

Modification-free amperometric biosensor for the detection of wild-type p53 protein based on the *in situ* formation of silver nanoparticle networks for signal amplification

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ABSTRACT

Sensitive and accurate quantification of wild-type p53 protein is of great importance for biological research and clinical diagnosis. Herein, a modification-free amperometric biosensor was proposed for sensitive detection of wild-type p53 protein by the signal amplification of silver nanoparticles (AgNPs) networks formed *in situ* on electrode surface. Double-stranded DNA (dsDNA) probe containing two consensus sites was immobilized on gold electrode surface to capture wild-type p53 protein. The cysteine thiol and amine groups on the exterior of the protein allowed for the attachment of bare AgNPs through the Ag–S or Ag–N interactions. Meanwhile, benzene-1,4-dithiol (BDT) molecules in solution triggered the assembly of more AgNPs on electrode surface through the Ag–S interactions, thus leading to the *in situ* formation of AgNPs networks for signal amplification. The target at the concentration as low as 0.1 pM can be readily determined. This method was further applied to determine wild-type p53 protein in spiked human serum and cell lysates with satisfactory results. Moreover, the biosensor is regenerative and does not require the modification of AgNPs with recognition element for signal readout. The modification-free strategy can potentially be applied to develop novel biosensors for detection of other biological macromolecules.

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1. Introduction

p53 protein, a tumor suppressor and transcription factor encoded by TP53 gene, plays important roles in multiple central cellular processes (e.g. DNA repair, cell cycle control and apoptosis) [1]. Concentration of wild-type p53 protein in normal cells is usually ultralow [2,3]. Mutation or inactivation of p53 gene in more than 50% of human cancers can cause the decrease of wild-type p53 protein and the accumulation of mutated p53 protein [4]. Following the death and destruction of tumor cells, p53 protein is released from cancer cells and circulated in the blood of cancer patients. Serum levels of p53 can accurately indicate tissue alteration at the gene and/or protein level [5]. Therefore, it is of great significance to achieve sensitive detection of wild-type p53 protein at ultralow concentration in human serum for clinical research and early diagnosis of cancers.

The classic methods for detection of p53 protein, such as enzyme-linked immunosorbent assay (ELISA) [6], electrophoretic mobility shift assay [7], immunostaining [8] and immunohistochemistry [9], generally involve in semi-quantification, low sensitivity and tedious experiment steps. In order to meet the increasing demand for sensitive detection

of p53 protein, researchers have focused considerable efforts to establish novel analytical methods, including electrochemistry [10–25], electrochemiluminescence [26], photoelectrochemistry [27], localized surface plasmon resonance (LSPR) [28], surface enhanced Raman scattering (SERS) [29,30] [31], and immunochromatographic test strip [3]. Among them, electrochemical biosensors have attracted considerable attention because of their intrinsic advantages of simple operation, high sensitivity, speed response and low cost. However, the biosensors mainly based on the specific antibody–antigen interaction cannot distinguish the wild-type p53 and mutant p53. Human p53 protein composing of 393 amino acid residues is functionally divided into three distinct domains: the transactivation domain at the N-terminus, the central DNA-binding domain, and the oligomerization domain near the C-terminus. It has been revealed that four wild-type p53 molecules can assemble into a tetramer in solution by their C-termini. The resultant p53 tetramer can bind to sequence-specific sites in the consensus double-stranded DNA (dsDNA) but the mutant p53 protein loses the binding ability [32]. Several groups have investigated the dsDNA/p53 interaction and developed a few biosensors with the consensus dsDNA as the acceptor for selective detection of wild-type p53 protein [33–43]. Herein, we attempt to develop a simple and sensitive electrochemical biosensor for specific detection of wild-type p53 protein using the consensus dsDNA receptor.

