



An CuInS₂ photocathode for the sensitive photoelectrochemical determination of microRNA-21 based on DNA–protein interaction and exonuclease III assisted target recycling amplification

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Abstract

A photocathode is described for the determination of microRNA-21 by using CuInS₂ as an active photocathode material. Exonuclease III assisted target recycling amplification was employed to enhance the detection sensitivity. The TATA-binding protein (TBP) was applied to enhance steric hindrance which decreases the photoelectrochemical intensity. This strategy is designed by combining the anti-interference photocathode material, enzyme assisted target recycling amplification and TBP induced signal off, showing remarkable amplification efficiency. Under the optimized conditions, the detection limit for microRNA-21 is as low as 0.47 fM, and a linear range was got from 1.0×10^{-15} M to 1.0×10^{-6} M.

Keywords Photoelectrochemistry · TATA-binding protein · Photoresponse · Nanoflowers · Electrode · Aptamer

Introduction

The photoelectrochemical (PEC) process refers to photovoltaic conversion of photoelectrochemically active material from electron excitation to charge transfer under an applied light [1, 2]. Due to its separate source for photoirradiation and electrochemical detection, PEC assay has received substantial attention due to its low background signal and high sensitivity [3, 4].

From the way of electron transmission, PEC assay can be classified into photoanode assay and photocathode assay [5].

The n-Type semiconductors such as CdSe, CdS, ZnO and TiO₂ [6, 7] are mainly used in photoanode assay. These materials rely on the ejection of electrons from the conduction-band to the electrode, which supply of electrons from electron donors in solution. However, the presence of sacrificial electron donor may compete with reductive ingredient (for example ascorbic acid, dopamine, H₂O₂ and thio compounds). In this case, the false positive signal can be appeared when the detection is carried out in the real samples [8–10]. The p-type semiconductors (such as CuS, PbS, BiOI and CuInS₂) are always used as photoactive species for photocathode assay.

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The electrons transfer from the conduction-band to electron acceptor, and then the electrons are supplied from the electrode to the valence-band holes. In this process, electron acceptor (commonly use dissolved O_2) instead of electron donor is required for cathodic photocurrent. Hence, the reductive component in the complex samples cannot interfere with the cathodic photocurrent. On the other hand, holes are used as carriers for electron acceptor in photocathode assay, which lead to lower photocurrent intensity compared with that of photoanode assay [11, 12]. Thus, highly active photocathode material has attracted much attention as enhancing the detection sensitivity. Among these p-type semiconductors, $CuInS_2$ is highly attractive due to its small band gap and a large optical absorption coefficient [13, 14].

From the aspect signal format, the main established strategies for PEC assay can be classified as steric hindrance [15], biocatalytic precipitation [16], in situ generation of electron donor/acceptor [17], quenching effect [18], DNA conformational change [19] and introduction of photoelectric active materials [20]. Almost all these strategies utilize DNA for constructing PEC methods because DNA strands can bind with target molecular strongly and specifically. The interaction between DNA and protein plays important roles in controlling a variety of biological activities. Several works have been carried out on TATA-binding protein (TBP), which can bind to TATA sequence specifically [21–23]. With the advantage of the steric hindrance effect of TBP and easy amplification effect of DNA, a cathodic PEC method for detecting microRNA-21 was designed. In this strategy, $CuInS_2$ is used as an active photocathode material and TBP is employed to enhance steric hindrance.

It has been reported that microRNA-21 is upregulated in many cancers, such as lung cancer, breast cancer and pancreatic cancer et al. [24]. However, it is difficult for the microRNA detection due to the instability, low content and short length of microRNA, as well as the similar sequence in microRNA families [25]. Various kinds of methods were designed for microRNA detection, which had difficulty meeting the requirements for high sensitivity [26–28]. To increase the detection sensitivity, exonuclease aided recycling was applied here. At the same time, target RNA was converted to DNA for further detection. The principle of PEC method was described in Scheme 1. Firstly, cathodic photoelectrochemical materials $CuInS_2$ microflowers were dropped on the ITO glass, and heated to give a dense film. Then chitosan solution was added which provides amino groups for later modification and protects the $CuInS_2$ on the ITO electrode. Subsequently, carboxyl group modified DNA₂ was coupled on the electrode after reaction with chitosan through amide bond. If hairpin DNA₁ is recognized by microRNA-21, the enzyme Exo III cuts off the blunt 3' terminus of DNA₁. This triggers the recycling of microRNA-21 and the release of DNA₃ which can hybridize with DNA₂ on the electrode. Finally, the DNA₂-DNA₃

duplex is recognized by TBP. The decrease of photocurrent caused by steric hindrance can be used to quantify microRNA-21.

