1	Supplementary information
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3	An all-in-one enzyme-free fluorescent aptasensor
4	integrating localized catalyzed hairpin assembly
5	for sensing of antibiotic in food with improved detection efficiency
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- 21 **Table S1** Sequences of oligonucleotides used in this study.
- 22 The red sequence in S2 is the primer sequence, and the green sequence in S2 is the
- aptamer sequence.

Name	Sequences (5'-3')
S1	CCCACAGTTAGATTCTTTTTTTTTTTTTTTTTTTTTTTT
S2	AATCAACTGCGAGAATCTAACTGTGGGGGGTTGAGGCTAAGCCGA
H1	AGATTGAGGGTTTGGGTGATTTTCAGTTAGATTCTCGCAGTTGATTCCATGT
	GTAGAAATCAACTGCGAGAA
H2	TAGTTAGTATGCTTGGCTGATTTAGTTGATT(BHQ2)-CTACACATGGAATCAA
	CTGCGAGAACCATGTGTAGA-(Cy5)
А	ATCACCCAAACCCTCAATCTTTTACATTCCTAAGTCTGAAACATTACAGCTT
	GCTACACGAGAAGAGCCGCCATAGTA
В	TCAGCCAAGCATACTAACTATTTTATCACCAGGCAGTTGACAGTGTAGCAA
	GCTGTAATAGATGCGAGGGTCCAATAC
С	TCAACTGCCTGGTGATAAAACGACACTACGTGGGAATCTACTATGGCGGCT
	СТТСААААААААААААААААА
D	TTCAGACTTAGGAATGTGCTTCCCACGTAGTGTCGTTTGTATTGGACCCTCG
	CAT



Fig. S1. (A) Polyacrylamide gel electrophoresis (15%) characterization the fabrication
of dsDNA S1-S2. (B) Agarose gel electrophoresis (3%) characterization the assembly

- of DNA tetrahedron.



31

Fig. S2. DLS analysis of DNA tetrahedron and L-CHA-based all-in-one enzyme-free
 aptasensor.

<sup>35</sup> DLS characterization was performed to provide the sizes of DNA tetrahedron <sup>36</sup> before and after modification. DNA tetrahedron (50  $\mu$ L, 100 nM) and L-CHA-based <sup>37</sup> all-in-one enzyme-free aptasensor (50  $\mu$ L, 100 nM) were respectively added into <sup>38</sup> 50- $\mu$ L disposable cuvettes for DLS measurements. As shown in Fig. S2, the size of <sup>39</sup> DNA tetrahedron was 13.6 nm whereas the size of the all-in-one aptasensor was 20.7 <sup>40</sup> nm. The increase in size was attributed to the connection of S1-S2, H1, and H2 to the <sup>41</sup> DNA tetrahedron.

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Fig. S3. Optimization the reaction time of dsDNA S1-S2, hairpins H1, and H2 binding to DNA tetrahedron. Condition:  $C_{S1-S2} = C_{H1} = C_{H2} = C_{DNA \text{ tetrahedron}} = 100 \text{ nM},$ 

47  $C_{kanamycin} = 100 \ \mu g/mL$ . Error bars were the standard derivation (n=3).