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1 **Screen-printed electrochemical immunosensor based on a novel**
2 **nanobody for analyzing aflatoxin M₁ in milk**

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23

24

25 **Abstract**

26 This study aimed to devise a nontoxic electrochemical immunosensor to
27 quantitatively determine aflatoxin M₁ by chronoamperometry with novel
28 anti-idiotypic nanobody-functionalized screen-printed carbon electrodes (SPCEs).
29 Anti-idiotypic nanobodies (AIdnb) were developed to replace the high toxic
30 chemically synthesized antigen. AIdnb was immobilized on the surface of SPCE
31 via covalent coupling as capture reagent. The functionalized SPCEs were followed by
32 characterization using electrochemical impedance spectroscopy, fourier-transform
33 infrared spectroscopy, transmission electron microscopy mapping, and atomic force
34 microscopy. After optimizing experimental parameters, the assembled immunosensor
35 exhibited a good linearity range of 0.25–5.0 ng/mL, with the limit of detection of 0.09
36 ng/mL. The immunosensor showed a satisfactory selectivity to AFM₁, without
37 interference from analogs, including zearalenone, ochratoxin, and fumonisin B₁. For
38 practical application, the developed immunosensor was validated using real spiked
39 samples with the recovery range 82.0%–108.0% and relative standard deviation (RSD)
40 10.1%–13.0%, indicating that it could be used in milk samples.

41

42 **Keywords:** Aflatoxin M₁; anti-idiotypic nanobody; chronoamperometry; milk;
43 screen-printed carbon electrodes

44

45 **1. Introduction**

46 Aflatoxins are a class of *Aspergillus* fungal metabolites, among which aflatoxin B₁ is
47 the most concerned mycotoxin due to its strong teratogenicity, carcinogenicity, and
48 mutagenicity (Li, Liu, Zhang, Luo, Lin, & Jiang, 2021). AFM₁ is a product of the
49 hydroxylated metabolism of aflatoxin B₁, which has mutagenic and teratogenic effects
50 on humans, particularly infants. The World Health Organization and the International
51 Agency for Research on Cancer classified AFM₁ as Group 2B human carcinogen (Du,
52 Su, Yang, Pan, Liu, Gong, et al., 2016). AFM₁ is the major type of contaminant in
53 milk and milk products. Thus, the safety related to milk consumption has gained
54 immense attention worldwide. On account of the high toxicity of aflatoxins, strict
55 limits were set by most countries worldwide. For example, Europe set a limit of 0.25
56 ng/mL for AFM₁ in milk, while the US Food and Drug Administration has set the
57 maximum level for AFM₁ at 0.5 ng/mL in milk. It is rather essential to develop rapid
58 detection methods for protecting consumption safety so as to detect the
59 ultra-trace-level contamination in food as early as possible.

60

61 In recent years, competitive immunoassay has been widely employed for the rapid
62 detection of mycotoxins (Lai, Wei, Xu, Zhuang, & Tang, 2017; Lin, Zhou, & Tang,
63 2017; Lin, Zhou, Tang, Niessner, & Knopp, 2017; Lin, Zhou, Tang, Niessner, Yang, &
64 Knopp, 2016; Zhou & Tang, 2020). Immunosensors are used in AFM₁ measurements
65 by incorporating enzyme catalysis (Di Giovanni, Zambrini, Varriale, & D'Auria,
66 2019), chromatography (Han, Gong, Wang, Zhang, Jin, Zhao, et al., 2019),
67 electrochemistry (Abera, Falco, Ibba, Cantarella, Petti, & Lugli, 2019), and so forth.
68 Most of them provide satisfactory detection results. Immunosensors based on
69 electrochemistry can be a good way to detect AFM₁ due to their many advantages
70 including cost- and time-effective procedure, sensitivity, and so forth (Jia, Liao, Fang,
71 Jia, Liu, Li, et al., 2021; Wei, Sun, Gao, Yang, Ye, Ji, et al., 2021). In addition,
72 immunosensors can be integrated with different types of detection strategies,
73 especially low-cost devices such as SPCEs, which can be an excellent advantage for

74 mycotoxin detection. Compared with the traditional rod-type electrodes, SPCE
75 technology offers remarkable benefits but is not limited to disposability and
76 portability (Sharma, Istamboulie, Hayat, Catanante, Bhand, & Marty, 2017). As we
77 know, recognition elements (monoclonal antibody, single-chain antibody, nanobody,
78 aptamer, etc.) always decide the sensitivity and the limit of detection of
79 immunosensors. Thus, research works have been done on developing high-quality
80 antibodies to increase the sensitivity and feasibility of immunosensors.

81

82 Nanobody is a variable domain of heavy chain of heavy-chain antibody (VHH)
83 obtaining from Camelidae species and sharks. VHH emerges as a nanobody because
84 its molecular size is only about 15 kDa and the dimension is about $2.5 \times 3 \times 4 \text{ nm}^3$
85 (Yu, Xu, Wu, Jiang, Wei, Zulipikaer, et al., 2020). The advantages of nanobodies
86 include high affinity, selectivity, solubility, and yield and low-cost productions (Tang,
87 Li, Zhang, Zhang, Zhang, & Jiang, 2017). Due to another disulfide bond in the
88 complementary determining region 3, nanobodies exhibit high tolerance to organic
89 solvents and high temperatures (Simões, Guedens, Keene, Kubiak-Ossowska,
90 Mulheran, Kotowska, et al., 2021). AIdnb are developed by immunizing alpacas with
91 monoclonal antibodies; they have “mirror image” relationships with the targets. With
92 their favorable properties, AIdnb are considered as promising reagents that can
93 replace highly toxic traditional antigens and standard solutions. Previous studies used
94 AIdnb in enzyme-linked immunosorbent assay (ELISA), polymerase chain reaction
95 (PCR), and immunochromatography successfully (Table S1). However, the field of
96 application of AIdnb should be continuously expanded.

97

98 A screen-printed electrochemical immunosensor based on an alternative antigen was
99 developed in this study to explore a new platform technique of AIdnb. This was the
100 first assay established using a nontoxic SPCE electrochemical. It could perform the
101 rapid, low-cost monitoring of milk consumption safety.