Experimental section

Materials and reagents

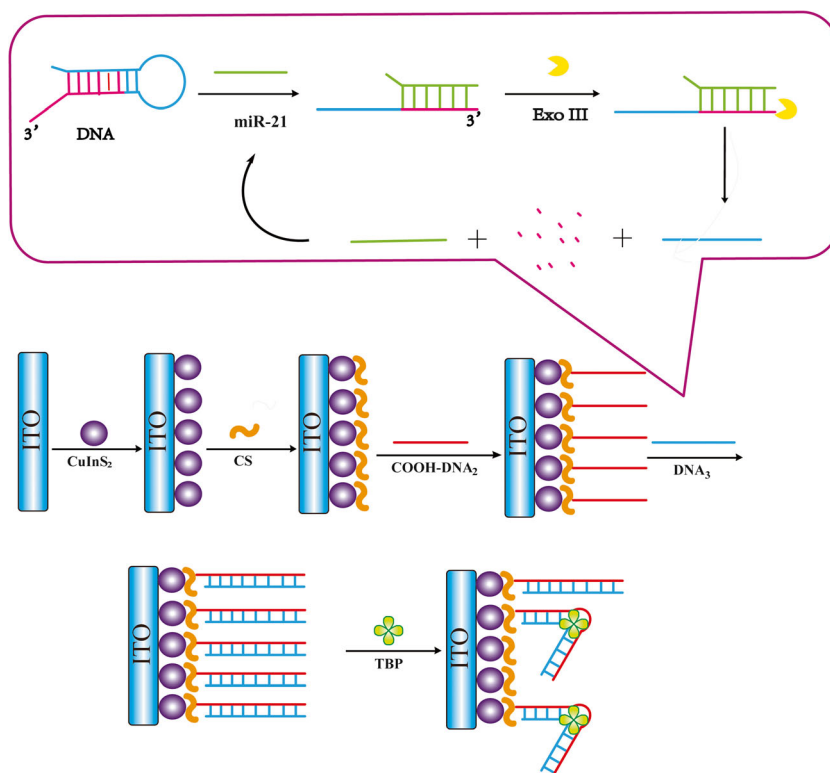
ITO electrodes were provided by South China Science and Technology Co., Ltd. (Yiyang, China) (website: <https://www.h-nxc.com>), with sheet resistance $\leq 10 \Omega/\text{square}$. Copper Chloride ($CuCl$) and indium chloride ($InCl_3$) were purchased from Sigma-Aldrich (website: <https://www.sigmaaldrich.com/china-mainland.html>). Thiourea (H_2NCSNH_2) and chitosan (CS) with degree of deacetylation $>95.0\%$ were the products of Aladdin Reagent Co., Ltd. (Shanghai, China) (website: <http://www.aladdin-e.com>). TATA-binding protein (TBP) and Exonuclease III (Exo III) were ordered from Sangon Biotech Co., Ltd. (Shanghai, China) (website: <http://www.sangon.com>). All other reagents used in the experiment were analytical grades and used directly. DNA oligonucleotides were gained from Sangon Biotech Co., Ltd. (Shanghai, China) (website: <http://www.sangon.com>) and the base sequences were as followed:

DNA₁: 5'-CCTCGACTGAGGTATAAAAGTCCTCC
TCAGTCTGATAAGCTA-3';
DNA₂: 5'COOH(CH₂)-AG GACTTTTATACCTC
AGTCGAGG-3';
DNA₃: 5'-CCTCGACTGAGGTAT AAAAGTCCT-3';
microRNA - 21 : 5' -
UAGCUUAUCAGACUGAUGUUGA-3';
M1 microRNA -21: 5'-UAGCUUAUAAGAC
UGAUGUUGA-3';
M2 microRNA -21: 5' -UAGCUUAU
AACACUGAUGUUGA-3';
M3 microRNA -21: 5' -UAGCUUAU
AACCCUGAUGUUGA-3';
miR-155: 5'-UUA AUGCUAAUCGUGAUAGGGU-
3';
TBA: 5'-GGTTGGTGTGGTTGG-3';
Apt-PDGF: 5'-TCAGGCTACGGCACG TAGAG
CATCACCATGATCCTG-3'

