A sensitive fluorescent method for the detection of Hg²⁺ was developed based on triple-helix molecular switch (THMS)-induced hybridization chain reaction (HCR) amplification. THMS was composed of a T-rich mercury-specific probe and an initiator probe, designed by the Watson-Crick and Hoogsteen base pairings and employed as a signal trigger. Two hairpin probes containing the G-quadruplex sequence were used as signal amplification elements. In the presence of Hg²⁺, the T-Hg²⁺-T mismatch resulted in disassembling the THMS and releasing the initiator probe. One of the hairpins was opened by the released initiator probe, which triggered a successive cross-opening of two hairpins based on the strand displacement principle, resulting in the formation of long-chain DNA with multiple G-quadruplex. When thioflavin T (ThT), a fluorophore, was bound to the G-quadruplex, an obvious fluorescence enhancement would occur. This sensing system enabled the highly sensitive and selective detection of aqueous Hg²⁺ with a limit-of-detection of 10.2 pM.

**Keywords** Hybridization chain reaction, mercury, triple-helix molecular switch, G-quadruplex, thioflavin T

(Received May 26, 2017; Accepted August 14, 2017; Published December 10, 2017)
hybridization chain reaction (HCR) amplification strategies. Su’s group\textsuperscript{32} report a fluorescent strategy based on metal-enhanced fluorescence (MEF) and hybridization chain reaction (HCR). THMS was designed according to the principle of Watson-Crick and Hoogsteen base pairings, and composed of a target specific aptamer flanked by two arm segments and a dual-labeled oligonucleotide serving as a signal transduction probe (STP).\textsuperscript{33,34} Aptamer/target binding results in the disassembling of the THMS and releasing the STP to achieve the purpose of detection. Compared with double-helix DNA molecular switches and molecular beacon-based signaling aptamers, the THMS presents distinct advantages, including sensitivity, high stability, preserved selectivity and original aptamer affinity.\textsuperscript{35} Recently, the THMS have been designed and adopted for the fluorescence,\textsuperscript{36,37} colorimetric\textsuperscript{38} and electrochemical\textsuperscript{39} biosensing strategies.

In this work, a target specific aptamer in the above-mentioned THMS was replaced with a T-rich probe (a thymine-rich mercury-specific oligonucleotide flanked by two arm segments) for selectively binding mercury(II) ion. A dual-labeled oligonucleotide in the above-mentioned THMS was substituted for an initiator probe for triggering a hybridization event. Two hairpin probes (H1, H2), which were functionalized at 5' and 3' ends of the stem regions with three-fourths and one-fourth of the G-quadruplex respectively, were used as signal-amplification elements. In the presence of Hg\textsuperscript{2+}, the T-Hg\textsuperscript{2+}T base pairings resulted in disassembling the THMS and releasing the initiator probe. One of the hairpins was opened by the released initiator probe, which triggered a successive cross-opening of two hairpins based the strand displacement principle, resulting in the formation of long-chain DNA with multiple G-quadruplex.

Thioflavin T (ThT), a commercially available dye which could selectively bind to the G-quadruplex structure with the fluorescence,\textsuperscript{40,41} was employed as the fluorophore to design a label-free THMS-HCR-based fluorescence sensor. Because the T-T base pair can only be stabilized by Hg\textsuperscript{2+}, this platform has excellent selectivity and sensitivity for the detection of Hg\textsuperscript{2+}.

