


Cite this: *RSC Adv.*, 2024, 14, 38279

Received 7th November 2024  
Accepted 27th November 2024

DOI: 10.1039/d4ra07922a

rsc.li/rsc-advances

# Organomercury oligonucleotide–polydopamine nanoparticle assemblies discriminate between target sequences by Hg(II)-mediated base pairing†

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A fluorescently tagged oligonucleotide hybridization probe incorporating a single 5-mercurocytosine residue was synthesized and found to adsorb on polydopamine nanoparticles much more strongly than its unmetallated counterpart. Hybridization with target sequences led to release of the probe from the nanoparticle to varying degrees depending on the nucleobase opposite to 5-mercurocytosine.

## 1. Introduction

Nanomaterials that exhibit fluorescence emitting or quenching properties are crucial for the advancement of molecular probes for the detection and investigation of nucleic acids.<sup>1–6</sup> The development of this class of fluorescent sensors for assaying biomolecules has recently received considerable attention due to their inherent advantages, such as high sensitivity, operational convenience and, especially, *in situ* imaging properties. The integration of oligonucleotides with nanoparticles has emerged as a promising avenue, offering remarkable opportunities for the design of advanced sensor architectures. Several nanomaterials, such as gold nanoparticles (AuNPs),<sup>7–9</sup> carbon nanotubes (CNTs),<sup>10</sup> graphene oxide,<sup>11,12</sup> metal–organic frameworks (MOFs),<sup>13</sup> MoS<sub>2</sub> nanosheets,<sup>14</sup> and polydopamine nanoparticles (PDANs)<sup>15,16</sup> have been utilized as quenchers in the development of fluorescent sensors.<sup>15–17</sup> Although considerable progress has been made in the use of nanomaterials in nucleic acid detection, several challenges remain in controlling the assembly and fluorescence emission properties, along with issues like background interference and salt-induced aggregation, which limit detection capabilities.

Among functional nanomaterials, PDANs have gained much attention in drug delivery, therapy, and biosensing due to their biocompatibility and surface-coating properties.<sup>18–20</sup> Like the majority of the nanomaterials discussed above, PDANs carry negative charges at neutral pH and, hence, repel the polyanionic phosphate backbone of DNA. The electrostatic repulsion can be alleviated by lowering the pH or masking the negative charges by increasing the electrolyte concentration. On the other hand, PDANs are capable of  $\pi$ – $\pi$  stacking and

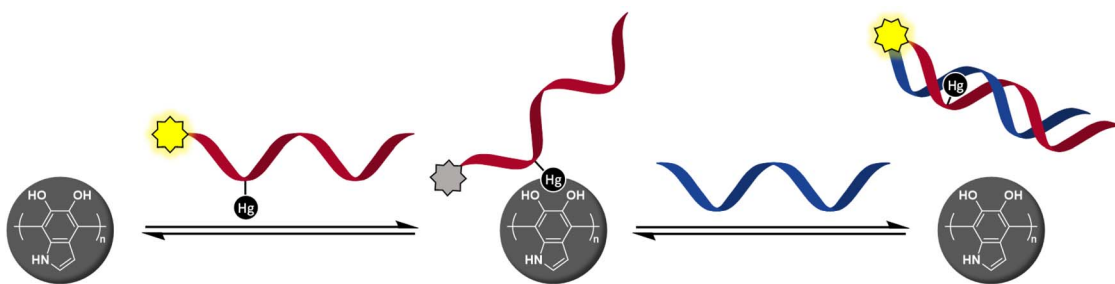
hydrogen bonding, both of which promote adsorption.<sup>15</sup> Furthermore, the catecholamine moieties of PDANs are strong ligands for various metal ions,<sup>21–23</sup> allowing much more efficient adsorption through metal coordination. Such metal-mediated adsorption of DNA has been harnessed to create DNA/PDAN sensor assemblies for the detection of nucleic acids in biological media.<sup>16</sup>