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To meet the urgent demand of ultrasensitive detection of biomarkers, electrochemical sensing strategies can be integrated with signal amplification to enhance the detection sensitivity. With the striking development of nanotechnology, various nanomaterials have been synthesized and used for designing of electrochemical biosensors with improving sensitivity, including noble metal nanoparticles (NPs), carbon nanostructures, metal-organic framework, metal oxide/sulfide NPs and other novel two dimensional (2D) nanomaterials [44–46]. The nanomaterials are usually employed as electrode modifiers to accelerate electron transfer and increase electrode surface area, as the carriers to load large numbers of enzyme labels or electroactive species, and/or as the reporters to amplify the electrochemical signal. For example, silver nanoparticles (AgNPs) modified with specific recognition element have been extensively utilized as the redox reporters to develop sensitive electrochemical biosensors for the analysis of cancer cells, DNA, microRNAs, proteins, drugs and metal ions because of their excellent electrochemical properties [47–57]. Wild-type p53 proteins in solution can form tetrameric oligomer. The tetramer can bind to dsDNA at the C-terminus with high binding affinity and each p53 monomer contains five surface accessible cysteine thiols on the exterior [38,58]. For this view, we developed an electrochemical biosensor for the detection of wild-type p53 protein based on the interaction of cysteine thiols and AgNPs. The signal was further amplified on the basis of benzene-1,4-dithiol (BDT)-induced *in situ* formation of AgNPs networks on electrode surface. The method has the advantages of high sensitivity and simplicity and does not require the modification of AgNPs nanolabels with recognition element for signal readout. Note that metal-free cysteine thiol and amine groups in most of proteins can attach onto the bare Au/Ag surface but dsDNA hybrids exhibit no interaction with bare Au/Ag nanoparticles [59]; the method may be extended to design novel biosensors by matching sequence-specific dsDNA acceptors.

2. Experimental

2.1. Chemicals and reagents

Tris(2-carboxyethyl)phosphine hydrochloride (TCEP), BDT, 6-mercapto-1-hexanol (MCH), tris-(hydroxymethyl)aminomethane hydrochloride (Tris-HCl), IgG, thrombin and bovine serum albumin (BSA) were purchased from Sigma-Aldrich (Shanghai, China). p53 ELISA kits were provided by SenBeijia Biological Technology Co., Ltd. (Nanjing, China). HPLC-purified oligonucleotides were ordered from Sangon Biotechn. Co., Ltd. (Shanghai, China). Their sequences were listed as follows: the thiolated single-strand DNA (ssDNA): 5'-HS(CH₂)₆-TTT TTA GAC ATG CCC AGA CAT GCC C-3' and the complementary DNA: 5'-GGG CAT GTC TGG GCA TGT CT-3'. Each p53 monomer binds to one-quarter site of the dsDNA with sequence of (A/T)(G/C)(A/T)(C/G)(A/T)(T/A)(G/C)(C/G)(C/G)(C/G). The DNA solution was prepared using TE buffer (20 mM Tris-HCl, 1 mM EDTA, pH 7.4). Prostate specific antigen (PSA) was provided by Linc-Bio Science Co. Ltd. (Shanghai, China). Serum sample was provided by Anyang Maternal and Child Health Care Hospital (Anyang, China). Silver nitrate (AgNO₃), ethylene diamine tetraacetic acid (EDTA), trisodium citrate and other reagents were obtained from Aladdin Reagent Company (Shanghai, China). All chemicals were of analytical grade and used without further purification. All solutions were prepared with ultrapure water (18.2 MΩ) obtained from a Milli-Q water purifying system (Milli-Q, Millipore).

2.2. Instruments

All electrochemical measurements were conducted on a CHI 660 electrochemical workstation (Shanghai CH instrument Inc., China). The conventional three-electrode system comprised the reference electrode (Ag/AgCl), the auxiliary electrode (platinum wire) and the work

electrode. Scanning electron microscopy (SEM) images were taken with a SU8010 scanning electron microscope (Hitachi, Japan).

2.3. Preparation of citrate-capped AgNPs

Citrate-capped AgNPs was synthesized according to our previously reported procedure [56]. In brief, 1 mL of 10 mM AgNO₃ solution and 1 mL of 10 mM trisodium citrate solution were added to 36.8 mL of ultrapure water under vigorous stirring. This is followed by addition of 1.2 mL of 10 mM freshly prepared NaBH₄. The solution color turned to yellow, indicating the formation of AgNPs. After stirring for additional 10 min, the solution was stored at 4 °C for use.

