model A, the DNAzyme can rotate around the bottom end, and tends to lie flat on the surface, making the effective distance of the methylene blue from the electrode shorter than Model B. The distance change between the electrochemically active group and the electrode in Model A is larger than that of Model B, so Model A exhibits higher detection performance.

Selectivity and specificity for detecting lead ions

Since Model C shows the best detection performance, it was chosen as an optimal detection model. As shown in Figure 5, the specificity of the sensor was determined by challenging it with the divalent metal ions, Co²⁺, Mg²⁺, Cd²⁺, Mn²⁺, and Ba²⁺ at a concentration of 100 μM. Figure 5, A shows that signal change of the positive group was respectively significantly larger than the blank control and the five negative control groups containing different ions ($P < 0.01$). As shown in Figure 5, B, the signals from the two positive groups separately or mixed contained Pb²⁺ is effectively distinguishable from the negative control groups when exogenously metal ions were added. The sensor was demonstrated selective enough to employ with a mixed ion solution, which also suggests its stability from the interferences of other ions.

Detection of actual samples

Our biosensor (Model C) can detect Pb²⁺ in tap water and river water as shown in Figure 6. Electrochemical detection was performed by adding 0.1 or 10 μM lead ions. No significant signal change was observed with the blank group, although a clear signal change was found with the experimental group. In contrast, however, we found that the detection signal of tap water changed significantly more than that of river water. For example, the signal drop rate for the tap water-detection limit was significantly lower than that of tap water at a Pb²⁺ concentration of 10 μM. We speculate that the composition of river water is

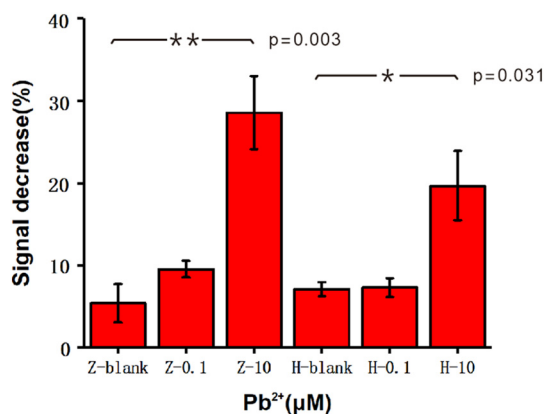


Figure 6. The detection of Pb²⁺ in tap water (TW) and river water (RW) samples with Model C. We added 0.1 or 10 μM lead ions for electrochemical detection, as indicated. No significant signal change was observed for the blank group. The positive group had a larger signal change. *P* values are displayed above the histogram.

complex, containing microorganisms or RNases that could cleave and inactivate the detection system. Therefore, among the blank groups samples, the river water showed more of a signal decrease than the tap water. The above test results suggest that the biosensor platform was also suitable for complex system samples from river and tap water, suggesting its stability.

Moreover, we also test its detection performance towards lead ions in soil extracts. As shown in Figure S5 in Supporting Information, the result presents statistically different between the blank group and the target group (*P* = 0.003). However, the signal difference for soil extracts is not so significant as that for tap or river samples. The interference might be from the possible negative effect of RNases present in soil extracts.

Conclusion

In summary, we designed a new biosensing interface to detect Pb²⁺ using a well-defined 3D DNA self-assembling nanostructure with one edge comprised of a DNAzyme. By comparing three models of DNA tetrahedron-based DNAzyme sensors, we found the impact of the tetrahedral DNA structure and the MB electrochemical active position on the signal response to Pb²⁺. Along with the distance change between the electroactive group and the electrode surface, the limit of detection (LOD) concentration and the linear range of semi-logarithm could reach 0.01 μM and 0.01 μM-100 μM respectively. The sensitivity of our optimal detection method was improved by an order of magnitude (reaching 0.01 μM) versus the non-nanomaterial modified electrochemical method for Pb²⁺. Model C not only showed good specificity for Pb²⁺ over other ions such Co²⁺, Mg²⁺, Cd²⁺, Mn²⁺, and Ba²⁺, but also enabled Pb²⁺ detection with samples of tap water and river water. Our design strategy improved the performance of DNA tetrahedra for detecting Pb²⁺, which could also enhance the understanding of how the tetrahedral structure and the position of electroactive groups affects electrochemical sensing of other analytes, based on the DNA tetrahedron-detection platform.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.nano.2019.102035>.

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