Our previous studies on organomercury oligonucleotides as hybridization probes<sup>24–28</sup> have revealed an increased hybridization affinity and in some cases robust discrimination of all the canonical bases, 2-thiothymine and 4-thiothymine through metal-mediated base pairing.<sup>29–34</sup> The state-of-the-art methods for the identification of single-nucleotide polymorphisms (SNPs) require as many as four different hybridization probes interrogating the polymorphic base through Watson–Crick base pairing.<sup>35,36</sup> In contrast, we have been able to obtain clearly different UV melting temperatures<sup>24,26,28</sup> or fluorescence signals<sup>25</sup> for all canonical nucleobases using only a single organomercury hybridization probe. After learning of the success of metal-mediated adsorption in developing DNA/PDAN sensors,<sup>16</sup> it occurred to us that the covalently bound Hg(II) ions of organomercury oligonucleotides<sup>37</sup> might also exhibit sufficient affinity to the catecholamine ligands on the surface of PDANs to promote adsorption (Scheme 1). On hybridization of the organomercury oligonucleotide with a complementary sequence, the Hg(II) ion would become engaged in Hg(II)-mediated base pairing and thus be no longer available for catecholamine coordination, leading to dissociation of the resulting duplex from the PDAN. As an important advantage over the previously reported DNA/PDAN sensors based on coordinative interactions with metal ions, the organomercury oligonucleotide remains undissociated even at high dilution in aqueous media, as exemplified by numerous prior studies.<sup>37,38</sup> The present article describes the preparation and testing of a novel sensor based on this reasoning.

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† Electronic supplementary information (ESI) available. See DOI: <https://doi.org/10.1039/d4ra07922a>





Scheme 1 Hg(II)-promoted adsorption of a fluorescently tagged oligonucleotide probe quenches emission and hybridization with a target sequence restores it.

## 2. Results and discussion

### 2.1 Synthesis of the polydopamine nanoparticles

The PDANs were synthesized according to a well-established literature procedure, which involves an oxidative self-polymerization of dopamine.<sup>39</sup> The size of the PDANs was determined by dynamic light scattering (DLS) and transmission electron microscopy (TEM) (Fig. S5 and S6 in the ESI†). An average diameter of approximately 170 nm was obtained by both methods.

### 2.2 Oligonucleotide synthesis

For the sensing component, oligonucleotide **ON1c-Hg**, bearing a fluorescent tag and a single 5-mercurycytosine residue, was synthesized. The mercurated nucleobase was positioned seven nucleotides away from the 5'-terminal fluorophore, sufficiently far to prevent direct interaction, but at the same time, sufficiently near so that Hg(II)-mediated adsorption on the PDAN would cause a quenching of the fluorescence. The 24-mer sequence (Table 1) of **ON1c-Hg** was first assembled by the conventional phosphoramidite strategy on an automated synthesizer. A 6-bromohexyl group<sup>25</sup> was introduced at the 5'-end for subsequent coupling of the fluorophore. To prevent unwanted mercuration, all cytosine building blocks were replaced with 5-methylcytosines, except for the single residue designated for mercuration. Additionally, the order of the post-synthetic transformations had to be planned carefully to

prevent mercuration of the FAM-6 fluorophore. Accordingly, after the synthesis of the 5'-bromo oligonucleotide sequence, the 5'-bromide was displaced by an azide group by treatment with tetramethylguanidinium azide (TMGA) on solid support (Scheme 2). The resulting 5'-azide oligonucleotide intermediate was then cleaved from the solid support with aqueous ammonia and purified by RP-HPLC.

Mercuration of the azide oligonucleotide intermediate was accomplished by treatment with aqueous mercuric acetate at 55 °C for 48 h. The product mixture was fractionated by RP-HPLC and identity of the desired oligonucleotide intermediate was confirmed by mass spectrometry. Finally, the fluorophore was introduced by strain-promoted azide-alkyne cycloaddition (SPAAC) of the 6-isomer of FAM-DBCO with the 5'-azide-functionalized oligonucleotide. The unmercurated reference oligonucleotide **ON1c** was synthesized in the same way except that the mercuration step was omitted. The complementary oligonucleotides **ON2a**, **ON2c**, **ON2g**, **ON2t**, **ON2s<sup>2</sup>t** and **ON2s<sup>4</sup>t** (Table 1), in turn, were synthesized from commercially available phosphoramidite building blocks following recommended procedures. The final products were purified by RP-HPLC (Fig. S1 and S2 in the ESI†), characterized mass spectrometrically (Fig. S3 and S4 in the ESI†) and quantified UV spectrophotometrically.