Instruments

The electrochemical impedance spectroscopy (EIS) and photocurrent were performed on an electrochemical workstation (Zahner Zennium, Germany) using ITO conductive glass, saturated $Ag/AgCl$ and Pt wire as working electrode, reference electrode and auxiliary electrode, respectively. The UV-vis

Scheme 1 Schematic Illustration of the PEC assay for microRNA -21 detection



absorption spectra were taken using a Cary 50 UV/vis-NIR spectrophotometer (Varian). Powder X-ray diffraction (XRD) patterns were recorded on an X-ray diffractometer with monochromatized Cu K α radiation (Brook AXS, Germany). Scanning electron microscopy (SEM) image was taken with a JEM-7500F scanning electron microscope (JEOL Ltd., Japan).

Preparation of CuInS₂ microflowers and modification of ITO electrode

CuInS₂ microflowers were synthesized according to previous method with a little modification [29]. Briefly, to 100 mL Teflon-lined autoclave, 40 mL of the ethylene glycol solutions containing 0.03 M InCl₃, 0.03 M CuCl, and 0.12 M H₂NCSNH₂ was added. The mixture was heated at 200 °C for 24 h, followed by washing with ethanol and deionized water and dried at 100 °C over night to give power products.

The ITO electrodes were ultrasonically cleaned in acetone, 2.0 M NaOH and deionized water successively for 20 min each, and then dried with nitrogen gas.

12.0 mg CuInS₂ product was dissolved in 2.0 mL deionized water and ultrasonicated to obtain homogeneous suspension. Then 20.0 μ L suspension was coated onto a bare ITO slices and dried at room temperature. The gained electrode was heated at 170 °C for 2.0 h followed by gradual cooling to room temperature. Subsequently, 20 μ L of 0.1 wt% CS solution containing 1% acetic acid was spread onto the

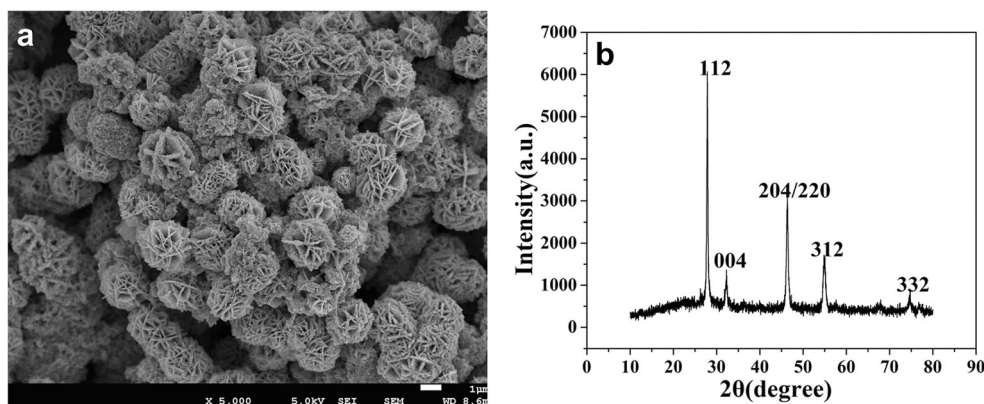
electrode. Finally, the electrode was dried at 50 °C, and washed with deionized water for three times to give ITO/CuInS₂/CHIT electrode.

Detection of microRNA -21

Carboxyl functionalized DNA₂ was assembled on ITO electrode surface through amidation coupling reactions. Firstly, 10 μ L DNA₂ was mixed with 90 μ L PB buffer containing 7.0 mg EDC and 3.5 mg NHS. Then 25 μ L mixture was spread on the electrode surface and incubated at 4 °C overnight. After washing with 10 mM PB, ITO/CuInS₂/CHIT/DNA₂ electrode was got.