102

103 **2. Experimental**

104 **2.1 Chemicals and reagents**

105 Anti-aflatoxin M₁ monoclonal antibody (mAb) 2C9 was developed in our laboratory,
106 which had cross-reaction with AFB₁, AFB₂, AFG₁, and AFG₂ of <0.01%. Aflatoxin B₁,
107 B₂, G₁, G₂, M₁ standard, bovine serum albumin (BSA), isopropyl
108 β-D-thiogalactopyranoside (IPTG), and 3, 3', 5, 5'-tetramethylbenzidine (TMB) were
109 obtained from Sigma (MO, USA). A LeukoLOCK total RNA isolation system was
110 obtained from Applied Biosystems (CA, USA). *Escherichia coli* ER 2738 competent
111 cells from the ER2673 line of *E. coli* were purchased from Lucigen Corp. (WI, USA).
112 TOP10F' competent cells were procured from Life Technologies (NY, USA). Helper
113 phage M13KO7, *sf*I, and T4 DNA ligase were obtained from New England Biolabs
114 (MA, USA). QIAprep Spin MinPrep Kit, QIAquick Gel Extraction Kit, and QIAquick
115 PCR Purification Kit were obtained from Qiagen. Tween 20 was obtained from J&K
116 Scientific (Beijing, China). Mouse anti-M13 monoclonal antibody conjugated to
117 horseradish peroxidase (HRP) was purchased from Sino Biological (Beijing, China).
118 Goat anti-mouse monoclonal antibody conjugated to HRP was purchased from
119 Solarbio (Beijing, China). The Costar 96-well EIA/RIA plate was purchased from
120 Corning Incorporated (Corning, NY, USA). Potassium ferrocyanide [K₄Fe(CN)₆],
121 potassium ferricyanide [K₃Fe(CN)₆], BSA, N-hydroxysuccinimide (NHS),
122 1-ethyl-3-(3-dimethylamino-propyl)-carbodiimide (EDC), sodium phosphate dibasic
123 (Na₂HPO₄), potassium phosphate monobasic (KH₂PO₄), magnesium chloride (MgCl₂),
124 potassium chloride (KCl), sulfuric acid (98%), sodium chloride (NaCl), ethanol (98%),
125 sodium nitrite (NaNO₂), 4-aminobenzoic acid, ethanolamine, HRP-conjugated
126 secondary antibody and aflatoxin M₁ were purchased from Sigma–Aldrich (France).
127 Ultrapure water from a Milli-Q system was used for preparing aqueous solutions and
128 rinsing procedures.

129

130

131 **2.2 Apparatus and electrodes**

132 SPCEs, which consisted of three-electrode configurations (working, counter, and
133 reference electrodes), were fabricated using a DEK 248 screen-printing system (BAE
134 laboratory, Perpignan, France). Electrochemical measurements were performed on an
135 Autolab PGSTAT100 potentiostat/galvanostat station equipped with a Nova 2.1
136 analyzer system (Metrohm Autolab B.V., Kanaalweg, The Netherlands).

137

138 **2.3 Chemical modification of SPCEs**

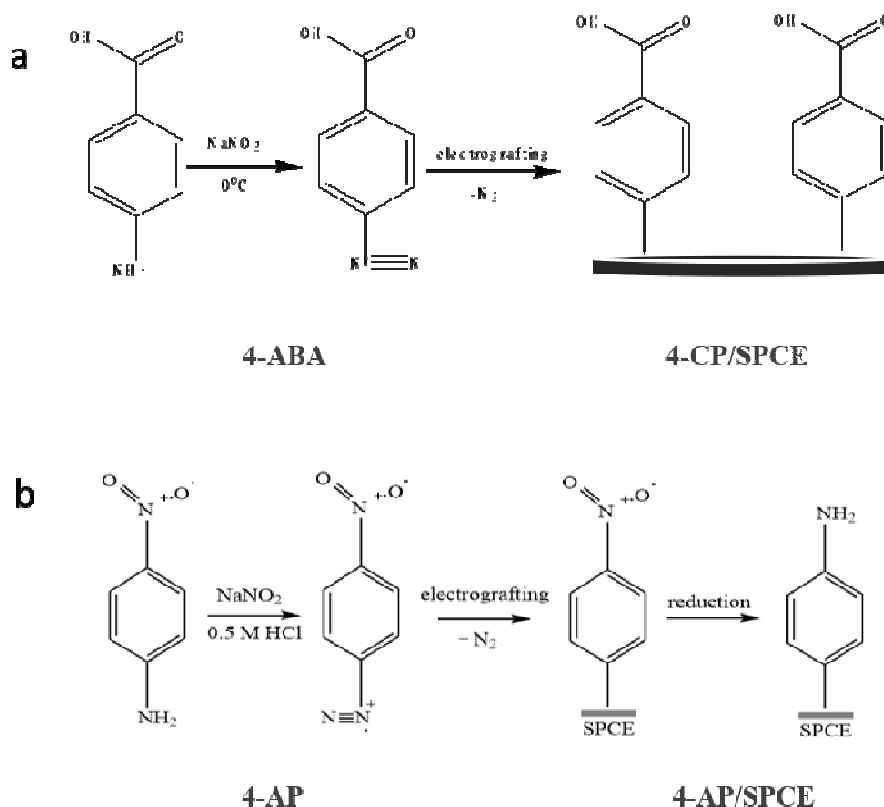
139 SPCEs were cleaned by applying six potential cycles between 1.5 to -1.0 V/pseudo
140 Ag reference electrode with 100 mVs^{-1} scan rate in the mixture of 0.5 mol/L sulfuric
141 acid and 0.1 mol/L KCl until clean SPCEs were obtained. The clean SPCEs were
142 chemically modified according to the reactions shown in Scheme 1. As shown in
143 Scheme 1a, the SPCEs were modified using 2 mmol/L sodium nitrite and 2 mmol/L
144 4-aminobenzoic acid (ABA) in 0.5 mol/L HCl. The mixture was allowed to react for 5
145 min for the maximum generation of diazonium salt at 4°C . Then, $150 \mu\text{L}$ of the salt
146 solution was electrografted on SPCE to generate 4-carboxyphenyl diazonium salt
147 SPCE (4-CP/SPCE) via linear sweep voltammetry (LSV) (potential: 0.4 V to -0.8 V ;
148 scan rate: 50 mVs^{-1}). The 4-CP/SPCEs were rinsed with phosphate-buffered saline
149 (PBS) (0.01 mol/L) in the subsequent step.

150 The other modification method (Scheme 1b) involved preparing the 4-aminophenyl
151 diazonium salt SPCE (4-AP/SPCE). The clean SPCEs were modified by the following
152 procedure. In brief, $4 \mu\text{L}$ of 0.1 mol/L NaNO_2 was added to $196 \mu\text{L}$ of 2 mmol/L
153 4-nitroanilineaminobenzoic acid in 0.5 mol/L HCl, stirred, and reacted for 5 min at
154 4°C . Then, the 4-AP/SPCEs were generated by LSV from 0.4 to -0.5 V . The
155 4-AP/SPCEs were rinsed with PBS (0.01 mol/L) and used for the subsequent step.

