

Construction of an ultrasensitive electrochemical sensing platform for microRNA-21 based on interface impedance spectroscopy

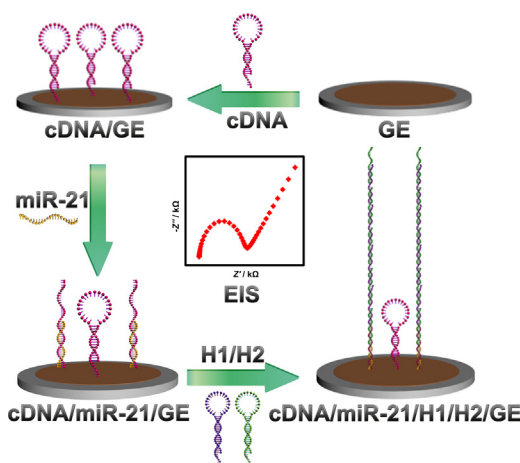
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HIGHLIGHTS

- An electrochemical sensing platform is constructed for detecting miR-21 based on interface impedance spectroscopy.
- The biosensor shows low limit of detection and wide dynamic correlation of miR-21.
- The proposed method is enzyme-free and PCR-free strategy.

GRAPHICAL ABSTRACT



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ABSTRACT

A hybridization chain reaction (HCR) amplification-based electrochemical impedimetric biosensor is fabricated for the quick, sensitive, and specific detection of miRNA-21 (miR-21) via monitoring of electrode interfacial property changes in real-time. Two sequences of H1 and H2 are adopted to trigger HCR amplification. A large amount of linear DNA concatemer are formed which could change the interfacial properties of the electrode. Interfacial charge transfer resistance difference (R_{ct}) is probed via electrochemical impedance spectroscopy (EIS) and Randles equivalent circuit. After amplifying via HCR, oligonucleotides with negatively charged repelling $[\text{Fe}(\text{CN})_6]^{3-/4-}$ ions can form a spatial blockage. HCR amplification strategy markedly enhanced the electrochemical signal with a limit of detection (LOD) down to 4.63 fM ($S/N = 3$). This strategy exhibited excellent selectivity for three different miRNAs: miR-199a, miR-141, and miR-155. Moreover, results show that the proposed method can be applied to miR-21 detection in the total RNA extracted from five cells. This work presents an enzyme-free and label-free EIS nucleic acid sensor for sensitively and selectively detecting miR-21, offering a promising approach in early diseases diagnosis.

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1. Introduction

Electrochemical biosensors have been widely utilized in energy resources, the life sciences, and other fields because of their simplicity of operation, quick responsiveness, inherent sensitivity, and good selectivity [1–4]. Currently, electrochemical biosensors have been efficiently applied in the detection of various miRNAs [5,6]. Among different electrochemical methods, electrochemical impedance spectroscopy (EIS) is an efficient method for investigating electrode surface change through the sinusoidal electric potential application with a variable frequency [7,8]. Utilizing equivalent circuits allows the experimental spectra to be fitted with the theoretical curve that corresponds to the selected circuit model, thereby obtaining the resistance values [9]. However, EIS-based label-free biosensors possess lower sensitivity than those with labeling methods. Therefore, nanomaterials and various amplification strategies have been utilized to enhance impedimetric sensing sensitivity [7]. Although most developed methodologies that relying various labels offer enhanced sensor sensitivity, they also increase the assay procedure complexity, time, and cost [10]. Moreover, the EIS can be seen as a label-free electrochemical method which can achieve transduction in the absence of labeled species. EIS have successfully been utilized to detect various biomolecules, such as toxins, bacteria, different kinds of DNA, and miRNAs during the last few years due to steric hindrance and electrostatic repulsion. For example, Miranda-Castro et al. constructed an impedimetric sensor to detect sensitively the uropathogenic *E. coli* cells by utilizing artificial recognition sites [11]. Azzouzi et al. successfully fabricated an integrated dual-functional recognition/amplification biosensor for the sensitive detection of microRNA-21 (miR-21) via EIS [12]. Kannan et al. constructed a cobalt oxide porous nanocubes-based impedimetric biosensor for the sensitive detection of hepatitis B virus DNA [13].