At the same time, Exo III-assisted amplification was performed using 66 mM Tris-HCl (containing 0.66 mM MgCl₂, pH = 8.0) as a reaction buffer. Specifically, 500 nM probe DNA₁ was incubated with different concentration of target microRNA -21. 60 U Exo III was added to the mixture and incubate at 37 °C for 2 h. Then the enzyme was deactivated after heating at 75 °C for 10 min. Subsequently, 25 μ L of mixture was dropped on the electrode surface and incubated for 2 h at 37 °C. Finally, 25 μ L of 1000 ng mL⁻¹ TBP was added onto the surface of the above electrode and incubated at 37 °C for 1.0 h in a humid condition. After each modification, the electrode was thoroughly washed for three times. The photocurrent measurements were performed in 0.1 M PB at a potential of 0 V. An excitation light source with a wavelength of 415 nm was switched on and off every 20 s.

Fig. 1 **a** SEM images and **b** XRD patterns of synthesized CuInS_2



Results and discussion

Characterization of materials

The gained CuInS_2 was characterized by SEM. In Fig. 1a, spherical particles of microflowers morphology were observed with average size of 1.5–2.0 μm . Such rough surface caused by the layer structure of CuInS_2 provides a lot of surface area for photogenerated electron exchange. As a result, PEC signal can be enhanced accordingly. The crystal structure of gained CuInS_2 microflowers was further identified by XRD. As presented in Fig. 1b, five diffraction peaks are observed, which are assigned to the (112), (004), (204/220), (312), (332) crystal planes of CuInS_2 [JCPDS Card No. 27–0159]. So, pure crystalline CuInS_2 was successfully synthesized. These sharp peaks indicate the excellent crystallinity character of synthesized CuInS_2 .

The feasibility of the designed strategy

To illustrate the feasibility of the PEC assay, the photocurrent responses for the assemble processes were measured in 0.1 M PB (pH 7.4). As presented in Fig. 2, a remarkable photocathode response of ITO/ CuInS_2 is showed, due to the efficient charge transport and broad visible light absorption of the CuInS_2 (curve a) [15]. After blocking with CS, which can protect the CuInS_2 from dissolving to the detection solution as well as providing amino-group for further modification, the photocurrent intensity decreased dramatically (curve b). When carboxylated DNA_2 was modified on the electrode (curve c) and hybridized with DNA_3 (curve d), the photocurrent signal decreased slightly owing to the steric hindrance effect of DNA. After incubating with TATA-binding protein (TBP), the photocurrent intensity decreases obviously (curve e). This might be explained that TBP can bend the TATA sequence at an 80 degree angle through inserting amino acid side chains to base pairs of dsDNA sequence [22, 23], which blocked the transfer of electrons and led to a sharp decay in photocurrent intensity. Therefore, the characteristics of the

photocurrent variation demonstrate the feasibility of the designed method.

The fabrication step of the PEC nanoprobe was also characterized by EIS. As exhibited in Fig. 3, the unmodified ITO shows a small semicircle (curve a). After modification of CuInS_2 microflowers, the Ret value increased (curve b), indicating the poor conductivity of CuInS_2 . When CHIT was coated, the Ret value further increased due to the non-conductivity of chitosan (curve c). The assembly of the carboxyl functionalized DNA_2 (curve d) and hybridization with DNA_3 (curve e) inhibited the charge transfer furtherly, owing to the steric hindrance and the electrostatic repulsion between the negatively charged $\text{DNA}_2/\text{DNA}_3$ phosphoric acid backbone and the negatively charged $\text{Fe}(\text{CN})_6^{3-/4-}$. After recognition by TATA binding protein, the Ret value increased greatly (curve f), indicating the recognition between the protein and its aptamer as well as subsequent steric hindrance. Therefore,