156

157

158



160

161 Scheme 1. Modification of SPCE with 4-aminobenzoic acid (a) and 4-aminophenyl diazonium
 162 salt (b).

163

164 **2.4 Development of anti-idiotypic nanobody of AFM₁**

165 The nanobody with a His tag at its C-terminal end was developed as depicted earlier.
 166 Briefly, a 2-year-old male alpaca was immunized with 100 µg/mL monoclonal
 167 antibody against AFM₁ (2C9). Total RNA was extracted and transcribed into cDNA.
 168 Primers were used to amplify the VHH and then ligated to the pComb3X. The
 169 pComb3X/VHH were transformed into *E. coli* ER 2738. The positive clones were
 170 screened by a complete elution method. During elution, 500, 20, and 5 ng/mL of
 171 aflatoxin M₁ in 10% methanol–PBS was used as the elution buffer for each round.
 172 Phagemids of positive clones were transformed into nonsuppressor *E. coli* strain
 173 TOP10F' cells, expressed, and purified with Ni-NTA metal affinity chromatography.
 174 Finally, the nanobody was stored at –20°C to use.

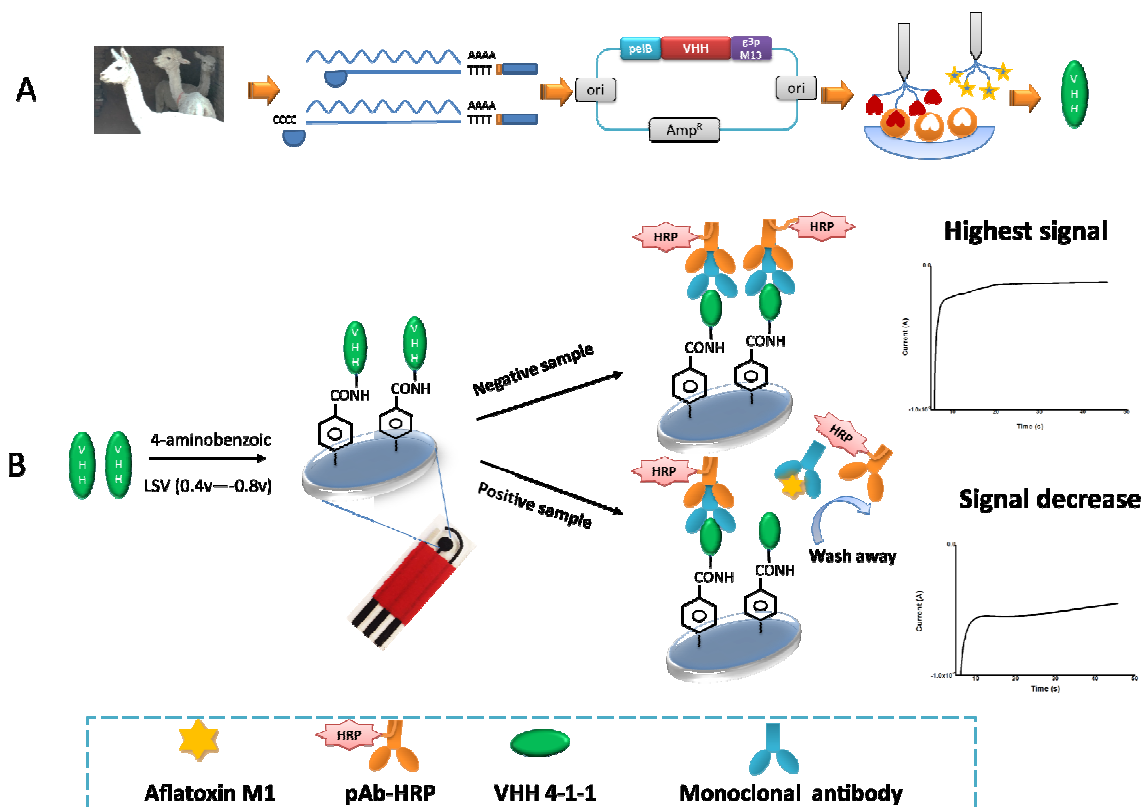
175

176

177 **2.5 Immunosensor for detection of AFM₁**

178 The principle of the immunosensor was based on the competitive reaction on SPCE.
179 In the negative samples, the mAbs were captured by the nanobody on the working
180 electrode surface while in the positive samples, the mAbs reacted with the AFM₁
181 resulting in less mAbs captured on the working electrode surface, the electrochemical
182 signal decreased as a result of increased concentration of AFM₁. The assembly steps
183 of the AFM₁ immunosensor and detection mechanism are illustrated in Scheme 2.
184 Firstly, the modified SPCEs were activated by incubation with 20 μ L of
185 EDC/NHS/nanobody in 4 mg/mL N- methyl piperidine (MES) buffer (pH 6.8) for 180
186 min. After incubation, SPCEs were rinsed with PBS (0.01 mol/L, pH 7.4) and stored
187 at 4°C for further use. The competition step was performed by incubating antibody
188 2C9 and equivalent volumes from a series of concentrations of AFM₁ on the working
189 electrode surface for 30 min at 37°C, with shaking at 250 rpm. The surface was rinsed
190 six times with PBST (PBS containing 0.05% Tween 20). Subsequently, 15 μ L of
191 HRP-labeled secondary antibody was added to the working electrode surface,
192 incubated for another 30 min, and rinsed six times with PBST. Then, 50 μ L of TMB
193 solution was used to cover the working counter and reference electrodes. The enzyme
194 activity was measured electrochemically using chronoamperometry (CV).

195 Scheme 2



196

197 Scheme 2. (A) Development of nanobody VHH 4-1-1. (B) Fabrication process mechanism of a
 198 nanobody-based electrochemical immunosensor for AFM₁ detection.

199

200 2.6 Electrochemical measurements

201 The impedimetric measurements (electrochemical impedance spectroscopy, EIS) were
 202 used for characterizing the immunosensor surface using 1mM ferri/ferrocyanide/PBS
 203 ($[\text{Fe}(\text{CN})_6]^{4-/3-}/\text{PBS}$) with a frequency range of 10 kHz to 0.5 Hz. The Nyquist plots
 204 were recorded with a cyclic voltammetry amplitude of 5 mV and a sampling rate of
 205 20 points. The surfaces of SPCEs were characterized using EIS after each
 206 modification step. Chronoamperometry measurements were performed for 40 s at 0.4
 207 V. Cyclic voltammetry measurements were obtained between maximum and
 208 minimum voltages of 0.8 V and -0.6 V (vs Ag/AgCl reference electrode), respectively,
 209 at a scan rate of 0.1 V/s.