2.4. Preparation of sensor electrode

Prior to modification, the gold electrode (AuE) with 2 mm diameter was polished with alumina suspensions down to 0.05 μm, followed by cleaning ultrasonically the electrode with ethanol and water. After that, the cleaned electrode was transferred into the TE buffer solution containing 50 μM thiolated ssDNA probe, 0.1 M NaCl and 100 μM TCEP for 12 h at room temperature. The probes were immobilized on the electrode surface through the Au—S interactions. Subsequently, the modified electrode was rinsed thoroughly with water and soaked in a 1 mM MCH solution for 30 min to block the unmodified region of electrode and remove non-specifically adsorbed ssDNA (Fig. S1). Before the assay, the probe-covered electrode was incubated with 10 μM complementary DNA for 2 h in the TE buffer solution. Finally, the consensus dsDNA modified electrode was rinsed with TE buffer and used for the capture of wild-type p53 protein.

2.5. p53 protein detection

10 μL of wild-type p53 protein at a given concentration in phosphate buffer (10 mM, pH 7.2) was casted onto the consensus dsDNA-modified electrode surface for 20-min incubation. Then, the electrode was washed thoroughly with the buffer and subsequently exposed to 25 μL of AgNPs suspension in a homemade plastic cell. After incubation for 5 min, 25 μL of BDT solution was added to the AgNPs suspension and incubated for 15 min again. After being rinsed with water, the electrode was placed in 1 M KCl solution for linear-sweep voltammetry (LSV). After each assay, the sensor electrode was regenerated with 10 mM NaOH.

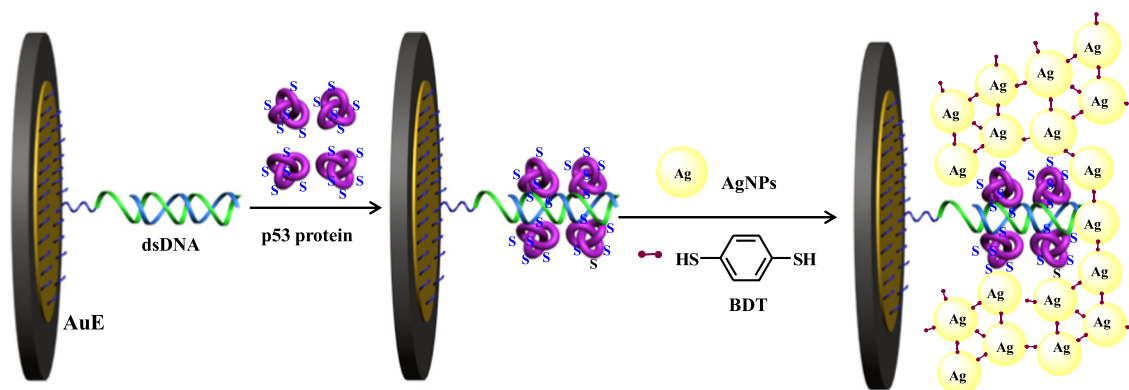
2.6. Analysis of cell lysates

The procedure for the preparation of cell lysates followed the previous report [38]. Briefly, MCF-7 and HeLa cancer cells and PC12 nerve cells were incubated in DMEM (Dulbecco's Modified Eagle Medium) for 24 h. The living cells were washed three times with ice-cold phosphate buffer and then lysed with lysis buffer. The lysed cells were collected and centrifuged at 4 °C for 10 min. The supernatants were diluted by 10-fold with phosphate buffer and then analyzed immediately.

