

## Supplementary Materials

### **In situ formation of DNA-templated copper nanoparticles as fluorescent indicator for hydroxylamine detection**

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***DNA Sequences were used in this study:***

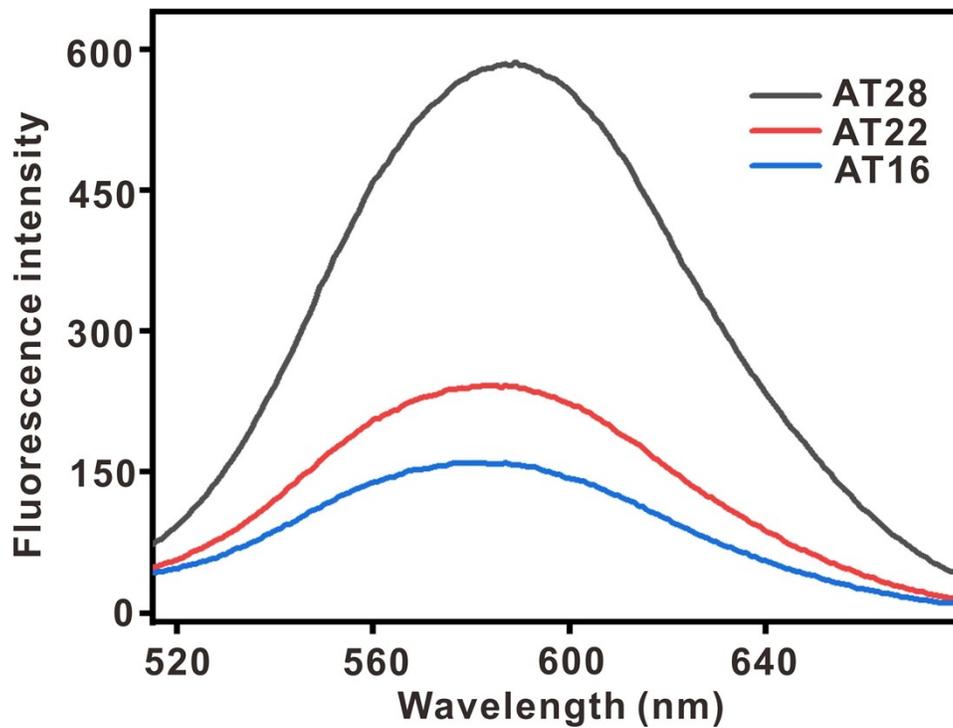
<b>Code</b>	<b>Sequence (5'→3' / 3'→5')</b>	<b>Length (bp)</b>
<b>AT16</b>	ATATATATATATATAT / TATATATATATATATA	<b>16</b>
<b>AT22</b>	ATATATATATATATATATATAT / TATATATATATATATATATA	<b>22</b>
<b>AT28</b>	ATATATATATATATATATATATATATATA / TATATATATATATATATATATATATAT	<b>28</b>
<b>AT34</b>	ATATATATATATATATATATATATATATATATATAT / TATATATATATATATATATATATATATATATA	<b>34</b>
<b>AT40</b>	ATATATATATATATATATATATATATATATATATATA TATAT / TATATATATATATATATATATATATATATATATAT ATATA	<b>40</b>

