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1	Supporting information						
2	Fluorescence immunoassay based on the enzyme cleaving ss-DNA to						
3	regulate the synthesis of histone-ds-poly(AT) templated copper						
4	nanoparticles						
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25 Figure S1. Optimization of synthesis CuNPs. A. Optimization of CuSO₄ in the presence 26 of 2.50 mM of AA, 5 mg/mL of histone with 5 μ M of ds-poly(AT). B. Optimization of 27 AA in the presence of 2.50 mM of CuSO₄, 5 mg/mL of histone with 5 μ M of ds-28 poly(AT). C. Optimization of histone in the presence of 2.50 mM of CuSO₄, 5 mM of 29 AA with 5 μ M of ds-poly(AT).

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43 Figure S2. A. Optimization of ds-poly(AT) in the presence of 2.50 mM of CuSO₄, 5
44 mM of AA, 10 mg/mL of histone. B. Calibration curve of ds-poly(AT) for synthesizing
45 of CuNPs. C. Photo show of CuNPs in the absence (left) and presence (right) of CuSO₄.
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56 Figure S3. Absorbance changes upon the interaction of TMB with different concentration of HRP ranging from 0.67 fM to 2750 fM. The LOD value of the HRP to TMB was calculated as 7.00 fM, which was defined as the lowest concentration of HRP that generated a higher absorbance than the blank absorbance plus 3 standard deviations.

7879 HRP-based conventional immunoassay for AFB₁ detection

80 The 96-well microplates were first coated with 100 µL of protein G (25 µg/mL) in bicarbonate buffer (0.10 M, pH 8.60) at 4 °C overnight and then 81 blocked with 1 mg/mLof BSA solution at 37 °C for 1 h. After washing three 82 times with PBS buffer containing 0.05% Tween-20 (PBST, pH 7.40), 100 µL of 83 anti-AFB1 mAbs (75 ng/mL) was added and incubated at 37 °C for 1 h. After 84 85 washing with PBST three times, 50 µL of HRP@AFB₁ solution (0.50 µg/mL) and 50 µL of sample solution were added into each plate well. After incubation 86 at 37 °C for 60 min, the microplates were washed with PBST five times, and then 87 100 µL of TMB solution was added. After incubation for 15 min at room temperature, 88 the reaction was terminated with 50 µL of 2 M H₂SO₄, and the absorbance was 89 90 measured at 450 nm using a microplate reader.





92 Figure S4. Calibration curve of the conventional immunoassay. The dynamic linear 93 range of AFB_1 concentration is 0.04 - 10 ng/mL. Vertical bars indicate the standard 94 deviation (n = 3).

	Dilution ratio of GOx-AFB ₁	Dilution ratio of anti-AFB ₁ mAbs (500 µg/mL)				
(588 μg	(588 µg/mL)	2000	4000	8000	16000	32000
	200	1.03	1.43	1.16	1.54	6.27
	400	1.04	1.00	2.47	3.73	6.36
	800	1.36	1.37	2.51	3.86	6.14
	1600	3.77	3.98	4.19	4.20	6.25
96	3200	4.45	6.39	6.40	6.42	6.57
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Table S1. The values of FI in the checkerboard titration.

110 UPLC-FLD Analysis Method

111 The reliability and practicability of the developed fluorescence immunoassay was further confirmed by UPLC-FLD. Samples pretreatment and UPLC-FLD operation 112 were conducted according to the national standard GB 5009.96-2016 (China) with 113 114 some modification. Briefly, corn samples (1g) was extracted by ultrasound with 5 mL of 60% (v/v) methanol/water for 20 min. After centrifugation at 6000 rpm 115 116 for 15 min, 3 mL of supernatant solution was further diluted three folds with 6 117 mL PBS. Extraction of samples were filtered with 0.22 μ M cellulose membrane and spiked with different concentrations of AFB₁. Then, samples were cleaned by immune 118 119 affinity column and further employed for UPLC-FLD analysis. Chromatographic 120 separation was achieved with an ACQUITY UPLC BEH C18 ($1.70 \mu m 2.10 \times 100 mm$) column using mobile phase methanol-water (30-70), Ex365 nm and Em436 nm. The 121 flow rate was 0.20 mL/min, column temperature was 30°C and the injection volume 122 was 1 µL in the full injection mode. The monitoring appearance time was chosen as 123 124 2.731 min (quantitation time, standards) and 2.745 (qualitative time, samples).