3. Results and discussion

3.1. Detection principle of the method

The modification-free amperometric biosensor for detection of wild-type p53 protein was developed based on the high affinity of wild-type p53 protein toward consensus dsDNA. The detection principle is illustrated in Scheme 1. The mixed dsDNA/MCH self-assembled monolayers (SAMs) are performed on the surface of gold electrode as the bioreceptor to capture wild-type p53 protein and eliminate the non-specific adsorption. After the formation of the p53 protein/dsDNA complex, the five metal-free cysteine thiols and amine groups at the surface



Scheme 1. Schematic representation of the biosensor for detection of wild-type p53 protein by the *in situ* formation of AgNPs networks for signal amplification.

of wild-type p53 protein allow for the capture of AgNPs through the Ag—S or Ag—N interactions. The signal was then amplified by BDT-induced assembly of AgNPs *via* the formation of Au—S bonds. In brief, the surface-anchored AgNPs allow for the attachment of BDT molecules through the Ag—S interactions. Then, the anchored BDT molecules facilitate the capture of more AgNPs through the same Ag—S interactions. Thus, BDT and AgNPs in solution circularly recruit more dissociative AgNPs on the electrode surface, leading to the *in situ* formation of the AgNPs-based networks. Finally, the network architecture of AgNPs produced an amplified electrochemical signal by the solid-state Ag/AgCl process. Note that dsDNA cannot adsorb onto the bare Au/Ag surface [59]; the dsDNA/MCH-covered electrode will not capture AgNPs, thus decreasing the nonspecific adsorption of signal labels and improving the detection specificity.

3.2. Feasibility

Electrochemical impedance spectroscopy (EIS) is sensitive to small charge change at the solid-liquid interface. The stepwise fabrication process of sensor electrode was first characterized by EIS technique. The Nyquist curves corresponding to the modification of electrode surface at different steps were presented in Fig. 1A. The semicircle diameter in high-frequency region was equal to the electron-transfer resistance (R_{et}), displaying the electron transfer dynamics of $[\text{Fe}(\text{CN})_6]^{3-/4-}$ on the electrode surface. The bare gold electrode showed a relatively small R_{et} (curve a). After immobilization of ssDNA onto the electrode surface, the R_{et} increased greatly (curve b), which is due to the repulsion between negatively charged ssDNA backbone and $[\text{Fe}(\text{CN})_6]^{3-/4-}$. When the ssDNA-modified electrode was covered with MCH, a