210

211

212 **2.7 Application of the immunosensor for real sample analysis**

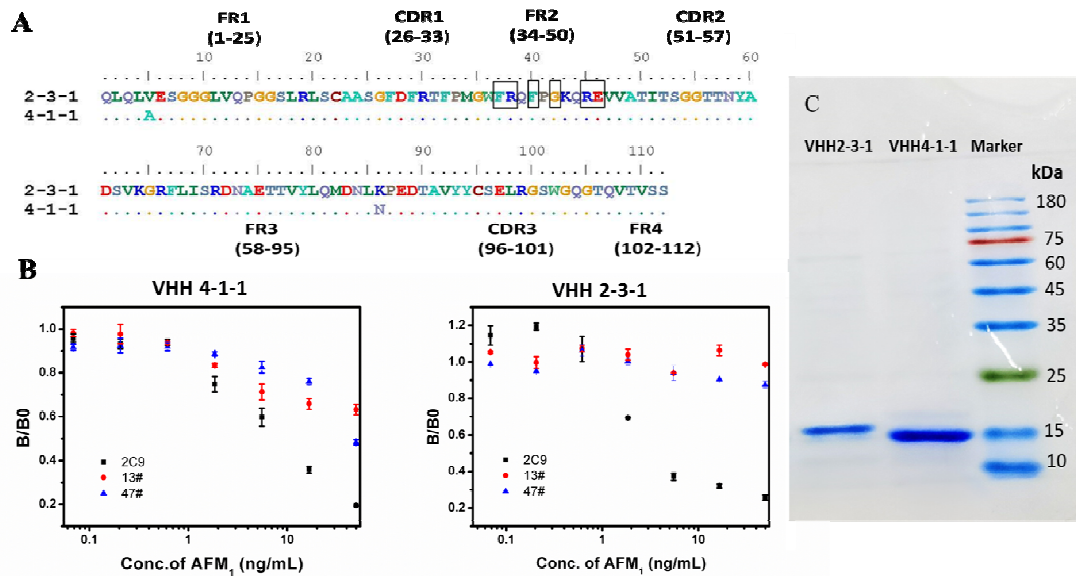
213 The milk sample was spiked with different concentrations of AFM₁ of 0, 0.25, 0.5,
214 1.25, 2.5, and 5.0 ng/mL. The milk samples were cultured at 40°C for 30 min and then
215 demulsified with ethanol (1/3, v/v). Further, 800 μL of milk was centrifuged at 6000
216 rpm for 5 min, followed by adding 90.0 μL of PBS to 10.0 μL of supernatant. Based
217 on the optimized parameter detection, the current signal value corresponding to each
218 concentration was determined. By calculating the ratio between different
219 concentrations of AFM₁ and the blank one (without AFM₁ standard), the standard
220 curve between the ratio and the corresponding AFM₁ concentration was established.

221

222 **3. Results and Discussion**

223 **3.1 Nanobody characterization**

224 The total RNA was extracted from the blood of immunized alpaca, and then the
225 heavy-chain IgG2 and IgG3 sequences were amplified (Fig. S1), followed by the
226 insertion of the amplified sequence into the vector pComb 3X. The constructed
227 nanobody library exhibited a size of 2.7×10^7 (Fig. S2). We utilized a competitive
228 panning strategy to select the specific nanobody. After three rounds of panning, two
229 sequences were obtained, defined as VHH 4-1-1 and VHH 2-3-1. Both contained 37 F,
230 38 R, 40 F, 42 G, 45 R, and 46 E in FR2, which explained the solubility of nanobodies
231 (Fig.1A). The two plasmids were transformed to TOP10F⁺ for expression, and the
232 nanobody with 6× His tag was purified using an Ni-NTA affinity column (Fig.1C).
233 The nanobodies were tested for their sensitivity and specificity by indirect
234 competitive ELISA. The IC₅₀ of nanobodies against AFM₁ and 2C9 was 8.54 and
235 3.33 ng/mL, respectively. The cross-reactivity of VHH 4-1-1 and VHH 2-3-1 with
236 other mAbs against AFM₁ (2C9, 13# and 47#) was 100%, <0.1%, and 17.7% and
237 100%, <0.1%, and <0.1%, respectively (Fig.1B). The stability of nanobody was
238 investigated. The results showed that the gene sequence of VHH 2-3-1 changed
239 during subculture, the 5 V changed to 5 A, and 86 K changed to 86 N. Therefore,
240 VHH 4-1-1 was selected as the recognition reagent to establish the detection method.



241

242 Figure 1. Nanobody selection and characterization. (A) Two classes of nanobodies containing
243 different amino acid sequences were determined, named VHH 2-3-1 and VHH 4-1-1. (B)
244 Cross-reactivity of nanobodies with mAb 2C9, 13# and 47#. (C) VHH 2-3-1 and VHH 4-1-1 were
245 purified and analyzed by SDS-PAGE.