### 2.3 Affinity of the oligonucleotide hybridization probes for the polydopamine nanoparticles

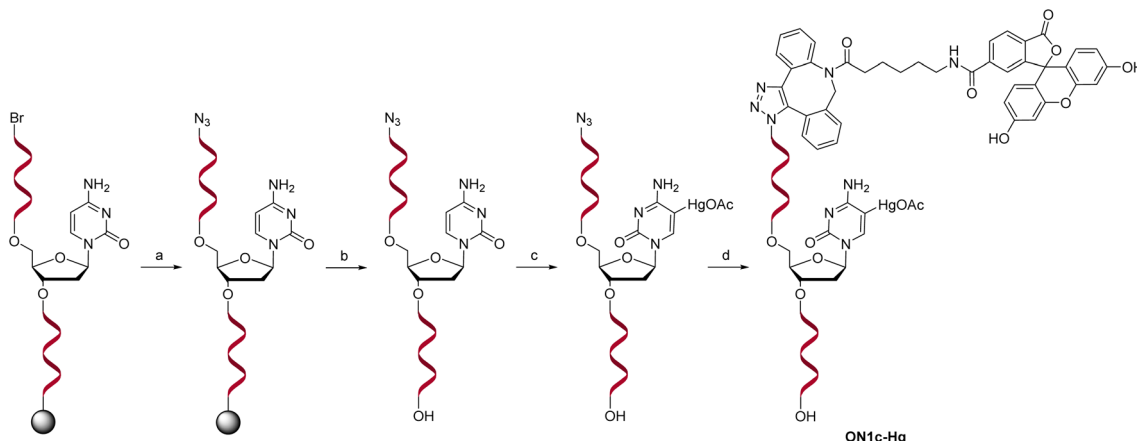
While inorganic Hg(II) shows high affinity for polydopamine,<sup>40</sup> it was not certain that a single 5-mercurycytosine residue would suffice to bind the oligonucleotide probe to the PDAN under conditions in which the unmetallated nucleic acids remain unbound. Therefore, we compared the PDAN-binding affinity of the mercurated oligonucleotide **ON1c-Hg** and its unmercurated counterpart **ON1c** by measuring their fluorescence emission at increasing PDAN concentrations. As anticipated, both oligonucleotides exhibited high fluorescence emission at 517 nm in the absence of PDANs. The fluorescence intensity decreased on increasing PDAN concentration but the profiles obtained for **ON1c-Hg** and **ON1c** (Fig. 1) were very different. With **ON1c**, the decrease was linear, suggesting a low binding affinity. Almost 90% of the emission persisted even at the highest PDAN concentration used. In contrast, with **ON1c-Hg**, fluorescence emission initially decreased steeply with increasing PDAN

Table 1 Sequences of the oligonucleotides used in this study

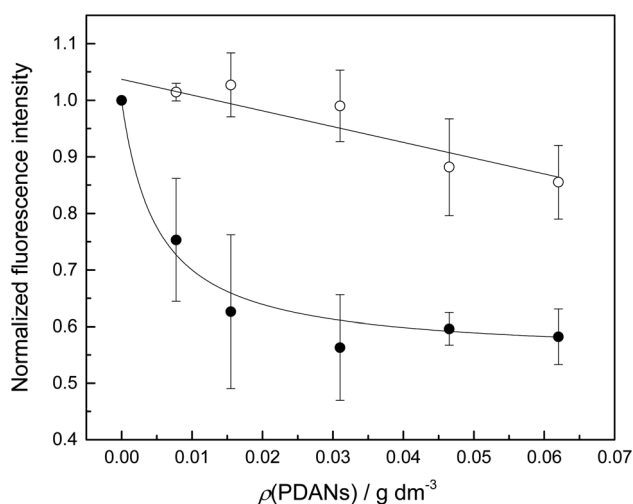
Oligonucleotide	Sequence (5'–3') <sup>a</sup>
<b>ON1c</b>	6-FAM-AC <sup>m</sup> G C <sup>m</sup> AT CTG TGA AGA GAA C <sup>m</sup> C <sup>m</sup> T GGG
<b>ON1c-Hg</b>	6-FAM-AC <sup>m</sup> G C <sup>m</sup> AT C <sup>Hg</sup> TG TGA AGA GAA C <sup>m</sup> C <sup>m</sup> T GGG
<b>ON2a</b>	CCC AGG TTC TCT TCA CAA ATG CGT
<b>ON2c</b>	CCC AGG TTC TCT TCA CAC ATG CGT
<b>ON2g</b>	CCC AGG TTC TCT TCA CAG ATG CGT
<b>ON2t</b>	CCC AGG TTC TCT TCA CAT ATG CGT
<b>ON2s<sup>2</sup>t</b>	CCC AGG TTC TCT TCA CAs <sup>2</sup> T ATG CGT
<b>ON2s<sup>4</sup>t</b>	CCC AGG TTC TCT TCA CAs <sup>4</sup> T ATG CGT

<sup>a</sup> C<sup>m</sup> refers to 5-methylcytosine, C<sup>Hg</sup> to 5-mercurycytosine, s<sup>2</sup>T to 2-thiothymine and s<sup>4</sup>T to 4-thiothymine. In each sequence, the residue varied in the hybridization experiments has been underlined.