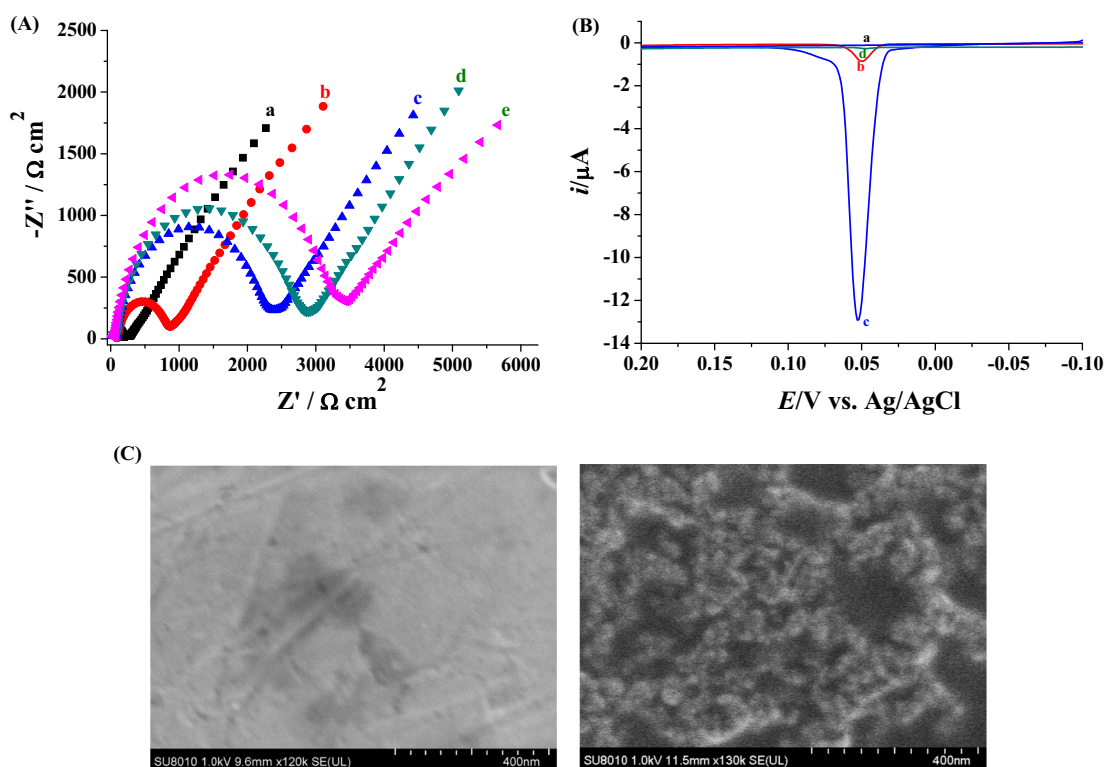


Fig. 1. (A) EIS of AuE (a), AuE/ssDNA (b), AuE/ssDNA/MCH (c), AuE/dsDNA/MCH (d) and AuE/dsDNA-p53/MCH (e). (B) LSV responses acquired at the sensor electrode after incubation with various solutions: AgNPs/BDT (a), wild-type p53 + AgNPs (b), and wild-type p53 + AgNPs/BDT (c). Curve d corresponds to that acquired at AuE/ssDNA/MCH after successive incubation with wild-type p53 protein and AgNPs/BDT. (C) SEM images of the p53-captured electrode before (left) and after (right) incubation with AgNPs/BDT.

significant increase of R_{et} (curve c) was observed. The result is attributed to the steric hindrance and electrostatic repulsion by the compact ssDNA/MCH SAMs. Hybridization of complementary DNA to form the consensus dsDNA on electrode surface led to a further increase in R_{et} (curve d). In addition, the R_{et} increased after the dsDNA/MCH-covered electrode was incubated with wild-type p53 protein, indicating that the target was captured by the sensor electrode.

Fig. 1B shows the LSV responses of the dsDNA/MCH-covered electrode after incubation with various solutions. Without capture of target p53 protein, no apparent LSV peak was detected even if the electrode was incubated with AgNPs/BDT (curve a), indicating that no or undetectable AgNPs were attached on the electrode surface. This is acceptable since dsDNA cannot interact with AgNPs or BDT. However, a small LSV peak was observed when the electrode was successively incubated with target p53 protein and AgNPs only (curve b), demonstrating that a few AgNPs were bound to the sensor electrode. Interestingly, when the p53-captured electrode was incubated with mixture of AgNPs and BDT, an intensive and well-defined LSV peak at 50 mV was observed (curve c). The sharp peak is generated from the highly characteristic solid-state Ag/AgCl reaction from AgNPs since no LSV peak was observed in the absence of AgNPs. The peak current is approximately 16-fold higher than that incubated with AgNPs only (curve b), indicating that the signal was amplified by the formation of AgNPs-based networks. The result was also confirmed by SEM (Fig. 1C): obvious AgNPs networks were found when the sensor electrode was incubated with target p53 protein and AgNPs/BDT. To prove that the binding of target is dsDNA-dependent, the ssDNA-modified electrode was incubated with p53 protein and the mixture of AgNPs and BDT. As a result, no discernible peak was observed (curve d). This confirmed that the nonspecific interaction between ssDNA and wild-type p53 protein was negligible. Therefore, the distinct and amplified LSV responses enabled the proposed electrochemical biosensor with the potential ability to quantify wild-type p53 protein.

3.3. Optimization of experiment conditions

BDT as the linker can trigger the assembly of metal nanoparticles by the metal-S interactions. The BDT concentration may have a great influence on the AgNPs assembly and the performance of the biosensor. Thus, we first investigated the influence of BDT concentration on the peak current. As shown in Fig. S2, the anodic peak current (i_{pa}) increased with the increasing concentration of BDT and reached to the maximum value at 5 μ M. When the BDT concentration was too low, the signal was small, demonstrating that only few AgNPs network architecture was produced on the electrode surface. However, when the concentration was higher than 5 μ M, i_{pa} began to fall down. This can be attributed to the competition between free BDT and cysteine thiols of p53 protein to bind with AgNPs. The incubation time for BDT-induced assembly of AgNPs was also optimized. As shown in Fig. S3, the current was intensified with the increase of incubation time and began to level off beyond 15 min, which is indicative of the completion of the assembly of AgNPs on electrode surface.