246

247 3.2 Characterization of the immunosensor for AFM₁

248 3.2.1 Modification of SPCEs

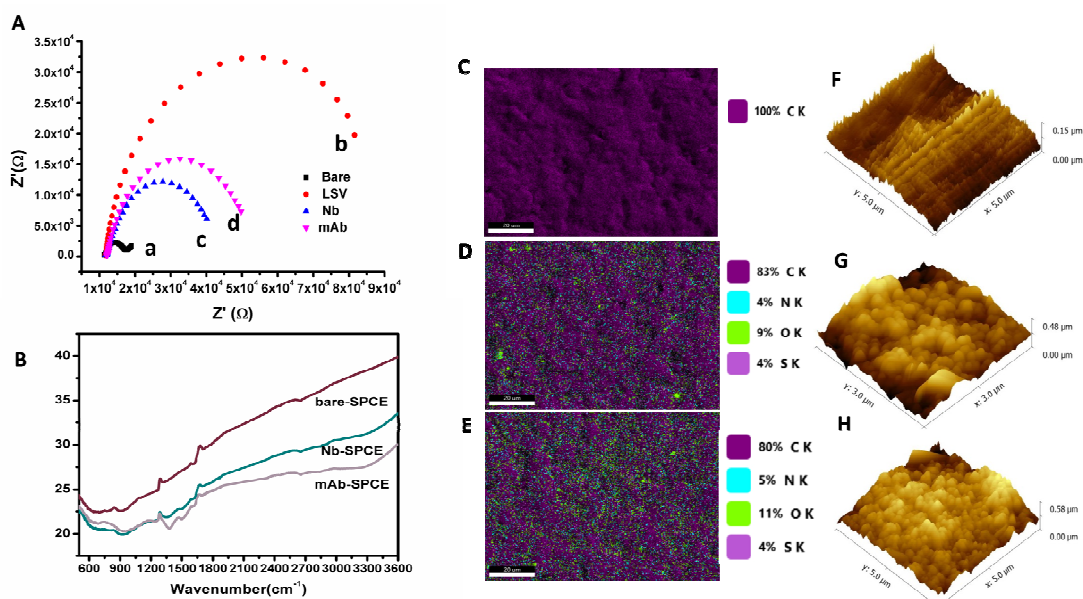
249 In the classical immunoassay, the antibody/antigen should immobilized on the solid
250 phase to ensure immune reaction successfully. Two nanobody modification methods,
251 4-aminobenzoic acid SPCE (SPCE-4ABA) with -COOH and 4-aminophenyl
252 diazonium salt SPCE (SPCE-AP) with -NH₂, were compared to ensure that the active
253 sites of nanobodies were exposed to the antibody. The results showed that using the
254 SPCE-4ABA modification, the current signal decreased with the increase in AFM₁
255 standard solution, while no signal changed using SPCE-AP modification. The
256 N-terminal covalent coupling with the functional SPCE-4ABA yielded better results.
257 Previous studies have shown that localized and oriented immobilization of antibodies
258 could avoid interference with the structure and function of antibodies (Sivaram,
259 Wardiana, Howard, Mahler, & Thurecht, 2018). In this study, we concluded that using
260 functional SPCE-4ABA, the epitope of nanobody was almost fully exposed to ensure

261 maximum binding with the antibody. On the contrary, using functional SPCE-AP, the
262 epitope of nanobody was hidden by itself or SPCE, and it was difficult to capture the
263 antibody effectively.

264

265 3.2.2 Electrochemical characterization

266 EIS offers detailed data of surface change after each modification step of the SPCE
267 electrode. Figure 2A shows the semicircle portion of SPCE, which responded to the
268 electron-transfer resistance. The bare SPCE exhibited the smallest semicircle, which
269 had a low electron-transfer resistance. After diazotization of the SPCE electrode
270 surface with 4-ABA and NaNO_2 , the semicircle increased obviously, revealing that
271 the deposition of the negative terminal on the SPCE surface led to an increase in
272 electron-transfer resistance. The Nb-SPCE activated with EDC/NHS induced a
273 decrease in the semicircle ascribed to the replacement of the carboxyl group by the
274 nanobody, leading to the decrease in electron-transfer resistance. After modifying the
275 mAb, the semicircle increase was attributed to the blocking behavior of mAb, which
276 hindered the diffusion of the redox probe. The results of EIS confirmed the successful
277 modification of the SPCE. LSV and CV were provided to identify the successful
278 modification of SPCE. As shown in Figure S3, a significant signal peak was observed,
279 implying that the diazonium salt was reduced on the SPCE. The bare SPCE and
280 SPCE-4ABA were detected using CV scanning to further verify whether benzoic acid
281 was modified to the surface of SPCE successfully. As shown in Figure S4, the results
282 showed a pair of reversible redox peaks in the bare SPCE scanning curve: EPA =
283 +0.5542 V; EPC = -0.21973V; and the difference between the oxidation peak and the
284 reduction peak $\Delta EP = 0.774$ V, corresponding to the redox reaction of $[\text{Fe}(\text{CN})_6]^{4-/3-}$.
285 The redox peak current of SPCE-4ABA was significantly lower than that of the bare
286 SPCE because the benzoic acid modified on the SPCE hindered the electron transfer
287 of $[\text{Fe}(\text{CN})_6]^{4-/3-}$ (Fig. S4).



288

289 Figure 2. (A) Nyquist plots of modified SPCEs detected with $1\text{mM Fe(CN)}_6^{4-/3-}/\text{PBS}$. (a) Bare
 290 SPCE, (b) SPCE-4ABA, (c) Nb-SPCE, and (d) mAb-SPCE. (B) FT-IR characterization of bare
 291 SPCE, Nb-SPCE, and mAb-SPCE. (C) TEM mapping of bare SPCE, (D) Nb-SPCE, and (E)
 292 mAb-SPCE. (F) AFM image of bare SPCE, (G) Nb-SPCE, and (H) mAb-SPCE.

293

294 3.2.3 FT-IR, TEM mapping, and AFM characterization

295 FT-IR, SEM, and atomic force microscopy (AFM) were performed for the
 296 characterization of bare SPCEs, nanobody modification SPCEs (Nb-SPCEs), and
 297 mAb modification SPCEs (mAb-SPCEs).

298 The results of the FT-IR spectrum confirmed that the Nb-SPCEs had absorption bands
 299 at $1229\text{--}1301\text{ cm}^{-1}$ and vibration bands at 3300 cm^{-1} (str.,-NH), which revealed the
 300 presence of nanobody after coupling conjugation. Then, after incubation with mAb,
 301 much clear absorption bands were observed at amide III ($1229\text{--}1301\text{ cm}^{-1}$) and 3300
 302 cm^{-1} , which strongly confirmed the mAb binding with the nanobody (Fig. 2B).

303 Transmission electron microscopy (TEM) mapping was used to characterize
 304 elemental distribution on the working electrode of SPCEs. Figure 2C shows 100%
 305 elemental C located on the working electrode of bare SPCE. Figure 2D and 2E
 306 represents the appearance of new elemental, including N, O, and S located on the
 307 surface of the electrode, indicating that the nanobody and mAb were conjugated on
 308 the surface of SPCE successfully. AFM is a surface analysis technique capable of

309 imaging the topographical and morphological changes in a specimen surface. The
310 AFM topographies of Nb-SPCEs showed an overlap of proteins and a uniform surface
311 compared with the bare SPCEs (Fig. 2F and 2G), indicating that the nanobody was
312 modified on the working electrode successfully. After incubating mAb, the surface
313 morphology of SPCE was changed due to the nanobody and antibody recognition
314 mechanism, the surface of working electrode overlap protein and becoming uniform
315 further (Fig.2H).