**Scheme 2** Post-synthetic transformations to afford the mercurate probe oligonucleotide **ON1c-Hg**. Reagents and conditions: (a) TMGA, NaI, DMF, 60 °C, 90 min; (b)  $\text{NH}_3$ ,  $\text{H}_2\text{O}$ , 55 °C, 16 h; (c)  $\text{Hg}(\text{OAc})_2$ ,  $\text{H}_2\text{O}$ , 55 °C, (d) 48 h; 6-FAM-DBCO, DMF, 25 °C, 16 h.



**Fig. 1** Fluorescence emission of oligonucleotides **ON1c** (○) and **ON1c-Hg** (●) as a function of PDAN concentration;  $T = 25\text{ °C}$ ;  $\text{pH} = 7.4$  (20 mM cacodylate buffer);  $I(\text{NaClO}_4) = 0.10\text{ M}$ ; [oligonucleotides] = 10 nM;  $\lambda_{\text{ex}} = 493\text{ nm}$ ;  $\lambda_{\text{em}} = 517\text{ nm}$ .

concentration before reaching a plateau. Assuming that the number of binding sites on the surface of the PDANs greatly exceeds the number of mercurated oligonucleotides in the sample, the dependence of fluorescence intensity on PDAN concentration can be expressed by eqn (1).

$$I = I_{\infty} + (I_0 - I_{\infty}) \frac{1}{1 + K\rho} \quad (1)$$

$I_0$  and  $I_{\infty}$  are fluorescence intensities in the absence and at an infinite concentration of PDANs,  $K$  is the adsorption affinity constant and  $\rho$  the mass concentration of the PDANs. Values of  $I_{\infty} = 0.55 \pm 0.02$  and  $K = 200 \pm 70\text{ g}^{-1}\text{ dm}^3$  were obtained by non-linear least-squares fitting, whereas  $I_0$  was defined as = 1. The residual emission at high PDAN concentration probably stems from flexibility of the single-stranded oligonucleotide, allowing the fluorophore to spend some time away from the

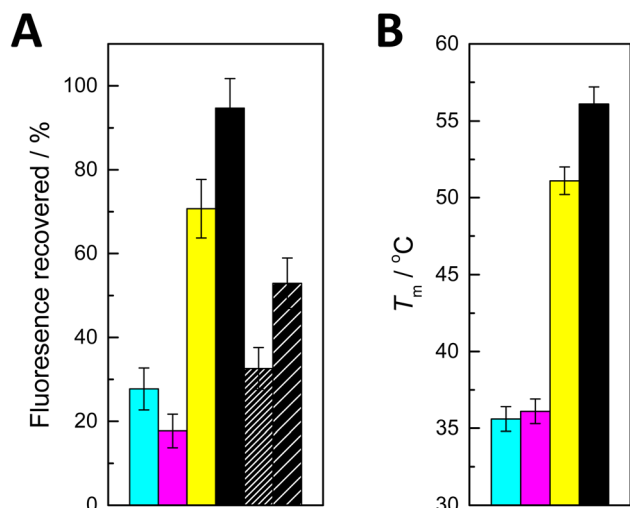
PDAN even when the oligonucleotide is adsorbed. Incomplete adsorption of **ON1c-Hg** appears a less likely explanation given the observed saturation at high PDAN concentration. The relatively high affinity of the mercurated probe **ON1c-Hg** (compared to its metal-free counterpart) for the PDANs, probably attributable to mercury coordination by the catechol moieties, was encouraging from the point of view of the intended sensor application. In contrast, the low affinity of the unmercurated probe **ON1c** precludes a meaningful analysis.

#### 2.4 Sequence selectivity of the polydopamine nanoparticle—oligonucleotide assemblies

For assessing the potential of the PDAN—**ON1c-Hg** assembly as a sensor, the PDAN concentration was fixed at the value where the difference in the fluorescence intensity of **ON1c** and **ON1c-Hg** peaked ( $0.016\text{ mg dm}^{-3}$ ), while keeping the other parameters the same as in the affinity measurements. **ON1c-Hg** and the PDANs were first incubated for approximately 10 min, after which an equimolar amount of one of the complementary oligonucleotides **ON2a**, **ON2c**, **ON2g**, **ON2t**, **ON2s<sup>2</sup>t** and **ON2s<sup>4</sup>t** was added. Hybridization of **ON1c-Hg** with these oligonucleotides places each of the canonical nucleobases, as well as 2- and 4-thiothymine, opposite to the 5-mercuricytosine residue. With **ON2a**, fluorescence emission was recorded 10 min, 20 min and 24 h after the addition and found to be unchanged. All subsequent measurements were performed 10 min after addition of the complementary oligonucleotide. Recovery of the fluorescence, expressed by eqn (2), was observed in each case, indicating duplex formation and subsequent dissociation from the PDAN.