The dsDNA hybrid and dsDNA/p53 complex are susceptible to denaturation and disassembly at low pH. By washing the electrode with acidic solution, we found that the electrode remained over 90% of the initial response after ten regeneration cycles. This suggested that the sensor electrode can be conveniently regenerated. Thus, multiple samples can be measured at one electrode. The relative standard deviations (RSDs) for assays of each sample at three chosen electrodes are all below 10.7%, indicating that the proposed method has acceptable reproducibility. Thus, multiple electrodes can also be prepared in parallel for assays of different samples.

3.4. Sensitivity

The normal concentration of p53 protein in serum circulatory system is very low. Thus, high sensitivity is required for the practical

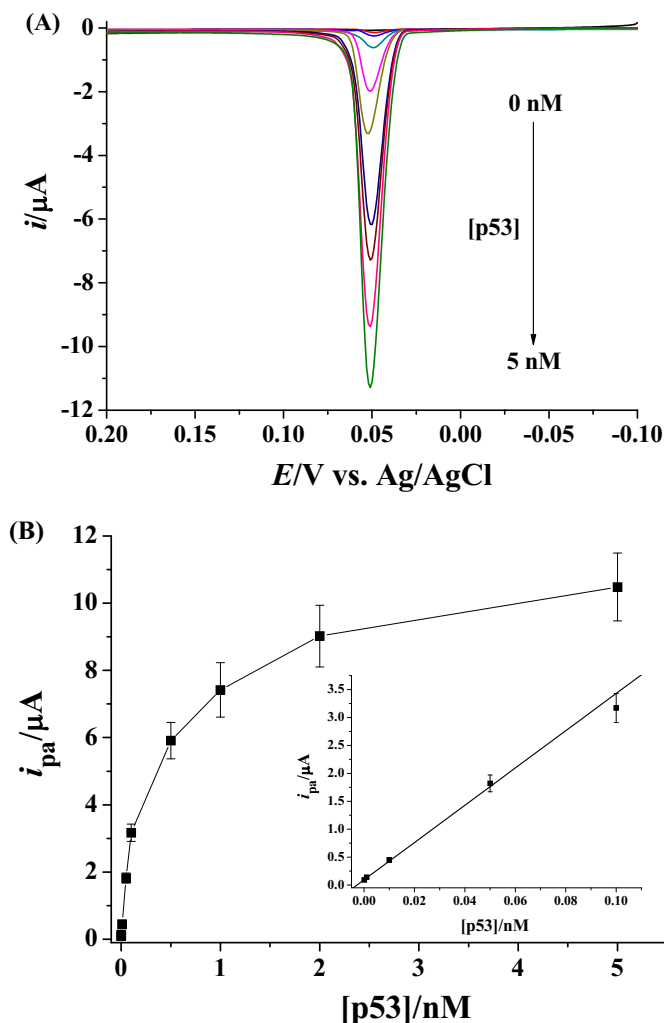


Fig. 2. (A) LSV responses for detection of various concentrations of wild-type p53 protein (0, 0.0001, 0.001, 0.01, 0.05, 0.1, 0.5, 1, 2 and 5 nM). (B) Dependence of i_{pa} on p53 concentration. The inset shows the linear portion of the curve.