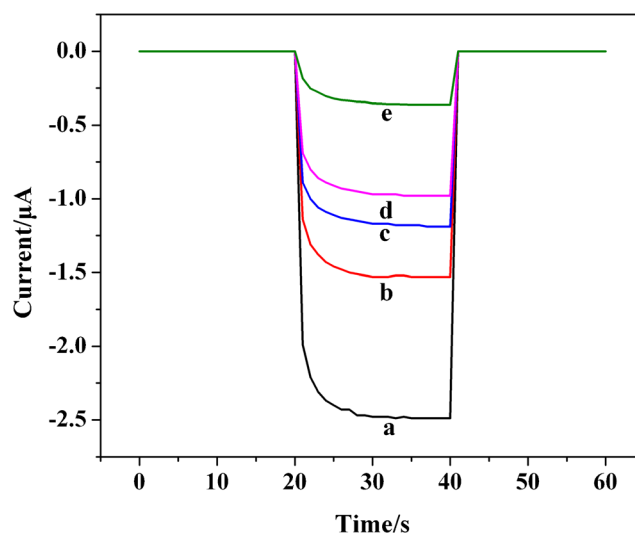


Fig. 2 Photocurrent responses of different ITO electrodes recorded in 0.10 M PB at a bias potential of 0 V for (a) ITO/ CuInS_2 ; (b) ITO/ CuInS_2 /CS; (c) ITO/ CuInS_2 /CS/ DNA_2 ; (d) ITO/ CuInS_2 /CS/ DNA_2 / DNA_3 ; (e) ITO/ CuInS_2 /CS/ DNA_2 / DNA_3 /TBP incubated with 1.0×10^{-6} mol/L microRNA -21. The PEC tests were performed in 0.10 M PB (pH = 7.4) at a bias voltage of 0 V with 415 nm light excitation

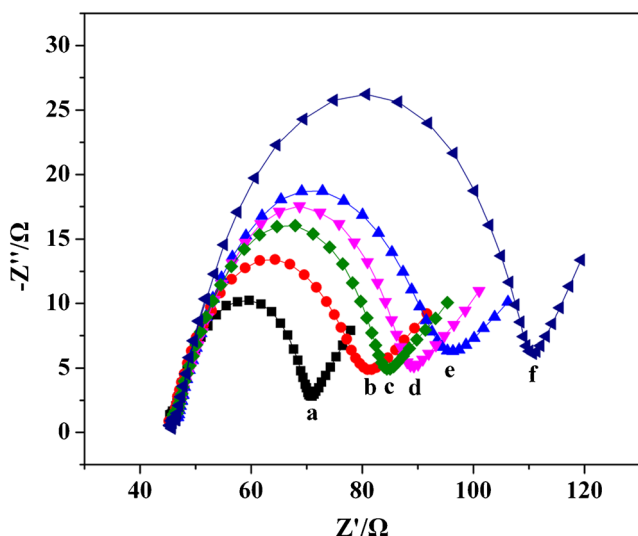


Fig. 3 Nyquist plots of different electrodes in 0.1 mol/L KCl containing 5.0 mM $K_3Fe(CN)_6/K_4Fe(CN)_6$ (1:1): **a** bare ITO substrate; **b** ITO/CuInS₂; **c** ITO/CuInS₂/CS; **d** ITO/CuInS₂/CS/DNA₂; **e** ITO/CuInS₂/CS/DNA₂/DNA₃; **f** ITO/CuInS₂/CS/DNA₂/DNA₃/TBP incubated with 1.0×10^{-6} M microRNA-21. The frequency range for EIS measurements was from 0.01 to 100,000 Hz at an applied potential of 0.2 V

the characteristics of the ESI also demonstrate the successful preparation the electrode interface.

The characterization of enzymatic cleavage by polyacrylamide gel electrophoresis (PAGE)

Exo III has been widely used to recycle the target with signal amplification. Exo III recognizes a blunt 3' terminus in double stranded nucleic acid substrate including duplex DNA-RNA and DNA-DNA [30–33]. The DNA strand in the duplex with 3'-flat or concave terminal is digested by Exo III to produce short oligonucleotide fragments, which released the target

RNA with recycle for further improving sensitivity in various assay. To demonstrate the enzymatic cleavage process, polyacrylamide gel electrophoresis (PAGE) was applied. As shown in Fig.S1, lane a and lane b represent stem loop probe DNA₁ and target microRNA -21, respectively. The bright band in lane c is attributed to the hybridization product of DNA₁ with target microRNA -21. When incubating the DNA₁/ microRNA -21 hybrid with Exo III, DNA₁ was cleaved from blunt 3' terminus to release the DNA₃ and intact microRNA -141 simultaneously, which has similar number of bases (lane d). So, the image of PAGE shows the cycle amplification aided by Exo III digestion was performed successfully.