**Experimental**

**Reagents and chemicals**

Thioflavin T and Hg(NO\textsubscript{3})\textsubscript{2} were purchased from Sigma-Aldrich (St. Louis, MO). Tris-HNO\textsubscript{3} buffer (25 mM, containing 400 mM NaNO\textsubscript{3}, 20 mM KNO\textsubscript{3}, 10 mM Mg(NO\textsubscript{3})\textsubscript{2}, pH 6.5) was used for the fluorescence detection of Hg\textsuperscript{2+}. The oligonucleotides used in this work were synthesized and HPLC-purified by Sangon Biotech. Co., Ltd. (Shanghai, China) and their sequences are shown in Table S1 (Supporting Information). Cu(NO\textsubscript{3})\textsubscript{2}, Ca(NO\textsubscript{3})\textsubscript{2}, Ni(NO\textsubscript{3})\textsubscript{2}, 6H\textsubscript{2}O, Fe(NO\textsubscript{3})\textsubscript{3}, 9H\textsubscript{2}O, Zn(NO\textsubscript{3})\textsubscript{2}, 6H\textsubscript{2}O, Cr(NO\textsubscript{3})\textsubscript{3}, 9H\textsubscript{2}O, Ce(NO\textsubscript{3})\textsubscript{3}, 6H\textsubscript{2}O, Al(NO\textsubscript{3})\textsubscript{3}, 9H\textsubscript{2}O, and AgNO\textsubscript{3} were bought from Sinopharm Group Co., Ltd. (Shanghai, China). All other reagents were of analytical reagent grade. All stock solutions were prepared using ultrapure water (18.2 M\textsubscript{Ω} cm), which was obtained from Milli-Q water purification system (Millipore Corp, Bedford, MA).

**Apparatus**

All fluorescence measurements were carried out on an F-2700 fluorescence spectrophotometer (Hitachi, Japan) with excitation at 425 nm and emission at 492 nm for ThT.

**Fluorescent detection for sample solutions**

Firstly, the stock solutions of hairpin probes H1 and H2 were, respectively, heated to 95°C for 5 min and then allowed to cool down slowly to room temperature at least for 2 h to form hairpin structure before use. Meanwhile, 1 μL of the T-rich probe (50 nM final concentration) and 1 μL of the initiator probe (50 nM final concentration) were incubated in 74 μL of binding buffer (25 mM Tris-HNO\textsubscript{3}, containing 400 mM NaNO\textsubscript{3}, 10 mM Mg(NO\textsubscript{3})\textsubscript{2}, pH 6.5) for 25 min at room temperature to form triple-helix molecular switch (THMS).

Secondly, 1 μL of the sample solution to be tested (containing Hg\textsuperscript{2+} or other metal ions) was added to the above-mentioned binding buffer; the mixture was incubated at 37°C for 15 min. Then, 10 μL of H1 (200 nM final concentration), 10 μL of H2 (200 nM final concentration) and 2 μL of KNO\textsubscript{3} (20 mM final concentration) were added to the above mixture. After incubation at 37°C for 2 h, 1 μL of ThT (10 μM final concentration) was added and the resulting mixture was incubated for another 5 min. The fluorescence emission intensity for the obtained mixture was measured at 492 nm (excited at 425).

**Results and Discussion**

**Strategy for Hg\textsuperscript{2+} detection**

The designed strategy for Hg\textsuperscript{2+} detection is depicted in Scheme 1. This sensing system includes a triple-helix molecular switch (THMS), two HCR hairpin probes (H1 and H2), and thioflavin T (ThT). The THMS was designed by the Watson-Crick and Hoogsteen base pairings, and composed of a T-rich probe for the recognition of Hg\textsuperscript{2+} and an initiator probe. Hairpin 1 is functionalized at its 5’ and 3’ ends with three-fourths of the G-quadruplex (domain I) and one-fourth of the G-quadruplex (domain II) sequences respectively. Domain I is linked to sequence IV and partially hybridized with domain III. The domain III in hairpin 1, which is complementary to the initiator probe, is hybridized with parts of domain IV at the stem region, resulting in a stable hairpin configuration. Hairpin 2 includes at its 5’ and 3’ ends with one-fourth of the G-quadruplex (domain II) and three-fourths of the G-quadruplex (domain I) sequences, respectively. Sequences of domains III’ and IV’ in hairpin 2, which are partially hybridized at the stem region, are complementary to domains III and IV in hairpin 1, respectively.\textsuperscript{62} The THMS is employed as a signal trigger. In the absence of Hg\textsuperscript{2+}, the structure of THMS, H1 and H2 remain stable in solution, the selfassembly of ThT/G-quadruplex cannot put into practice due to the fact that the sequence of domain I is locked in the stem region of H1 and H2. Since the interactions between ThT and THMS, H1 and H2 are weak, the fluorescence of the background is low. In the presence of Hg\textsuperscript{2+}, the T-Hg\textsuperscript{2+}T base pairings resulted in disassembling the THMS and releasing the initiator probe. The released initiator probes hybridize with the domain III in H1, and impel H1 to be opened so that the single-stranded domain IV and the conserved three-fourths of the G-quadruplex (domain I) are released. The opened domain IV in H1 hybridizes with domain IV’ in H2, resulting that domain III’ and I in H2 are released. The released domain III’ in H2 hybridizes with domain III in the next H1, where the G-quadruplex fragments (domains I and II) will self-assemble into the G-quadruplex structure, and the next H1 will be opened. Thus, H1 and H2 are cross-opened autonomously and self-assembled into long-chain DNA incorporated with multiple G-quadruplexes, and the formation of the long-chain DNA incorporated with multiple G-quadruplexes depends on the concentration of Hg\textsuperscript{2+}. Because thioflavin T (ThT) can be bound to G-quadruplexe and its fluorescence intensity become
enhanced, the long-chain DNA with multiple G-quadruplexes will make the fluorescence intensity of ThT increase greatly. Based on this, the concentration of Hg^{2+} can be detected by monitoring the fluorescence changes of the solution. To demonstrate the feasibility of this approach, a series of experiments were carried out. As shown in Fig. 1, only H1 or H2 existed, the fluorescence intensity of the solution was low. In the absence of Hg^{2+}, THMS, H1 and H2 remained stable; the fluorescence intensity of the solution was not high. The fluorescence intensity would increase significantly after Hg^{2+} was brought into the solution. The results indicated that our proposed method could be used to detect Hg^{2+}.