316

317 **3.3 Experimental parameter optimization**

318 The important parameters, including concentrations of nanobody, mAb, and HRP-IgG,
319 required time for incubation, and blocking method, were studied to obtain the most
320 sensitive and effective analytical performance of immunosensors for AFM₁ detection.
321 All the parameters were optimized by measuring the different absolute values of the
322 current signal. The larger the difference, the higher the sensitivity of the analysis,
323 which meant the higher the binding rate of the antibody. The different absolute values
324 of the current signal were estimated directly using the following formula:

325

$$326 \quad \Delta\text{Current} = |\text{Current}_{\text{blank}}| - |\text{Current}_{\text{AFM1}}|$$

327 where $\text{Current}_{\text{blank}}$ is the current response value of mAb 2C9 and PBS incubated;
328 $\text{Current}_{\text{AFM1}}$ is the current response value of mAb 2C9 and AFM₁ incubated.

329

330 3.3.1 Nanobody and antibody concentration

331 The amounts of antigens and antibodies were considered as the key factors in
332 developing immunoassay methods. In theory, the lower the antigen concentration used
333 in the competitive reaction, the higher the sensitivity. Different concentrations of
334 4.375, 8.75, and 17.5 $\mu\text{g/mL}$ of VHH 4-1-1 used as surrogate antigens were
335 immobilized on the working electrode. As observed in Figure 3A, the value of the
336 $\Delta\text{Current}$ was directly proportional to the VHH4-1-1 concentration until 17.5 $\mu\text{g/mL}$.
337 The optimization of mAb 2C9 was carried out using different concentrations of 0.50,

338 0.33, and 0.25 $\mu\text{g/mL}$. The maximum value of the $\Delta\text{Current}$ was obtained at 0.33
339 $\mu\text{g/mL}$, which then decreased at higher concentrations, indicating that the more
340 amounts of mAb 2C9 needed much more free targets to match with, leading to a
341 decrease in sensitivity on AFM_1 analysis (Fig. 3B). The best concentration of
342 HRP-IgG was used at 1:5000 dilution (Fig. 3D).

343

344 3.3.2 Blocking method and immunoreaction time optimization

345 In a traditional immunoassay, it is necessary to block any remaining active sites of
346 AFM_1 ; otherwise, it is considered that the incomplete blockage will affect the
347 accuracy of the immunosensor. Therefore, the effect of the blocking method on the
348 detection sensitivity was investigated. Two blocking buffers, 0.1 mol/L
349 ethanolamine/Tris HCl (pH 7.4) and 1% BSA/PBS (*w/v*) were compared with the
350 control group having no blocking. The $\Delta\text{Current}$ of both blocking methods did not
351 show any obvious change with the AFM_1 increase, considering that ethanolamine
352 blocked the whole active sites, including the sites of the nanobody reacting with the
353 mAbs. The molecule of BSA was too big and blocked all the binding active sites of
354 the nanobody. Therefore, the working electrode was without any blocking in this
355 experiment (Fig. 3E).

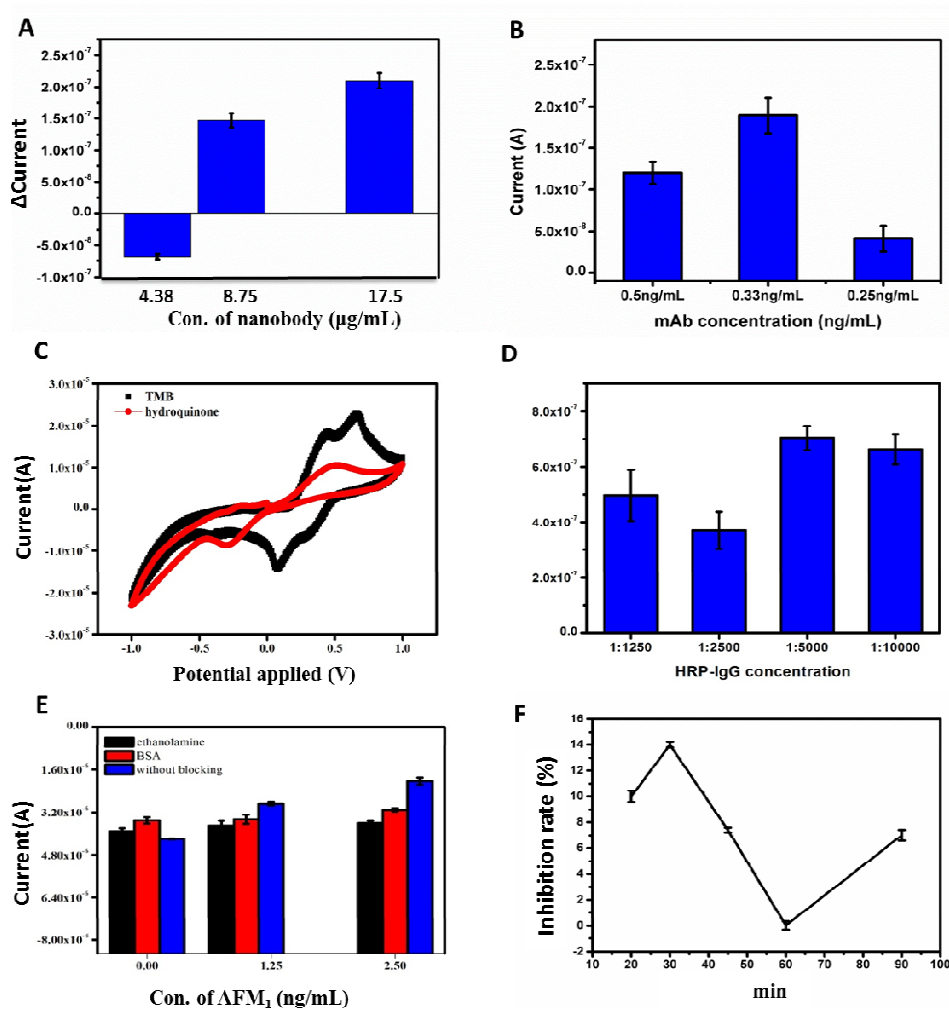
356 With the other factors fixed, the optimization of the immunoreaction time was
357 performed using the immunosensor (nanobody concentration of 0.33 $\mu\text{g/mL}$; mAb
358 2C9 concentration of 17.5 $\mu\text{g/mL}$; HRP-IgG 1:5000 dilution) with AFM_1 at a
359 concentration of 10 ng/mL for different time durations. The results showed that the
360 inhibition rate reached the maximum when the incubation time was 30 min, indicating
361 that much higher amounts of AFM_1 were needed to match with the mAb after that
362 time (Fig. 3F).