Sensitivity of the PEC assay

The concentrations of Exo III and TBP were optimized to improve the performance of designed PEC assay. The results (Fig. S2, supporting information) suggest an optimized concentration of 60 U for Exo III and 1000 ng·mL⁻¹ for TBP. Under these optimum conditions, target microRNA 21 was detected quantitatively. As shown in Fig. 4, with increasing concentration of microRNA, the produced photoelectrochemical intensity of ITO/CuInS₂/CHIT/DNA₂/DNA₃ decrease accordingly. The photocurrent response is proportional to the logarithm of microRNA concentrations across the range from 1.0×10^{-15} M to 1.0×10^{-6} M. The linear regression equation was expressed as I (nA) = $-0.0937 \lg C_{\text{microRNA}} (M) - 0.202$, and the correlation coefficient (R^2) was 0.999. The detection limit of microRNA -21 is estimated to be 0.47 fM ($S/N = 3$). A series of eleven repetitive measurements of 1.0×10^{-10} M microRNA 21 is used for estimating the precision, and the standard deviation (SD) is 0.27. Compared with some previous reports for microRNA -21

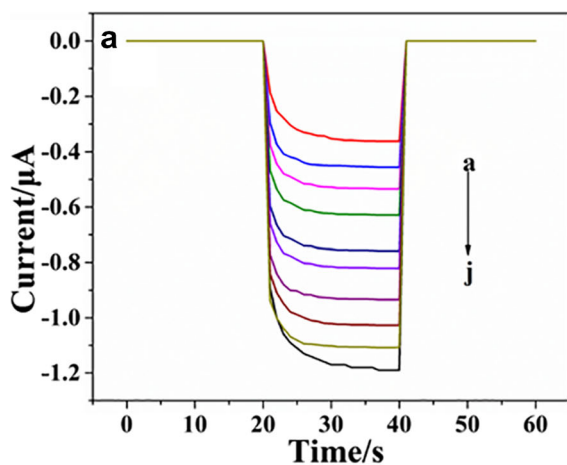
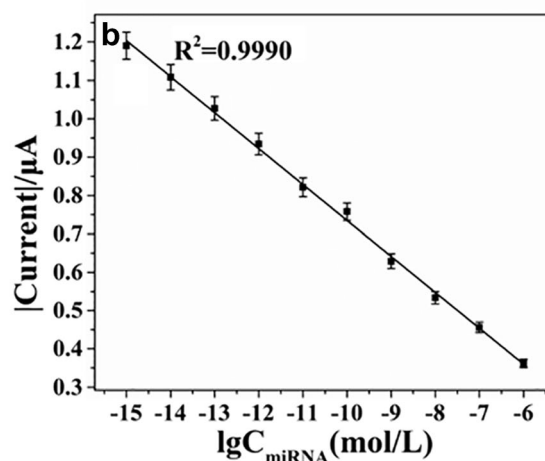


Fig. 4 a Photocurrent responses of the designed assay incubated with different concentrations of microRNA-21: (from a to j 1.0×10^{-6} , 1.0×10^{-7} , 1.0×10^{-8} , 1.0×10^{-9} , 1.0×10^{-10} , 1.0×10^{-11} , 1.0×10^{-12} , 1.0×10^{-13} , 1.0×10^{-14} , 1.0×10^{-15} M). **b** Calibration plot for microRNA-21



on the PEC assay. All photocurrent responses were measured in 0.10 M PB (pH = 7.4) at a bias voltage of 0 V with 415 nm light excitation. Error bars show the standard deviations of three replicate experiments

Table 1 Performance comparison of microRNA-21 detection by different detection methods