MicroRNA (miRNAs) is an endogenous, noncoding, single-stranded RNAs with approximately 18–24 nucleotides [14]. They are closely related to almost every cellular process, including post-transcriptional gene regulation, cell proliferation, differentiation, and apoptosis [15–18]. Sufficient evidence suggests that miRNAs can emerge as significant diagnostic markers due to miRNAs over-expression in the cellular process [19–22]. Among them, miR-21 is over-expressed in multiple malignant tumors, including lung cancer, breast cancer, uterine cancer, gastric cancer, pancreatic cancer, liver cancer, cholangiocarcinoma, colon cancer, and prostate cancer, as well as participating in tumor occurrence and development through complex regulation [23–25]. Furthermore, the miR-21 content derived from five different cancer cells including breast (MCF-7), cervical (HeLa), non-small cell lung (A549), mouse leukemia cells of monocyte-macrophage (RAW 264.7), and human umbilical vein endothelial (HUVEC) was investigated to analyze the feasibility of the proposed impedimetric biosensor. Among these, HUVEC was utilized as an endogenous reference gene for normalization purposes. Therefore, accurate and sensitive miR-21 detection can be a powerful tool for clinical diagnosis and therapy.

Recently, many target amplification assays were explored to achieve highly sensitive miR-21 electrochemical determinations. The most common methods for achieving the target amplification are as follows: loop-mediated isothermal amplification (LAMP) [26], catalytic hairpin assembly (CHA) [27,28], strand-displacement amplification (SDA) [29], isothermal exponential amplification reaction (EXPAR) [30], and hybridization chain reaction (HCR) [31–34]. In a typical HCR, the initiator triggers a self-assembly reaction of two hairpins with partial complementarities to form alternate nucleic acids concatemers. HCR amplification strategies have proven to be a particularly effective method for

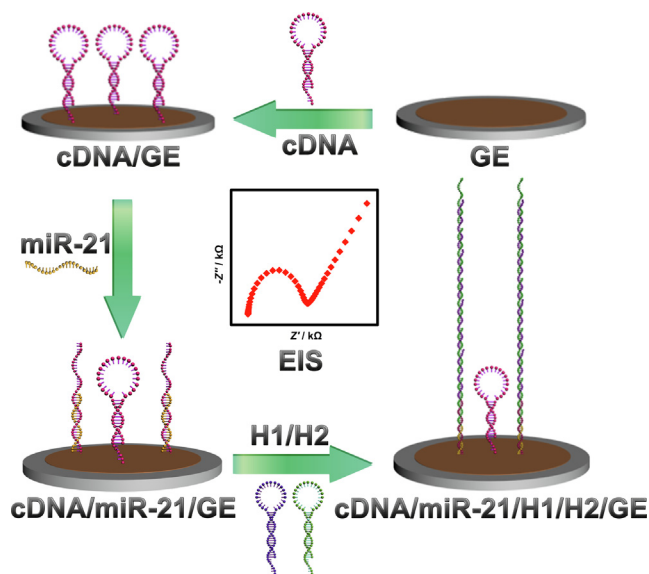
fabricating the electrochemical nucleic acid biosensor because of its isothermal, enzyme-free, high amplification efficiency, and mild reaction conditions. For example, Zhou et al. constructed a novel electrochemical assay for sensitively detecting of miRNAs via non-linear HCR [35]. From Zhang's report, HCR amplification strategy and MoS₂-loaded G-quadruplex molecular beacon probes were utilized to improve sensitive detection of miRNA-199a/b-3p and miR-21 [36]. Cheng et al. successfully constructed mesoporous silica container biosensors for detecting miR-21 via CHA/HCR [5]. Also, Li et al. have constructed a biosensor for sensitively detecting miRNA-141 via a one-step introduction of a target-triggered branched HCR circuit [37].