$$\text{Fluorescence recovered} = 100\% \frac{I - I_{\infty}}{I_0 - I_{\infty}} \quad (2)$$

$I$  is the fluorescence intensity recorded after addition of the complementary oligonucleotide and  $I_0$  and  $I_{\infty}$  are the limits determined during the affinity measurements. The values obtained are summarized in Fig. 2A.



**Fig. 2** (A) Recovery of fluorescence on addition of **ON2a** (cyan), **ON2c** (magenta), **ON2g** (yellow), **ON2t** (black), **ON2s<sup>2</sup>t** (black with dense hash) and **ON2s<sup>4</sup>t** (black with sparse hash) to PDAN-bound **ON1c-Hg**;  $T = 25\text{ }^{\circ}\text{C}$ ;  $\text{pH} = 7.4$  (20 mM cacodylate buffer);  $[\text{NaClO}_4] = 0.10\text{ M}$ ;  $[\text{oligonucleotides}] = 10\text{ nM}$ ;  $\rho(\text{PDAN}) = 0.016\text{ mg dm}^{-3}$ ;  $\lambda_{\text{ex}} = 493\text{ nm}$ ;  $\lambda_{\text{em}} = 517\text{ nm}$ . (B) Previously reported<sup>27</sup> UV melting temperatures of double-helical 11-mer DNA oligonucleotides incorporating a Hg(II)-mediated base pair between 5-mercurocytosine and adenine (cyan), cytosine (magenta), guanine (yellow) and thymine (black).

The effects of adding the oligonucleotides consisting solely of canonical nucleobases fell into two distinct categories. With **ON2a** and **ON2c**, recovery of fluorescence was modest (28 and 18%, respectively). In contrast, addition of **ON2g** and **ON2t** led to 71 and 95% recovery of the fluorescence, respectively. These results are in good agreement with the previously reported<sup>27</sup> thermal stabilities of corresponding short duplexes (Fig. 2B) and with the tendency of Hg(II) to displace a proton from the N1 of guanine and the N3 of thymine upon coordination. The differences in fluorescence recovery were sufficiently large for a robust discrimination between most of the canonical nucleobases. The sole exception was differentiating adenine from cytosine, which could not be achieved reliably with the present system.

Curiously, when either of the thiopyrimidine sequences **ON2s<sup>2</sup>t** and **ON2s<sup>4</sup>t** was added, the fluorescence recovery was lower compared to **ON2t** and **ON2g**, but higher compared to **ON2a** and **ON2c**. This result may seem unexpected given the high stabilities previously reported for duplexes involving Hg(II)-mediated base pairing with 2- and 4-thiothymine.<sup>28</sup> It should be pointed out, however, that in those cases the Hg(II) ion was offset from the centerline of the base pair, allowing it to coordinate to an exocyclic thio substituent with only minor distortion of the geometry. In the present case, coordination of the Hg(II) to an exocyclic thio substituent would lead to a considerable shear with either isomer of thiothymine.

### 3. Conclusion

In summary, a single 5-mercurocytosine residue allows efficient adsorption of oligonucleotide hybridization probes on PDANs

under conditions where respective metal-free oligonucleotides remain largely unabsorbed, in all likelihood through coordination of the Hg(II) ion to a catecholamine moiety. Addition of a complementary target oligonucleotide leads to partial dissociation of the probe and the equilibrium reflects differences in the strength of Hg(II)-mediated base pairing between 5-mercurocytosine and the nucleobase opposite to it within the target sequence. These results demonstrate the potential of organometallic oligonucleotide–PDAN assemblies as novel sensors for SNP genotyping. The advantages of our method include a simple and rapid procedure and the utilization of affordable and biodegradable nanoparticles.