Analytical methods	Linear range	Detection limit	Ref
ECL	10 aM - 1.0 pM	3.3 aM	[34]
ECL	10 fM - 1.0 pM	10 fM	[35]
Fluorescence	1.0 fM - 10 nM	1.0 fM	[36]
Electrochemical	1.0 fM - 10 pM	1.0 fM	[37]
ECL	1.0 fM - 1.0 pM	1.0 fM	[38]
Fluorescence	10 fM - 1.0 nM	10 fM (37 °C)	[39]
	1.0 aM - 10 fM	1 aM (4 °C)	
Electrochemical	10 aM - 10 fM	5 aM	[40]
PEC	0.2 nM - 20 fM	5.6 fM	[41]
PEC	10 pM - 1.0 fM	0.5 fM	[42]
PEC	1.0 μM - 1.0 fM	14.7 fM	This work

determination (Table 1), the designed PEC aptamer method shows a preferable detection performance. In contrast, an additional control experiment was performed without Exo III assisted amplification. DNA₃ was employed as the cleaved

product of the duplex strand by Exo III. The PEC signal was gained without Exo III by direct titration of DNA₃ on the DNA₂ modified electrode surface. The detection limit was approximately in the nM range (Fig. S3), which was nearly 6–7 orders of magnitude poorer than that of Exo III catalyzed amplification.

Specificity of the PEC assay

To investigate the selectivity of the designed PEC assay, other ssDNA or RNA including miR-155, thrombin aptamer (TBA) and platelet-derived growth factor aptamer (Apt-PDGF) were tested as an interfering substance. From the results shown in Fig. 5a, the relative photocurrent change for 1.0 μM microRNA -21 is obvious. However, as for TBA, miR-155 and Apt-PDGF in the same concentration, the relative photocurrents response was similar to that of the blank, indicating the excellent selectivity of the designed PEC method. When 1.0 μM microRNA -21 was mixed with equal amount of TBA, miR-155 and Apt-PDGF, the relative photocurrent