In this paper, an electrochemical nucleic acid biosensing approach for miR-21 detection is proposed based on HCR strategy signal amplification coupled with EIS (Scheme 1). The thiol-functionalized capture hairpins (cDNA) were immobilized on the gold electrode (GE) surface. When the target miR-21 was presented in the system, the cDNA hairpin structures were opened. Thereafter, the released cDNA sequences are utilized to initiate the HCR reaction between two hairpin probes (H1 and H2), and many linear DNA concatemers can eventually be formed. The steric hindrance and electrostatic repulsion between the inherent negatives were induced via linear DNA concatemers, leading to a great increase of the interfacial electron transfer resistance (R_{ct}). The impedimetric biosensor also achieved miR-21 detection in human serum. The impedimetric biosensor combining the HCR signal amplification and EIS method obtained a wide linear range, high sensitivity, and excellent specificity toward miR-21.

2. Experimental

2.1. Reagent and apparatus

Gelred and loading buffer were received from Sangon, Inc. (Shanghai, China). Ethylenediaminetetraacetic acid disodium salt (EDTA), trichloromethane (CHCl₃), HCl, orthoboric acid 6-Mercapto-1-hexanol (MCH), and isopropanol were provided by MACKLIN Reagent Co. Ltd. *N*-hydroxysuccinimide (NHS, 98%), *N*-(3-dimethylaminopropyl)-*N*-ethylcarbodiimidehydrochloride (EDC, 98%), ammonium persulphate (APS), and tris were provided



Scheme 1. A principle of impedimetric biosensing platform for detecting miR-21 based on HCR.

by Aladdin, Inc. (Shanghai, China). Diethyl pyrocarbonate (DEPC) and acrylamide were provided by Biosharp, Inc. (Hefei, China). *N,N,N',N'*-Tetramethylethylenediamine (TEMED) was provided by Sigma-Aldrich, Inc. (Beijing, China). All HPLC purified oligonucleotides were provided by Sangon, Inc. (Shanghai, China) (Table 1). All electrochemical measurements were carried out with an AUTO-LAB Electrochemical Workstation (Metrohm Instruments, Switzerland).

2.2. Extraction of total RNA in cells

The proposed method was investigated to detect the miR-21 content in total RNA, which were extracted from A549, HeLa, MCF-7, RAW 264.7, and HUVEC. In a typical trizol method, trizol can maintain RNA integrity during sample splitting or homogenization. The solution is divided into a water phase and an organic phase after chloroform is added. RNA is present in the aqueous phase, then, after taking out the aqueous phase, isopropanol can be utilized to precipitate and recover RNA. The trizol method has a high extraction efficiency. Also, the extracted total RNA can effectively avoid pollution by DNA and protein. Therefore, the trizol method has little influence on extracting RNA from cells. About 1×10^6 cells were rinsed by 0.1 M of PBS (pH 7.4) then dispersed in a non-enzymatic centrifuge tube containing 1 mL of trizol. Next, 200 μ L of CHCl_3 was added into the non-enzymatic centrifuge tube to separate RNA. Then, the supernatant was carefully transferred into a fresh non-enzymatic centrifuge tube and washed with isopropanol and ethanol. Finally, total RNA was dispersed in 15 μ L of DEPC then stored at -80°C before utilization.

2.3. Preparation of the target-triggered amplification electrochemical biosensor

The GE was initially polished carefully with 0.3 and 0.05 μm of alumina powder on a polishing cloth of chamois leather, then it was sonicated in deionized water and ethanol for 30 s. Before the experiment, the DNA hairpin probes including cDNA, H1, and H2 were denatured at 95°C for 5 min then gradually gradient cooled to room temperature to form a stem-loop structure before utilization. To reduce disulfide bonds, thiol-modified cDNA was dispersed in a TE buffer containing 10 mM TCEP for 1 h at room temperature. Then, 8 μ L of cDNA (2.5 μM) was incubated on the well-cleaned GE surface at 4°C for 14 h through the Au-S bond. Next, the prepared electrode was thoroughly washed with deionized water to remove physically adsorbed cDNA. Then, 10 μ L of 1.0 mM MCH was dropped onto the modified electrode for 40 min to prevent non-specific adsorption. Meanwhile, steric hindrance was minimized to enhance the modified electrode charge transfer. After rinsing with deionized water, 8 μ L of target miR-21 was dropped onto the prepared electrode and incubated for 60 min at 37°C . The mixed solution was prepared by mixing 8 μ L of 2.5 μM H1 and