***Preparation of dsDNA templates:***

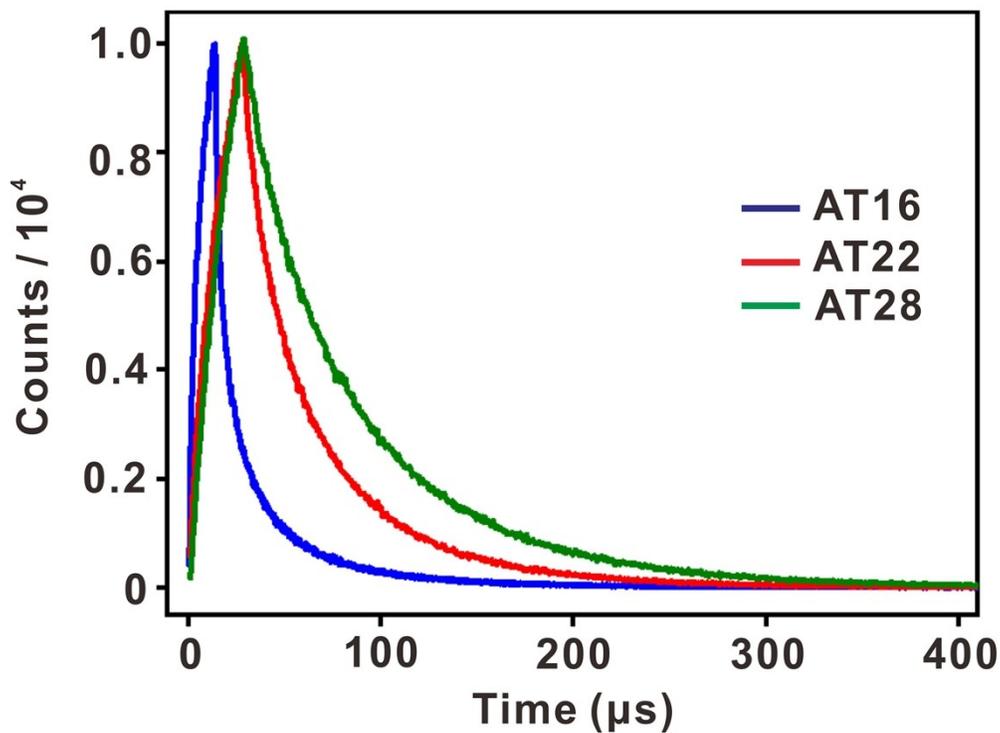
Firstly, both the oligonucleotide (1  $\mu\text{M}$ ) and its complementary strand were mixed together at 1:1 molar ratio in the MOPS buffered solutions (10 mM MOPS, 2 mM  $\text{MgCl}_2$ , 150 mM NaCl, pH 7.5). Then, the mixture was annealed at 95  $^\circ\text{C}$  for 10 min, and then slowly cooled down to the room temperature (25  $^\circ\text{C}$ ).

***Preparation of fluorescent DNA-CuNPs:***

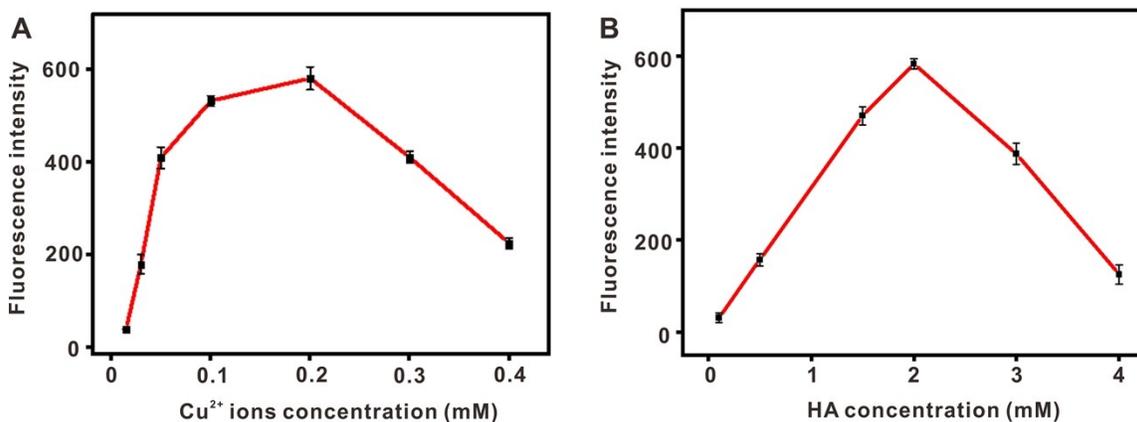
Firstly, 500  $\mu\text{L}$  MOPS buffered solutions and final concentration of hydroxylamine hydrochloride (2 mM) was mixed in 2 mL EP tubes. Then, after the solution was incubated for 1 min, final concentration of  $\text{CuSO}_4$  (100  $\mu\text{M}$ ) was introduced. Finally, the synthesis of fluorescent DNA-CuNPs were completed within 10 min incubation time at room temperature.