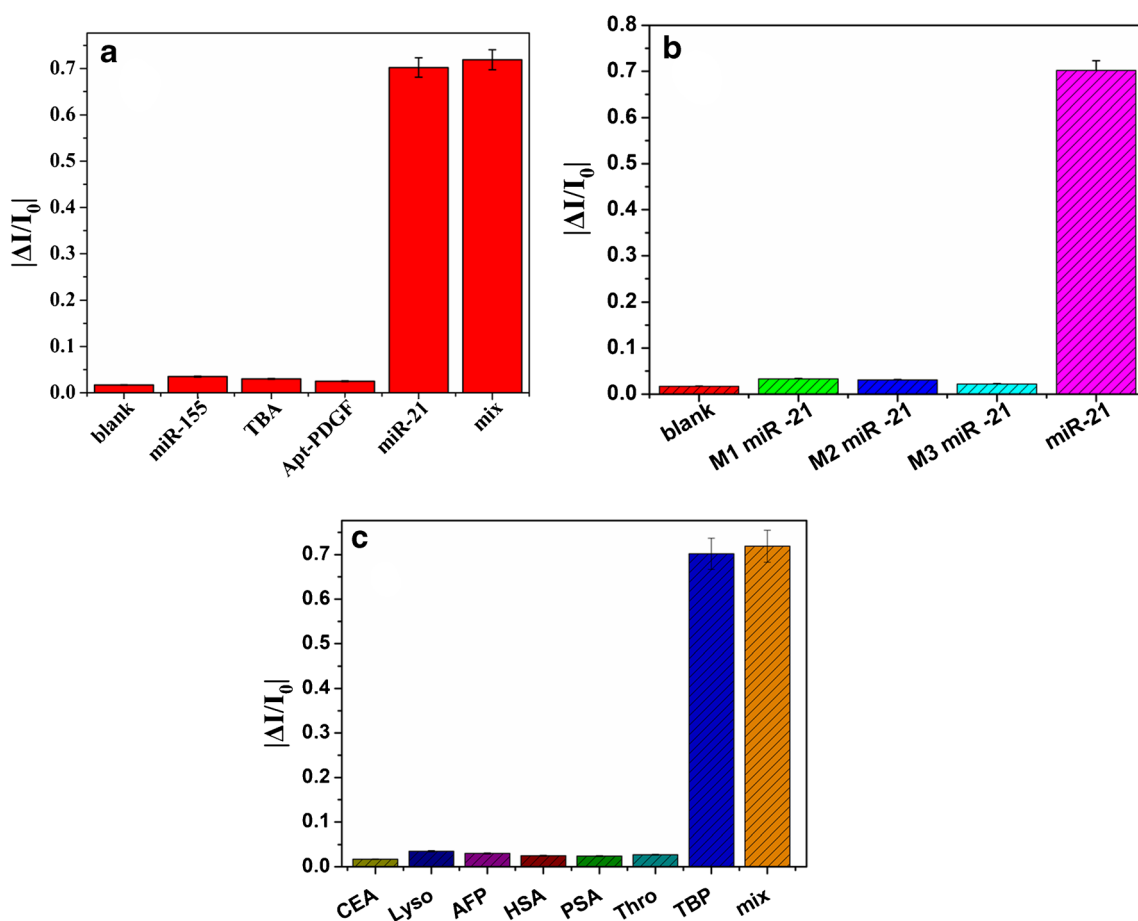


Fig. 5 **a** Selectivity experiment for microRNA-21 detection against microRNA-155, TBA, Apt- PDGF, and mixture. **b** The specificity of our assay in discriminating other RNA strands with one, two or three nucleotide mutations (M1 miR-21, M2 miR-21 and M3 miR-21). **c** Effects of other proteins on the detection of 1000 ng mL⁻¹ TBP in

0.10 M PB (pH = 7.4) at a bias voltage of 0 V with 415 nm light excitation. The bars represent 2000 ng mL⁻¹ of CEA, Lyso, AFP, HAS, PSA, Thro and a mixture of the seven kinds of proteins, respectively. The concentrations of other samples are 1.0 μM. The error bars represent the standard deviations for three replicates

Table 2 Recovery of microRNA -21 in normal human serum

Samples	Found microRNA-21 (pM)	Added microRNA-21 (pM)	Detected microRNA-21 (pM)	Recovery (%)	RSD (%) (n = 3)
1	0.011	0.01	0.0204	97.1	2.8
2		0.1	0.1128	101.6	4.3
3		1.0	1.0013	99.0	3.6

change was almost the same as that of pure microRNA -21. The results demonstrate that other DNA and RNA do not interfere with the detection of target object. (I_0 referred to the photocurrent of assembled electrode ITO/CuInS₂/CHIT/DNA₂, I referred to the final photocurrent response after addition different concentration of target microRNA -21.)

To investigate whether the mutation can be distinguished from the target microRNA, three different microRNAs were employed, including single-base mismatched microRNA-21 (M1 microRNA-21), two-base mismatched microRNA-21 (M2 microRNA-21) and three-base mismatched microRNA-21 (M3 microRNA-21). As displayed in Fig. 5b, the relative photocurrent change is occurred by the perfect target. The PEC signal outputs of M1 microRNA-21, M2 microRNA-21 and M3 microRNA-21 are similar to the blank signal as well. These results verify that one or few nucleotides mutation is easily distinguished by our designed strategy.

In addition, the interference study was evaluated with other proteins. The modified electrode was incubated in sample solutions containing different interfering proteins such as carcinoembryonic (CEA), lysozyme (Lyso), α -fetoprotein (AFP), human serum albumin (HSA), prostate specific antigen (PSA) and thrombin (Thro). The mixed solution of these proteins was also detected under the same conditions. Figure 5c shows that none of other proteins exhibit an obvious signal change, expect for the TBP and mixed sample. This phenomenon is due to the high affinity of the TBP toward the TATA box.

Practical application in the human serum

The practical application of the designed method in the human serum was tested by standard addition method. Healthy human serum samples spiked with different concentrations of target microRNA -21 were tested. The gained recoveries are found to vary from 97.1% to 101.6% (Table 2), the RSDs are less than 4.3%, demonstrating the excellent potential of the designed strategy in complex samples.

Conclusion

A novel PEC method was designed for the detection of microRNA. Several advantages of our method have been demonstrated: 1) A photocathode assay using CuInS₂ as a

photoactive species is synthesized, which can perform well in complex sample without the interference of reductive ingredient; 2) Exo III assistant amplification led to the recycling of the target microRNA and enhanced the detection sensitivity; 3) In this method, DNA-protein bioconjugates were used to enhance steric hindrance. One major limitation of this strategy is that the DNA probes immobilized on the electrode exhibited weak PEC signal reduction, which reducing the range of PEC signal response to TBP. The photocurrent variation produced by this recognition process can be realized easily. What's more, after carefully designing the recognition sequence, the designed PEC platform based on DNA amplification can also be used to detect other targets including DNA, protein and small molecule.

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Compliance with ethical standard

Conflict of interest The authors declare that they have no conflict of interest.

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