8 μ L of 2.5 μM H2. Subsequently, the mixed solution was introduced to the prepared electrode for 80 min at 37°C to trigger the HCR process, thereby producing numerous linear DNA concatamers. In the end, the self-assembly electrode was rinsed with deionized water then dried in nitrogen gas and stored in the air before utilization.

2.4. Native polyacrylamide gel electrophoresis (PAGE)

Fifteen percent native polyacrylamide gel (5 mL of 30% acrylamide-bisacrylamide, 2.4 mL of deionized water, 2.5 mL of $5 \times \text{TBE}$, 100 μ L of 10% APS, and 4 mL of TEMED) was utilized to characterize the the HCR process. The samples were dropped into each lane, then electrophoresis was run in $1 \times \text{TBE}$ buffer operated at 100 V. Next, the gel was dyed with gelred for 30 min. Finally, the gel was scanned via a Bio-Red Imaging System.

3. Results and discussion

3.1. Analysis of the HCR via PAGE

HCR assembly process feasibility has been investigated by PAGE (Fig. S1). Lanes 1–4 were loaded with 5 μM of cDNA, 5 μM of H1, 5 μM of H2, and 20 μM of miR-21. Only one well-defined band in lanes 1–4 is exhibited. Lane 4 shows a fast-migrated rate because of the low molecular weight of miR-21. The one bright band in lane 5 (without miR-21) can be observed, confirming no reaction occurred for it. In the presence of miR-21, lane 6 shows many bright bands, demonstrating the successful assembly of many H1 and H2 to form chain-like DNA polymers.

3.2. Electrochemical characterization of fabricated biosensor

EIS is effective for exploring GE stepwise assembly processes. An ideal Nyquist plot consists of a linear part and a semicircle portion, corresponding to diffusion from the solution to the interface and the electron transfer-limited process. Higher-frequencies diameter refers to the electron transfer resistance (R_{ct}). Electron transfer kinetics processes were analyzed via the Randles equivalent circuit. Fig. 1 shows the gradual modification processes of the biosensor; curve a shows the smallest semicircle of the bare GE, suggesting the fast charge transfer on the GE surface. The

Table 1
The sequences of all oligonucleotides.

Name	Sequence (5'–3')
cDNA	SH-(CH ₂) ₆ -TCAACATCAGTCTGATAAGCTA GCATCCTAGCTTATCAGACTGAAGCTA
H1	TATCAGACTGAAGCTATTCAAAGT TAGCTTCAGTCTGATAA GCTAGGA
H2	ACTTTGAATAGCTTCAGTCTGATATCTAGCT TATCAGACTGAAGCTATT
miR-21	UAGCUUAUCAGACUGAUGUUGA
miR-155	UUAAUGCUAAUCUGAUAGGGGU
miR-141	UAACACUGUCUGGUAAGAUGG
miR-199a	ACAGUAGUCUGCACAUGGUUA