363

364 3.3.3 Substrate optimization

365 Finally, the substrate is another critical factor that provides an amplification
366 electrochemistry signal of the immunoreaction. The efficiency of the substrates,
367 hydroquinone, and TMB was evaluated by the CA analysis of the HRP enzyme

368 activities. The results in Fig. 3C demonstrated the TMB radical cation and the
369 charge-transfer complex of the diamine and the diamine were formed in the first step.
370 Then, the intermediate product was further electro-oxidized to the complete oxidation
371 product quinone diimine in the second step. Since the oxidation peak located at 0.07
372 V was considered that of the final products of TMB (TMB^{2+}), the second peak current
373 was used as the optimal applied potential of TMB in the following experiment (Jin, Ko,
374 Kim, Tran, Son, Geng, et al., 2018). The optimal applied potential of hydroquinone
375 was -0.271 V, as assayed by CV. The use of hydrogen peroxide and benzoquinone
376 resulted in no differences in the $\Delta\text{Current}$ value, while TMB and H_2O_2 yielded a
377 $\Delta\text{Current}$ value of 3.69×10^{-7} (Fig. 4A, B). The absolute value of the current signal
378 collected in 30 s decreased, which was consistent with the results of competitive
379 detection in theory; that is, the higher the concentration of the target, the lower the
380 signal value. Therefore, HRP was used to catalyze the H_2O_2 oxidation of TMB for
381 electrochemical signal reaction.



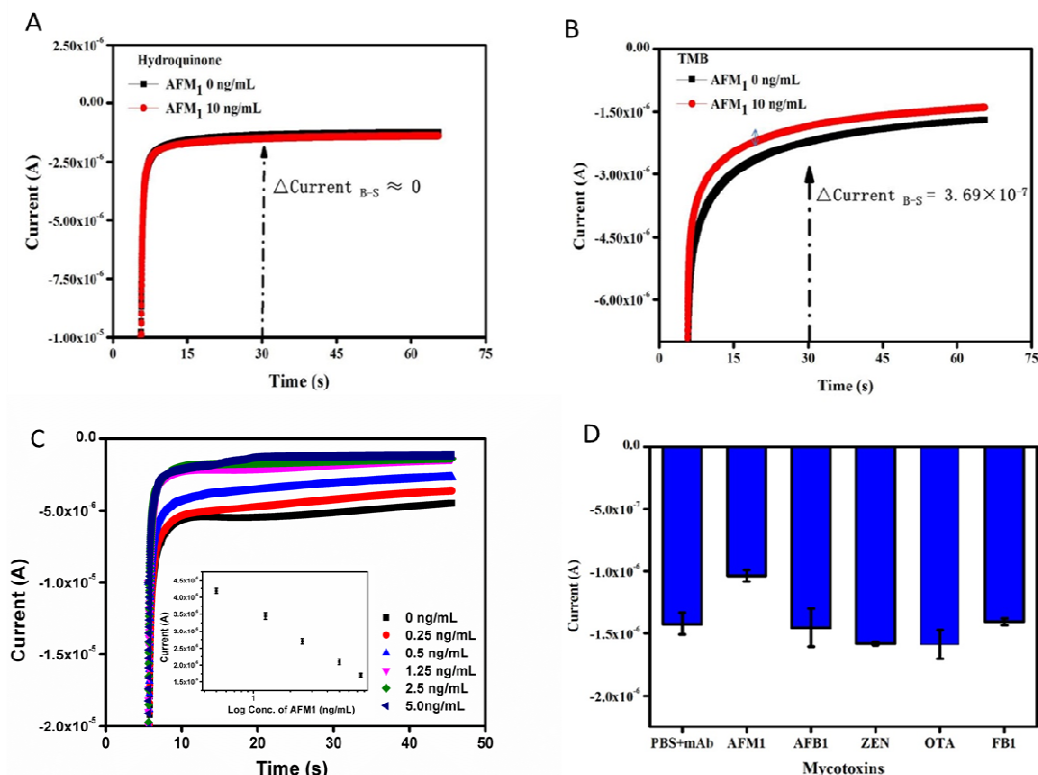
382

383 Figure 3. Currents values *via* CV on (A) nanobody concentration, (B) mAb concentration, (C)
 384 substrate reagents, (D) HRP-IgG concentration, (E) blocking methods, and (F) incubation time of
 385 samples.

386 3.4 Detection of AFM₁

387 With the optimization of the aforementioned factors, the immunosensor was incubated
 388 with different concentrations of AFM₁ to establish the standard curve. As shown in
 389 Figure 4C, a linear relation between the current responses and the logarithm of the
 390 AFM₁ concentration was observed in the range 0.25–5.0 ng/mL, which could be fitted
 391 into a linear regression equation: $y = -2E-06x + 4E-06$, $R = 0.9981$, where y is the
 392 value of current, and x is the Log of AFM₁ concentration. In the blank sample
 393 (without AFM₁), the HRP enzyme concentration is high and the redox reaction rate is
 394 fast, resulting in the faster current, so the absolute value of current is large. With the
 395 AFM₁ concentration increase in the sample, the HRP concentration decrease and the

396 reaction speed decrease, resulting in the slower current, so the absolute value of
397 current is smaller. The developed immunosensor demonstrated a sensitivity toward
398 AFM₁ with a detection limit at 0.09 ng/mL, which was calculated using 3SD of blank
399 samples. The sensitivity of the assembled immunosensor could meet the requirement
400 of the maximum residual levels in most countries, such as China, the EU, and so forth.
401 Especially the SPCEs were low cost and could be easily applied in developing
402 countries. There were other immunoassays using different kinds of electrodes for
403 AFM₁ detection. For example, Guo et al. constructed a covalent organic frameworks
404 TpBD on glassy carbon electrode using paminobenzoic acid and ethylenediamine as
405 connector, the developed electrochemical biosensor exhibited high selectivity and
406 sensitivity toward AFM₁ with a LOD of 0.15 ng/mL(Guo, Wang, Pang, Shen, Yang,
407 Ma, et al., 2021). Karczmarczyk et al. developed a strategy based on a competitive
408 immunoassay for the voltammetric detection of mycotoxins (OTA and AFM₁) using
409 modified gold screen printed electrodes with IC80 of 0.04 ng/mL for
410 AFM₁(Karczmarczyk, Baeumner, & Feller, 2017). Li et al. developed an inner filter
411 effect-based immunosensor incorporating fluorescent detection for AFM₁ residue
412 analysis with a LOD of 18.10 ng/kg (Li, Liu, Zhang, Luo, Lin, & Jiang, 2021). All the
413 established methods above obtained satisfied results, performing based on the
414 traditional antigens and antibodies reaction. In our work, the AIdnb were used to
415 replace the high toxin traditional antigens to modify the surface of working electrode.
416 In conclusion, compared with other kinds of electrodes, the SPCE used in this work
417 are not only modified with non-toxic alternative antigens to meet the needs of
418 environmental friendly, but also have high sensitivity. Moreover, SPCE has the
419 advantages of convenient use and low-cost, and has broad application prospects in
420 food safty detection.