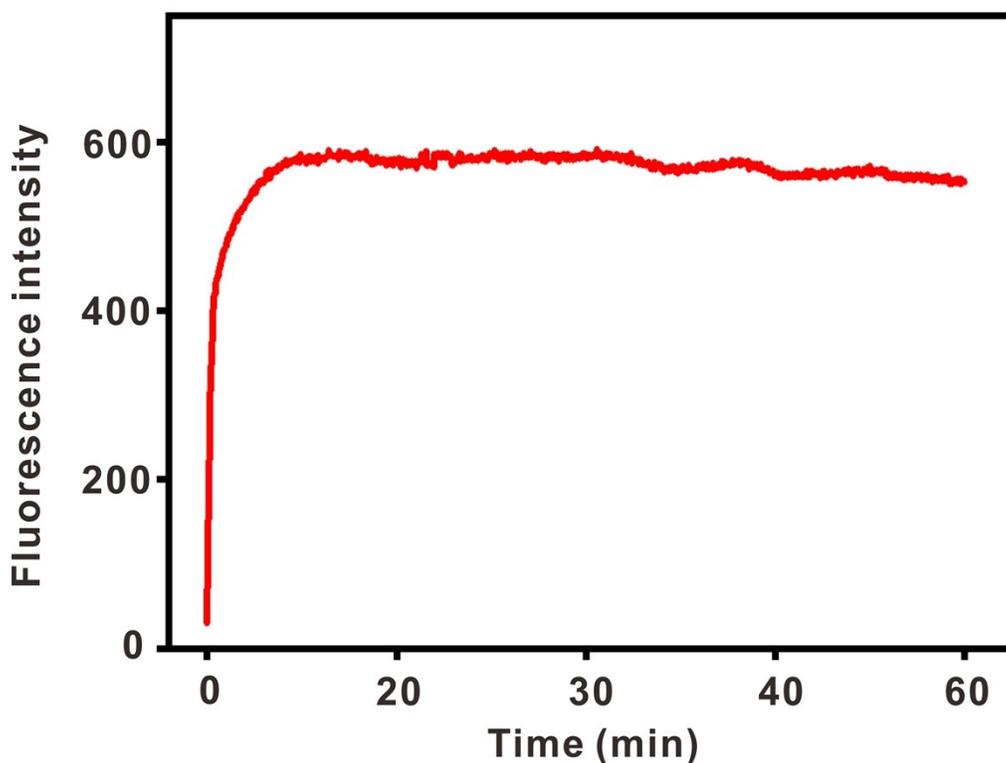
**Fig. S1.** Fluorescence spectra of CuNPs templated by various AT-rich sequence templates: AT16, AT22, and AT28.