application of biosensor for detection of wild-type p53 protein. To evaluate the sensitivity and detection limit of the method, LSV responses for the detection of various concentrations of target protein were collected. As shown in Fig. 2A, the peak current increased with the increase of target protein from 0 to 5 nM, demonstrating that the attachment of AgNPs on electrode was dependent upon p53 protein concentration. Fig. 2B displays the dependence of i_{pa} on the concentration of target protein. The linear regression equation was fitted as $i_{pa} = 33.4[p53] \text{ (nM)} + 0.094$ in the range of 0.1–100 pM. The target at the concentration as low as 0.1 pM was readily measured. The value is lower than that achieved by surface plasmon resonance (SPR) (1.06 pM) [39], chemiluminescence (3.8 pM) [41], fluorescence (8 pM) [43], electrochemistry (2.2 pM) [38], and enzyme-linked electrochemical amperometric immunosensor (14 pM) [16]. The sensitivity is comparable to (or even lower than) that of sandwich-type immunoassays of p53 proteins by different signal amplification strategies (Table 1). The high sensitivity can be attributed to the low background current, the well-defined silver stripping peak and the signal amplification of AgNPs-based networks. Moreover, the method does not require the pre-labeling of nanolabels with recognition element for signal readout, thus simplifying the test steps, reducing the detection complexity and decreasing the assay cost.

Table 1
Analytical performances of sandwich-type electrochemical biosensors for p53 protein detection.

Electrode modifier	Signal label	Detection limit	Linear range	Ref.
SAMs	Fc-AuNPs	2.2 pM	Non-linear	[38]
AuNPs/GO	HPR	0.03 pM	0.2–200 pM	[10]
Au NPs	GO-HPR	10 pM	20–2000 pM	[12]
Nano-SPEC	AP	50 pM	200–50,000 pM	[18]
MWCNT-PA6	HRP	0.02 pM	0.04–40 pM	[19]
Graphene-CHI	HRP	2 pM	4–200 pM	[60]
SAMs	AgNPs	0.1 pM	0.1–100 pM	This work

Abbreviations: Fc-AuNPs, ferrocene-capped gold nanoparticles; GO, graphene oxide; HRP, horseradish peroxidase; Nano-SPEC, nanostructured screen-printed carbon electrode; AP, alkaline phosphatase; MWCNT, multi-walled carbon nanotube; PA6, doped nylon 6; CHI, chitosan.

3.5. Selectivity and anti-interference

Selectivity is a critical factor to evaluate the applicable capability of electrochemical biosensors. To demonstrate the selectivity of the proposed method, we challenged the sensing system with four common proteins (IgG, thrombin, PSA and BSA) instead of wild-type p53 protein under the same experimental conditions. As shown in Fig. 3, the current caused by the tested proteins are close to background level, even at the concentration of at least 100-fold higher than that of target protein. Meanwhile, when introducing these non-target proteins as interferences to the detection system, there was no significant difference in the current value by comparison to that of target p53 protein only. The high selectivity and good anti-interference ability should be attributed to the specific interaction between wild-type p53 protein and dsDNA and the high antifouling ability of the dsDNA/MCH SAMs toward AgNPs.

3.6. Real sample analysis

To indicate the accuracy and amenability of the biosensor for real sample analysis, we determined the concentration of wild-type p53 protein in a serum sample. As a result, no or undetectable target protein was found. Thus, standard protein samples at the final concentrations of 1, 10 and 50 pM were spiked to the serum and then analyzed by the method. According to the above calibration curve, the concentrations of target protein were deduced to be 0.9 pM, 10.3 and 49.1 pM, respectively. The found contents are close to those of the known values and the recoveries vary in the range of 90–103%, indicating that the method is reliable for analysis of p53 protein in biological matrix sample. Thus,

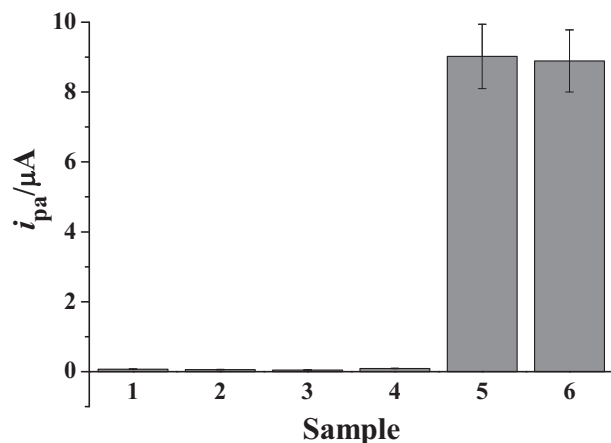


Fig. 3. Selectivity and anti-interference. Bar 1, IgG; bar 2, thrombin; bar 3, PSA; bar 4, BSA; bar 5, p53; bar 6, the mixture of 1–5. The concentration of wild-type p53 protein is 2 nM and that of other proteins is 0.1 μg/mL.