421

422 Figure 4. (A) Electrochemical current response of the immunosensor for the detection of AFM₁ at
 423 0 ng/mL and 10 ng/mL using hydroquinone and (B) TMB; (C) Current response of immunosensor
 424 for the detection of AFM₁ (ng/mL): (a) 0, (b) 2.5, (c) 0.5, (d) 1.25, (e) 2.5, and (f) 5.0 ng/mL.
 425 The inset represents the linear curve of the immunosensor obtained for AFM₁ concentrations. (D)
 426 Electrochemical current response of the immunosensor for detecting AFM₁, AFB₁, ZEN, OTA, and
 427 FB₁ at a concentration of 10 ng/mL. The error bars were obtained from five parallel experiments.

428

429 3.5 Specificity, stability, and validation of the immunosensor

430 The specificity of the proposed immunosensor was investigated by measuring the
 431 current responses to AFB₁, ZEN, OTA, and FB₁ at a concentration of 10 ng/mL. The
 432 current response exhibited no decrease compared with the blank (PBS), indicating that
 433 the immunosensor had satisfactory specificity without nonspecific reaction with AFB₁,
 434 ZEN, OTA, and FB₁ (Fig. 4D). The electrode stability of the immunosensor was
 435 investigated by detecting the influence of electrode storage time on the current
 436 response value. After incubating the nanobody on the electrode, the same patch
 437 electrode was stored in the dark for different periods before detecting the blank

438 sample and the sample spiked with 10 ng/mL of AFM₁. It exhibited that 95% of the
 439 initial current signal was retained after storage at 4°C for 9 weeks, indicating that the
 440 immunosensor had good stability. Compared with the monoclonal antibody, nanobody
 441 exhibit thermal stability and showed better tolerance to organic solvents (He, Nie, Yan,
 442 Zhu, He, Li, et al., 2022; He, Wang, Li, Zhang, Lei, Zhang, et al., 2014). Thus, in
 443 theoretically, the nanobody modified electrode can be regenerated. However, the
 444 actual regeneration results should be proved by experimental data further.

445 We examined the performance of the immunosensor at different concentrations of
 446 AFM₁ (0.25, 0.5, and 1.0 ng/mL); the recoveries of the assays were 82.0%–108.0%
 447 (Table 1). We detected the intra-assay variability on the same day and the inter-assay
 448 variability on different days with the immunosensor to study the stability of the
 449 immunosensor. The intra-assay variability was given by the average of five replicated
 450 SPCEs. The inter-assay was given by the average of five replicated SPCEs on
 451 different days. The relative standard deviation was 10.1%–11.5% for intra-assays and
 452 10.2%–13.0% for inter-assays, which showed that it was feasible to determine AFM₁
 453 using the immunosensor.

454 **Table 1.** Recovery analysis of the immunosensor for AFM₁ in milk samples

	Spiked concentration (ng/mL)	Mean±SD (ng/mL)	Recovery (%)	RSD (%)
Intraday (n=5)	0.25	0.27	108.0	10.6
	0.5	0.41	82.0	11.5
	1.0	0.85	85.0	10.1
Interday (n=5)	0.25	0.26	104.0	12.6
	0.5	0.43	86.0	10.2
	1.0	0.83	83.0	13.0

455

456 Furthermore, 10 naturally contaminated milk samples collected from local farms were
 457 detected using the immunosensor and high performance liquid chromatography-mass
 458 spectrometry (HPLC-MS/MS). The results obtained from the analysis are summarized
 459 in Table 2. A total of 4 samples (40% of 10 samples) were found to contain AFM₁

460 with a concentration of lower than 0.47 ng/mL. The results showed a satisfactory
 461 agreement between the immunosensor and HPLC-MS/MS, indicating that the
 462 proposed method was highly appropriate for the screening and quantitation of AFM₁
 463 in milk.

464

465 **Table 2.** Validation of the immunosensor by HPLC-MS/MS for AFM₁
 466 in natural milk samples

Sample	HPLC-MS/MS (n=5)	Immunosensor (n=5)
	Mean (ng/mL)	Mean (ng/mL)
1	ND ^a	ND
2	0.40±0.001	0.43±0.56
3	ND	ND
4	0.46±0.005	0.47±0.72
5	ND	ND
6	0.38±0.004	0.40±0.51
7	ND	ND
8	ND	ND
9	0.26±0.008	0.28±0.69
10	ND	ND

467 ^aND: not detected.

468 **4. Conclusions**

469 In summary, a nontoxic electrochemical immunosensor was developed based on novel
 470 anti-idiotypic nanobody-functionalized SPCEs, and applied to identify AFM₁ using
 471 the immunosensor. Due to the specificity and stability of the nanobody, the proposed
 472 immunosensor exhibited excellent analytical performance with a linear range
 473 (0.25–5.0 ng/mL), as well as a low limit of detection value (0.09 ng/mL), meeting the
 474 requirement of the regulatory standard. In addition, the immunosensor showed

475 excellent recovery (%) in samples, proving that the immunosensor had high accuracy
476 and repeatability for the quantitative determination of AFM₁ by chronoamperometry.
477 The proposed immunosensor could be further used in the naturally contaminated
478 samples for AFM₁ detection.
479

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549 **Declaration of competing interest**

550 The authors declare that they have no known competing financial interests or personal relationships
551 that could have appeared to influence the work reported in this paper.

552

553 **Credit authorship contribution statement**

554 **Xiaoqian Tang:** Conceptualization, Investigation, Formal analysis, Writing – original
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556 **Gaëlle Catanante:** Supervision, Methodology. **Xiaorong Huang:** Validation.

557 **Jean-Louis Marty:** Conceptualization, Visualization. **Qi Zhang:** Supervision, Formal
558 analysis, Methodology. Supervision. **Hong Wang:** Validation, **Peiwu Li:** Project
559 administration, Funding acquisition.

560