## 4. Experimental section

### 4.1 General experimental methods

HPLC elution buffers (pH 7.0) were prepared using freshly distilled analytical grade triethylamine ( $\text{Et}_3\text{N}$ ). All other reagents and solvents were obtained commercially and used without further purification. Mass spectra were recorded using a Waters ACQUITY RDa mass spectrometer, UV spectra on a Jenway Genova Nano microvolume UV/vis spectrophotometer and dynamic light scattering (DLS) spectra on a Zetasizer Nano 90 (Malvern) instrument. Transmission electron microscopy (TEM) images were captured with a JEM-1400 Plus TEM instrument. Fluorescence measurements were performed on an FLS 1000 (Edinburgh instruments) fluorescence spectrometer. X-ray photoelectron spectroscopy (XPS) measurements were conducted using a Nexsa XPS instrument (Thermo Scientific). Fourier-transform infrared spectroscopy (FTIR) was carried out using a Vertex 70 instrument (Bruker).

### 4.2 Oligonucleotide synthesis

Oligonucleotides **ON1c**, **ON1c-Hg**, **ON2a**, **ON2c**, **ON2g**, **ON2t**, **ON2s<sup>2</sup>t** and **ON2s<sup>4</sup>t** were synthesized using an ÅKTA oligopilot plus 10 DNA/RNA synthesizer on a controlled pore glass (CPG) support, employing the conventional phosphoramidite method with 5-(benzylthio)-1H-tetrazole as the activator and the previously reported<sup>25</sup> 6-bromohexyl phosphoramidite building block for introduction of the 5'-bromo group. The trityl response indicated that all coupling reactions achieved near-quantitative efficiency. Following synthesis, the support-bound oligonucleotides **ON1c**, **ON1c-Hg**, **ON2a**, **ON2c**, **ON2g**, **ON2t** and **ON2s<sup>2</sup>t** were treated with 25% aqueous  $\text{NH}_3$  at 55 °C for 16 h to cleave the linker and to remove the phosphate and base protecting groups. With **ON2s<sup>4</sup>t**, the cyanoethyl protecting groups of the phosphate linkages were removed first by treating with 1.0 M DBU in anhydrous acetonitrile at 25 °C for 2 h. This treatment was followed by full deprotection using 50 mM NaSH in 25% aqueous  $\text{NH}_3$  at 25 °C for 24 h. The manufacturer of the 4-thiothymidine phosphoramidite building block recommends this protocol to prevent ammonolysis of the S-cyanoethyl group. The crude oligonucleotides were purified by reverse-phase high-performance liquid chromatography (RP-HPLC) employing a Hypersil ODS C18 column (250 × 4.6 mm, 5  $\mu\text{m}$ ) and a Bio-Zen™ Oligo column (150 × 4.6 mm, 2.6  $\mu\text{m}$ ). A linear gradient





(5–45% over 20 min) of MeCN in 50 mM aqueous triethylammonium acetate was employed, with flow rates of 1.0 mL min<sup>−1</sup> for the Hypersil ODS C18 column and 0.6 mL min<sup>−1</sup> for the BioZen Oligo column.

For introduction of the fluorescent tag to **ON1c-Hg** and **ON1c**, the 5'-bromo group was converted to an azido group. This conversion was performed on solid support within the synthesis column by adding tetramethylguanidinium azide (TMGA) (32 mg, 100 eq.) and NaI (30 mg, 100 eq.) in dry DMF (2 mL), followed by heating the mixture at 60 °C for 90 min. The reaction mixture was periodically agitated by manually transferring the solution between two syringes. Subsequently, the support-bound oligonucleotides were washed sequentially with DMF, DCM, H<sub>2</sub>O and MeOH and dried under vacuum. Finally, the oligonucleotides were cleaved from the solid support and purified by RP-HPLC on a Hypersil ODS C18 column.

A portion (30 nmol) of the 5'-azido oligonucleotide was incubated with Hg(OAc)<sub>2</sub> (10 μmol) in 100 μL of H<sub>2</sub>O at 55 °C for 48 h. The resulting mercurated oligonucleotide was purified by RP-HPLC on a Hypersil ODS C18 column. Subsequently, the mercurated 5'-azido oligonucleotide was coupled to fluorescein-DBCO by strain-promoted alkyne-azide cycloaddition (SPAAC). For this, FAM-DBCO (13.5 mg) was dissolved in DMF (200 μL) and two equivalents (0.3 μL) were added to an aqueous solution of the oligonucleotide (15 nmol). The reaction mixture was vortexed at room temperature for 16 h, followed by RP-HPLC purification on a BioZen™ Oligo column (Fig. S2 in the ESI†). The rest of the 5'-azido oligonucleotide (30 nmol) was conjugated with fluorescein-DBCO and the crude product purified under the same conditions (Fig. S1 in the ESI†). Finally, both of the purified products (**ON1c** and **ON1c-Hg**) were characterized by mass spectrometry (Fig. S3 and S4 in the ESI†) and quantified by UV spectrophotometry.