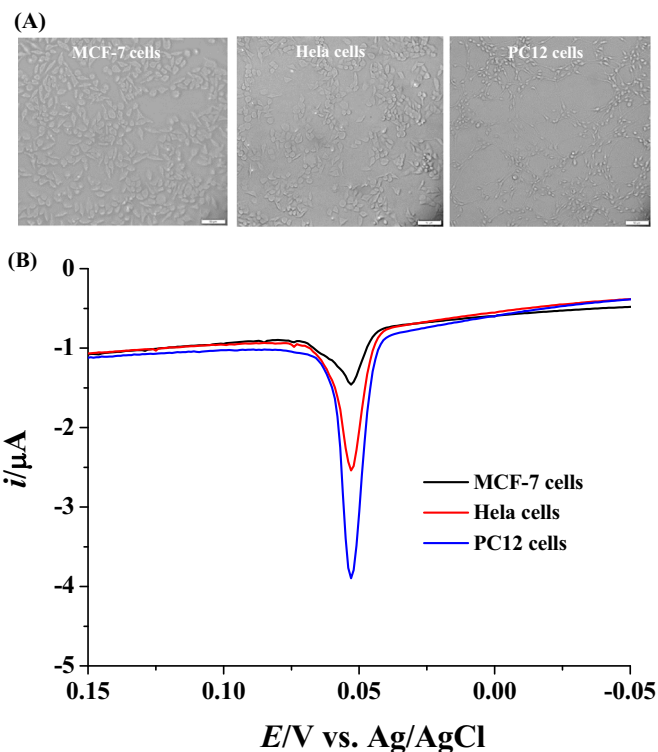


Fig. 4. (A) Confocal images of living cells and (B) LSV responses for analysis of cell lysates.

this method was further used to detect wild-type p53 protein in different cell lysates, including cancer cells of MCF-7 and HeLa and normal nerve PC12 cells. Fig. 4A depicts the confocal images of these living cells. Fig. 4B shows the LSV responses for detection of wild-type p53 protein in the supernatants of various cell lines. The found concentration of wild-type p53 protein in the cancer cells was much lower than that in the normal cells (Table 2). Meanwhile, the contents of total p53 proteins (wild-type and mutant combined) were also determined by ELISA kits with a calibration curve of $OD_{450} = 0.01 + 0.004[p53]$ (pM) (Fig. S4). It was found that the total p53 protein concentration in the cancer cells is higher than that in the normal cells. From the deduced difference between wild-type and total p53 proteins, the content of mutant p53 protein in the cancer cells was greatly higher than that in the normal cells. Note that the ratio between wild-type and mutant p53 proteins is a preferred criterion for cancer diagnosis; thus, the biosensor may be used as a promising tool for diagnosis of cancer development by measuring the level of wild-type p53 protein.

4. Conclusion

In summary, we developed a modification-free electrochemical biosensor for sensitive detection of wild-type p53 protein. AgNPs networks formed *in situ* on the sensor electrode surface were utilized as the electroactive reporters for signal readout. The well-defined electrochemical signal originated from the solid-state Ag/AgCl reaction of AgNPs-based networks. The biosensor exhibited a wide linear range and a detectable concentration as low as 0.1 pM. In contrast to other nanomaterials-based strategies for signal amplification, the biosensor

Table 2
Results for analysis of p53 protein in cell lysates.

Cell lysates	Wild-type p53 (pM)	Total p53 (pM)
MCF-7 cells	20.8 ± 0.21	102.4 ± 7.36
HeLa cells	49.5 ± 0.31	127.8 ± 6.52
PC12 cells	83.2 ± 6.82	85.5 ± 2.65

is simple and cost-effective and obviates the modification of nanolabels for molecular recognition. We believe that the method has promising applications in early clinical diagnosis and prognosis.

Declaration of competing interest

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

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.ijbiomac.2020.04.271>.

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