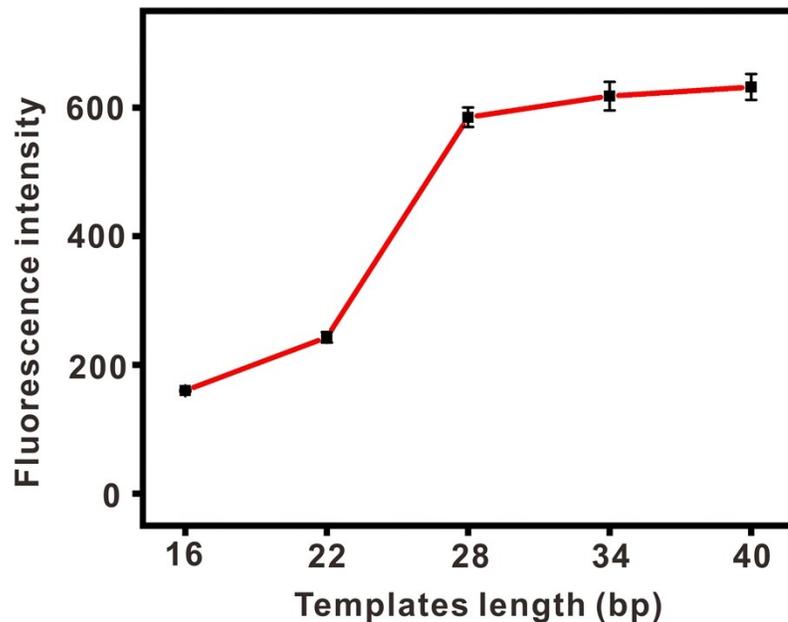
**Fig. S2.** Fluorescence decay curves of CuNPs templated by AT16, AT22, and AT28.



**Fig. S3.** Effect of the synthetic conditions at  $\lambda_{em} = 588$  nm ( $\lambda_{ex} = 340$  nm). A) The concentrations of  $\text{Cu}^{2+}$  ions were 0.015, 0.03, 0.05, 0.1, 0.2, 0.3, and 0.4 mM, respectively. B) The concentrations of hydroxylamine (HA) were 0.5, 1, 1.5, 2, 2.5, 3, and 4 mM, respectively. The error bars represent the standard deviation of three independent measurements.



**Fig. S4.** Time dependence of the fluorescence intensity of AT28-CuNPs ( $\lambda_{ex} = 340$  nm,  $\lambda_{em} = 588$  nm) in MOPS buffered solutions.



**Fig. S5.** The fluorescence intensity of CuNPs based on different templates length of AT-rich DNA: AT16, AT22, AT28, AT34, and AT40 ( $\lambda_{\text{ex}} = 340 \text{ nm}$ ,  $\lambda_{\text{em}} = 588 \text{ nm}$ ) in MOPS buffered solutions.

**Table S1.** Detection of HA from the local tap water, the local ground water, and the local river water near the factory.

	Samples	Added (mM)	Detected (mM)	Recovery (%)	RSD, n=9 (%)
Tap water	1	0.3	0.289	96.3	3.8
	2	0.8	0.782	97.8	3.5
	3	1.2	1.124	93.7	4.4
Ground water	1	0.3	0.278	92.7	3.7
	2	0.8	0.771	96.4	4.6
	3	1.2	1.108	92.3	5.8
River water	1	0.3	0.278	92.7	5.6
	2	0.8	0.788	98.5	4.8

	3	1.2	1.146	95.5	4.2
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**Table S2.** Comparison of the proposed method with other recent methods for HA detection.

	Method	Detection limits	Response Time	Reference
1	HPLC	0.05 mM	More than 1 h	W. D. Korte, <i>J. Chromatogr.</i> , 1992, 603, 145
2	Spectrophotometric determination by flow injection analysis	0.05 mM	More than 1 h	G. C. M. Bourke, G. Stedman, A. P. Wade, <i>Analytica Chimica Acta</i> , 1983, 163, 277-280
3	Fluorescent sensor: DNA-CuNPs	0.022 mM	10 min	This work
4	Kinetic spectrophotometric determination	0.58 $\mu$ M	More than 1 h	A. Afkhami, T. Madrakian, A. Maleki, <i>Anal. Sci.</i> , 2006, 22, 329.
5	Ion chromatographic determination	36 nM	More than 1 h	C. Xu, <i>Modern Agrochemicals</i> , 2017, 16, 36.
6	GC	14.4 nM	More than 1 h	Y. Seike, R. Fukumori, Y. Senga, H. Oka, K. Fujinaga, M. Okumura, <i>Anal. Sci.</i> , 2004, 20, 139.