Optimization of the experimental conditions

For the purpose of maximizing the performance of the Hg^{2+} assay, we investigated the effect of several parameters on the fluorescence response of the detection platform. The different stem lengths of T-rich probes, the pH of solution, the concentrations of T-rich probes, K^{+} and ThT were optimized. The difficulty level of releasing the initiator probe depends on the stem length of the T-rich probe. In order to find out the ideal stem length, different T-rich probes with the stem lengths of 6 to 9 bases were investigated. As shown in Fig. 2, the fluorescent signal ratio \((F - F_0)/F_0\) was plotted as a function of T-rich probes. The optimal K^{+} ions concentration was chosen to be 20 mM. The pH is a crucial parameter, which is vital to the stability of the
Hoogsteen base pairing to form a stable THMS structure.\textsuperscript{31} Figure S2 (Supporting Information) displays the fluorescence intensity ratio of the sensing platform in different pH solution. It was observed that the fluorescent ratio rose gradually with the increase of pH and then reached a peak at 6.5. Thus, the optimal pH was chosen to be 6.5. As shown in Fig. S3 (Supporting Information), the fluorescent response was increased with the increase of T-rich probe 3 concentrations and reached a plateau at 50 nM of T-rich probe 3, while keeping T-rich probe 3 and initiator probes with equal proportion. The optimal T-rich probe 3 concentration was chosen to be 50 nM. The concentration of ThT can also affect the sensing performance. The result in Fig. S4 (Supporting Information) shows that the fluorescent response enhanced with the increase of the ThT concentration and reached a plateau at 10 $\mu$M of ThT. The optimal concentration of ThT was chosen to be 10 $\mu$M.

**Sensitivity and selectivity**

Under the optimal conditions, we incubated the THMS-HCR sensing platform with various concentrations of Hg$^{2+}$ (0 - 100 nM). As shown in Fig. 3a, the fluorescence intensity increased with the increasing concentration of Hg$^{2+}$. It can be observed in Fig. 3b that the fluorescence enhancement of the sensing platform was proportional to the concentration of Hg$^{2+}$.

A linear correlation existed between the fluorescence intensity ratio ($\frac{(F - F_0)}{F_0}$) and the concentration of Hg$^{2+}$ in the range of 0.02 to 1 nM. The regression equation was $y = 0.872x + 0.453$ with a correlation coefficient of 0.993, the limit of detection (LOD) was 10.2 pM, which was relatively low comparable to the recently reported assays (Table S2, Supporting Information).