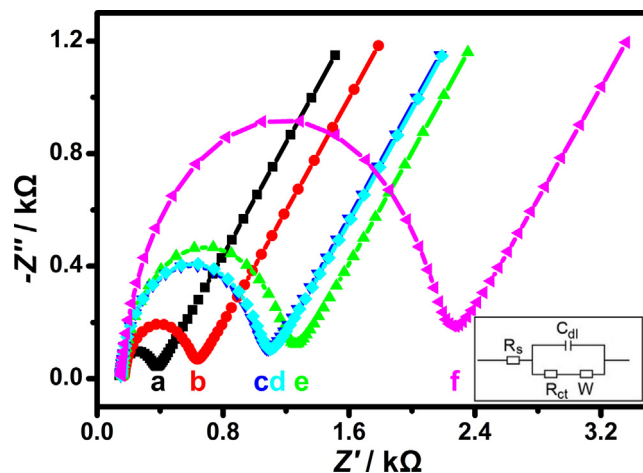


Fig. 1. Nyquist plots of stepwise assembly process: (a) bare GE, (b) cDNA/GE, (c) cDNA/MCH/GE, (d) cDNA/MCH/H1/H2/GE, (e) cDNA/MCH/miR-21/GE, and (f) cDNA/MCH/H1/H2/miR-21/GE. The detection was analyzed in 5.0 mM of $[\text{Fe}(\text{CN})_6]^{3-/4-}$ containing 0.1 M of KCl (0.1 Hz–100 kHz); Open current potential: 0.24 V, amplitude: 0.01, $n = 10$.

smallest R_{ct} (257 Ω) is also obtained through the Randles equivalent circuit. After introducing cDNA onto the bare GE surface, the R_{ct} (433 Ω) consequently increases, which is attributed to the steric hindrance formed by the self-assembly of cDNA (curve b). Increased R_{ct} also suggests the successful assembly between cDNA and bare GE. After MCH was modified on the electrode surface, the R_{ct} (882 Ω) further increased (curve c), which is due to the MCH film blocking the unreaction sites. Subsequently, the R_{ct} (1040 Ω) increases further when the target miR-21 was incubated on the modified GE (curve e). Curve d ($R = 895 \Omega$) exhibits a semicircle diameter similar to that of curve c, demonstrating that the HCR process cannot be triggered without miR-21. The cDNA/MCH/H1/H2/miR-21/GE (curve f) exhibits the largest R_{ct} (2230 Ω). The remarkable semicircle increase of the cDNA/MCH/H1/H2/miR-21/GE suggests a successful and acceptable GE hybridization strategy process. Fig. S2 shows the data of the red curve and the black curve are consistent, demonstrating that the equivalent circuit is justified for explaining the proposed sensors.

To achieve the best analytical performance of this proposed strategy, investigating some potentially effects is necessary. Different concentrations ranging from 0.5 to 10 μM of cDNA immobilized on the GE were optimized (Fig. 2A). The R_{ct} value originally increased with an increase of the cDNA concentration, then it become stable from 5 μM to 10 μM . Therefore, 5 μM was selected as the best cDNA concentration. Fig. 2B shows a similar trend, illustrating that the best H1 and H2 incubation concentration was 2.5 μM . Simultaneously, the effect of the hybridization time of cDNA was investigated, ranging from 10 h to 18 h (Fig. 2C). Results show that the maximum R_{ct} value was obtained at 14 h. Also, the

maximum R_{ct} value was generated when the miR-21 hybridization time reached 60 min (Fig. 2D), demonstrating that this is the optimal miR-21 hybridization time in this system. Moreover, the effect of the H1 and H2 incubation time was explored (Fig. 2E). Consequently, 90 min was selected as the incubation time for whole experiments.

EIS spectra were utilized to assess the sensitivity, linear range, and limit of detection (LOD) of the proposed biosensor. Fig. 3A shows that the R_{ct} value gradually increased as the miR-21 concentration increased. Fig. 3B shows that the calibration plots EIS response exhibited a good linear range from 10 fM to 50 pM; the linear equation is $\Delta R_{ct} = -0.01907 + 0.35955 \log C_{\text{miR-21}} \text{ (pM)}$ ($R = 0.9988$). The LOD is evaluated as 4.63 fM ($S/N = 3$), which is competitive with the values in previous reports (Table 2). Moreover, the EIS can be a label-free electrochemical method, achieving transduction in the absence of labeled species. This label-free electrochemical method also shortens biosensor preparation time. In some typical labeled electrochemical biosensors, various nanoparticles with electrocatalytic properties, such as Au nanoparticles, metallic oxide, and metal sulfide, should be prepared first, thereby expanding the time detection. Therefore, the proposed electrochemical impedimetric biosensor provided a novel method for detecting other biomarkers.