#### 4.3 PDAN synthesis and characterization

A solution containing 200 mg of dopamine in 100 mL of freshly prepared 8.5 mM aqueous NaOH was vigorously stirred at 50 °C in a well-ventilated fume hood. After 5 h reaction, the resulting black suspension was cooled and centrifuged at 9300 rpm for 20 min. The supernatant was removed and the pellet resuspended in H<sub>2</sub>O. The washing and centrifugation procedure was repeated four times, after which the PDANs were redispersed in H<sub>2</sub>O for subsequent experiments. The dynamic size and morphology of the PDANs were determined by DLS Zetasizer Nano 90 (Malvern) and TEM, respectively (Fig. S5 and S6 in the ESI†). Additionally, the synthesized PDANs were subjected to characterization *via* FTIR and XPS (Fig. S7 and S8 in the ESI†).

#### 4.4 Fluorescence measurements

Fluorescence emission spectra were obtained over a range of 500–700 nm using an excitation wavelength of 493 nm and a dwell time of 0.2 seconds per data point. The samples consisted of 10 nM probe oligonucleotide **ON1c** or **ON1c-Hg**, 20 mM cacodylate buffer, maintained at pH 7.4 with an ionic strength of 0.10 M (adjusted with NaClO<sub>4</sub>). With the mercurated probe **ON1c-Hg**, emission spectra were recorded both before and after

addition of a 10 nM concentration of each one of the complementary strands **ON2a**, **ON2c**, **ON2g**, **ON2t**, **ON2s<sup>2</sup>t** and **ON2s<sup>4</sup>t**.

## Data availability

The data supporting this article have been included as part of the ESI.†

## Author contribution

T. K. K.: conceptualization, data curation, formal analysis, funding acquisition, investigation, methodology, validation, visualization, writing – original draft, writing – review and editing; M. A.-W.: investigation, methodology, writing – review and editing; J. L.: methodology, supervision, writing – review and editing; T. L.: data curation, formal analysis, project administration, supervision, visualization, writing – review and editing.

## Conflicts of interest

The authors report no conflict of interest.

## Acknowledgements

The kind help of Mr Kari Eränen (Åbo Akademi University) with the DLS experiments is gratefully recognized.

## References

- 1 S. Song, Y. Qin, Y. He, Q. Huang, C. Fan and H. Y. Chen, *Chem. Soc. Rev.*, 2010, **39**, 4234–4243.
- 2 H. Wang, R. Yang, L. Yang and W. Tan, *ACS Nano*, 2009, **3**, 2451–2460.
- 3 J. J. Xu, W. W. Zhao, S. Song, C. Fan and H. Y. Chen, *Chem. Soc. Rev.*, 2014, **43**, 1601–1611.
- 4 Z. Lin, G. Wu, L. Zhao and K. W. C. Lai, *IEEE Nanotechnol. Mag.*, 2019, **13**, 4–14.
- 5 H. S. Wang, *Coord. Chem. Rev.*, 2017, **349**, 139–155.
- 6 H. Zhang, H. Zhang, A. Aldalbahi, X. Zuo, C. Fan and X. Mi, *Biosens. Bioelectron.*, 2017, **89**, 96–106.
- 7 D. J. Maxwell, J. R. Taylor and S. Nie, *J. Am. Chem. Soc.*, 2002, **124**, 9606–9612.
- 8 B. Dubertret, M. Calame and A. J. Libchaber, *Nat. Biotechnol.*, 2001, **19**, 365–370.
- 9 H. Li and L. J. Rothberg, *Anal. Chem.*, 2004, **76**, 5414–5417.
- 10 R. Yang, J. Jin, Y. Chen, N. Shao, H. Kang, Z. Xiao, Z. Tang, Y. Wu, Z. Zhu and W. Tan, *J. Am. Chem. Soc.*, 2008, **130**, 8351–8358.
- 11 C. H. Lu, H. H. Yang, C. L. Zhu, X. Chen and G. N. Chen, *Angew. Chem., Int. Ed.*, 2009, **48**, 4785–4787.
- 12 S. Li, A. N. Aphale, I. G. MacWan, P. K. Patra, W. G. Gonzalez, J. Miksovska and R. M. Leblanc, *ACS Appl. Mater. Interfaces*, 2012, **4**, 7069–7075.
- 13 X. Zhu, H. Zheng, X. Wei, Z. Lin, L. Guo, B. Qiu and G. Chen, *Chem. Commun.*, 2013, **49**, 1276–1278.