To demonstrate the selectivity of the proposed assay, a series of metal ions (Cu$^{2+}$, Ca$^{2+}$, Ni$^{2+}$, Fe$^{3+}$, Zn$^{2+}$, Cr$^{3+}$, Ce$^{3+}$, Al$^{3+}$, Ag$^{+}$, each at 200 nM) were investigated by this method. As shown in Fig. 4, although the concentrations of Hg$^{2+}$ were 20-times lower than other metal ions respectively, the fluorescence intensity ratio of Hg$^{2+}$ was significantly higher than other metal ions. The result indicated excellent selectivity of the presented sensing platform for detection of Hg$^{2+}$.

**Practical application**

To investigate the capability of this method for the determination of Hg$^{2+}$ in real samples, different concentrations of Hg$^{2+}$ spiked in tap water were further analyzed. Before detection, the tap water was briefly pretreated. Firstly, the tap water was heated and boiled for 15 min. Then, the tap water sample was filtered through a 0.22-$\mu$m membrane after the tap water cooled to room temperature. Finally, the processed tap-water sample was diluted 10 fold with 25 mM Tris-HNO$_3$. The treated tap-water sample was firstly analyzed by an atomic absorption spectrophotometer, and our proposed method, respectively; both results indicated that no mercury ions were detected. And then different amounts of Hg$^{2+}$ were spiked into the sample. The recovery rates for spiked Hg$^{2+}$ was measured by our proposed method. The results given in Table 1 show that the recovery rates were in the range of 98.13 to 107.01%, indicating that this biosensor was both reliable and practical.

<table>
<thead>
<tr>
<th>Tap water</th>
<th>Added/nM</th>
<th>Found/nM (mean ± SD)$^a$</th>
<th>Recovery, %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sample 1</td>
<td>0.1</td>
<td>0.10701 ± 0.0427</td>
<td>107.01</td>
</tr>
<tr>
<td>Sample 2</td>
<td>0.5</td>
<td>0.50037 ± 0.0491</td>
<td>100.07</td>
</tr>
<tr>
<td>Sample 3</td>
<td>1.0</td>
<td>0.98135 ± 0.0109</td>
<td>98.13</td>
</tr>
</tbody>
</table>

$^a$ Mean values and SDs were obtained from three independent experiments.

Fig. 3 (a) Fluorescence responses in the presence of different concentrations of Hg$^{2+}$. From bottom to top: 0, 0.02, 0.1, 0.2, 0.4, 0.6, 1.0, 5.0, 10.0, 20.0, 50.0, 100.0 nM Hg$^{2+}$. (b) Relative fluorescence intensity ratio of the THMS-HCR sensing platform in the presence of different concentrations of Hg$^{2+}$. The inset presents a linear relationship in the concentrations of Hg$^{2+}$ ranging from 0.02 to 1 nM.

Fig. 4 Selectivity of the Hg$^{2+}$-sensing. The concentration of Hg$^{2+}$ is 10 nM and other metal ion is 200 nM. Each data were performed in triplicate.
Conclusions

In order to detect Hg^{2+}, a novel label-free and highly sensitive fluorescence assay depending on the triple-helix molecular switch (THMS)-induced hybridization chain reaction (HCR) amplification was developed. Two hairpin probes containing the G-quadruplex sequence were used as signal-amplification elements. THMS is composed of a T-rich mercury-specific probe and an initiator probe, designed according to the principle of Watson-Crick and Hoogsteen base pairings. Hg^{2+} disassembles the THMS and releases the initiator probe, which triggered a successive cross-opening of two hairpins, resulting in the formation of long-chain DNA with multiple G-quadruplex.

The assay showed a high selectivity to Hg^{2+} with a detection limit of 10.2 pM. Instead of requiring complex experimental conditions, this THMS-HCR sensing platform could be generalized for the detection of Hg^{2+} ions in a wide range of biological, toxicological, and environmental samples.

Acknowledgements

This work was supported by National Natural Science Foundation of China (No. 21327009) and Scientific Research Fund of Hunan Provincial Education Department (Nos. 14A012 and 17C0033).

Supporting Information

Sequences of DNA used in this work, comparison of analytical methods for the detection of Hg^{2+} and the figures of optimized experimental conditions were listed in Supporting Information. This material is available free of charge on the Web at http://www.jsac.or.jp/analsci/.

References