- 14 C. Zhu, Z. Zeng, H. Li, F. Li, C. Fan and H. Zhang, *J. Am. Chem. Soc.*, 2013, **135**, 5998–6001.
- 15 W. Qiang, W. Li, X. Li, X. Chen and D. Xu, *Chem. Sci.*, 2014, **5**, 3018–3024.
- 16 Y. Meng, P. Liu, W. Zhou, J. Ding and J. Liu, *ACS Nano*, 2018, **12**, 9070–9080.
- 17 G. Bülbül, A. Hayat, F. Mustafa and S. Andreescu, *Sci. Rep.*, 2018, **8**, 1–9.
- 18 M. Liu, G. Zeng, K. Wang, Q. Wan, L. Tao, X. Zhang and Y. Wei, *Nanoscale*, 2016, **8**, 16819–16840.
- 19 Y. Liu, K. Ai and L. Lu, *Chem. Rev.*, 2014, **114**, 5057–5115.
- 20 M. E. Lynge, R. Van Der Westen, A. Postma and B. Städler, *Nanoscale*, 2011, **3**, 4916–4928.
- 21 N. Holten-Andersen, M. J. Harrington, H. Birkedal, B. P. Lee, P. B. Messersmith, K. Y. C. Lee and J. H. Waite, *Proc. Natl. Acad. Sci. U.S.A.*, 2011, **108**, 2651–2655.
- 22 X. Li, P. Gao, J. Tan, K. Xiong, M. F. Maitz, C. Pan, H. Wu, Y. Chen, Z. Yang and N. Huang, *ACS Appl. Mater. Interfaces*, 2018, **10**, 40844–40853.
- 23 H. Zeng, D. S. Hwang, J. N. Israelachvili and J. H. Waite, *Proc. Natl. Acad. Sci. U.S.A.*, 2010, **107**, 12850–12853.
- 24 A. Aro-Heinilä, T. Lönnberg and P. Virta, *Bioconjugate Chem.*, 2019, **30**, 2183–2190.
- 25 A. Aro-Heinilä, T. Lönnberg and P. M. Virta, *ChemBioChem*, 2020, **22**, 354–358.
- 26 A. Aro-Heinilä, A. Lepistö, A. Äärelä, T. A. Lönnberg and P. Virta, *J. Org. Chem.*, 2022, **87**, 137–146.
- 27 D. Ukale, V. S. Shinde and T. Lönnberg, *Chem.–Eur. J.*, 2016, **22**, 7917–7923.
- 28 T. K. Kotammagari, P. Tähtinen and T. Lönnberg, *Chem.–Eur. J.*, 2022, **28**, e202202530.
- 29 M. Hebenbrock and J. Müller, *Comprehensive Inorganic Chemistry III*, 3rd edn, 2023, vol. 1–10, pp. 664–713.
- 30 S. Naskar, R. Guha and J. Müller, *Angew. Chem., Int. Ed.*, 2020, **59**, 1397–1406.
- 31 J. Müller, *Coord. Chem. Rev.*, 2019, **393**, 37–47.
- 32 B. Jash and J. Müller, *Chem.–Eur. J.*, 2017, **23**, 17166–17178.
- 33 Y. Takezawa, J. Müller and M. Shionoya, *Chem. Lett.*, 2017, **46**, 622–633.
- 34 T. Dairaku, K. Furuita, H. Sato, J. Sebera, K. Nakashima, A. Ono, V. Sychrovsky, C. Kojima and Y. Tanaka, *Inorg. Chim. Acta*, 2016, **452**, 34–42.
- 35 D. M. Kolpashchikov, *Acc. Chem. Res.*, 2019, **52**, 1949–1956.
- 36 J. X. Chen, C. Shi, X. Y. Kang, X. T. Shen, X. Z. Lao and H. Zheng, *Anal. Methods*, 2020, **12**, 884–893.
- 37 D. Ukale and T. Lönnberg, *ChemBioChem*, 2021, **22**, 1733–1739.
- 38 T. K. Kotammagari, L. Y. Saleh and T. Lönnberg, *Chem. Commun.*, 2024, **60**, 3118–3128.
- 39 M. Zandieh and J. Liu, *Langmuir*, 2021, **37**, 8953–8960.
- 40 A. M. Potts and P. C. Au, *Exp. Eye Res.*, 1976, **22**, 487–491.