To validate the established biosensor accuracy, an examination was conducted of the miR-21 in 10% human serum. Real samples were spiked with certain amounts of miR-21 to detect and calculate recoveries. Table 3 shows that miR-21 recoveries range from 83.61% to 118.61% with RSD ranging from 0.96% to 9.67%, suggesting the proposed sensor accuracy was desirable.

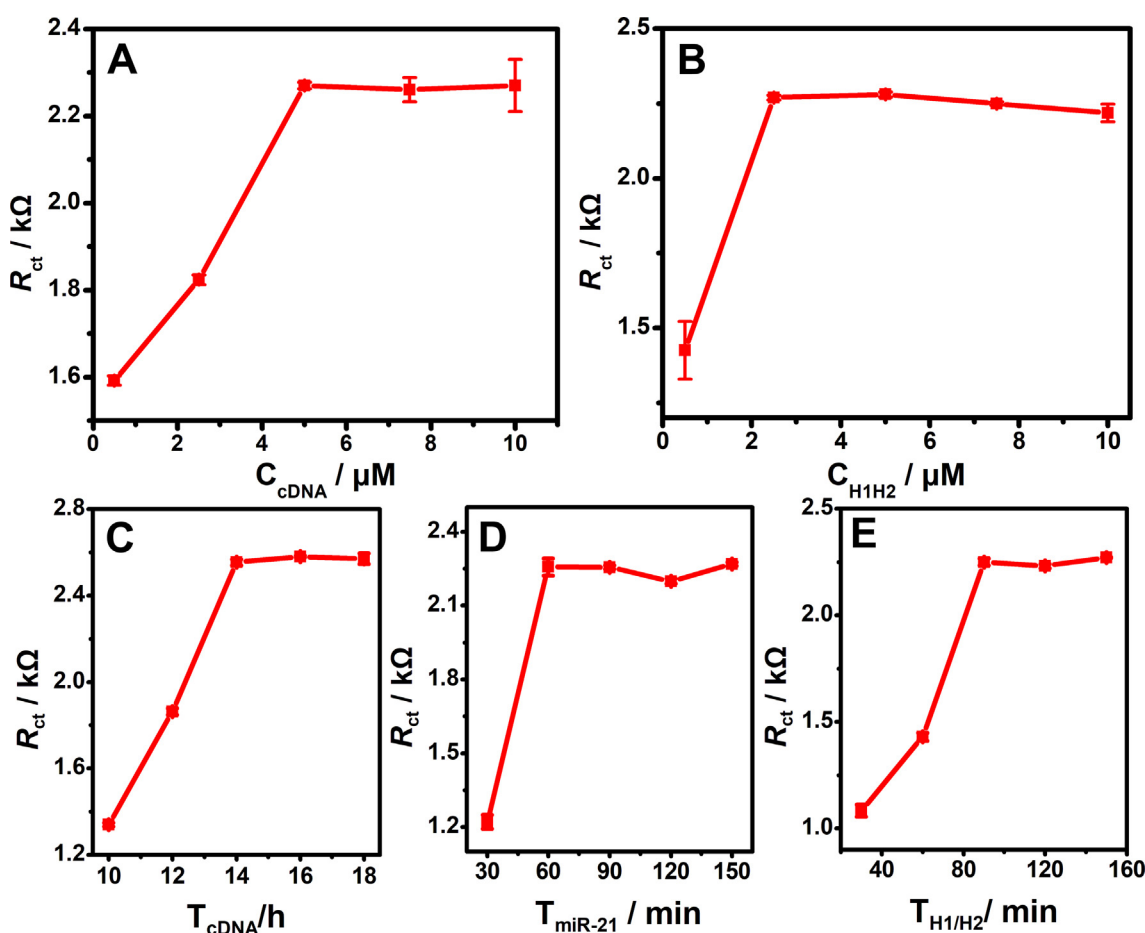


Fig. 2. Influence of the cDNA (A) and H1, H2 (B) concentrations. Effects of hybridization time of cDNA (C), miR-21 (D) and H1, H2 (E), $n = 10$.

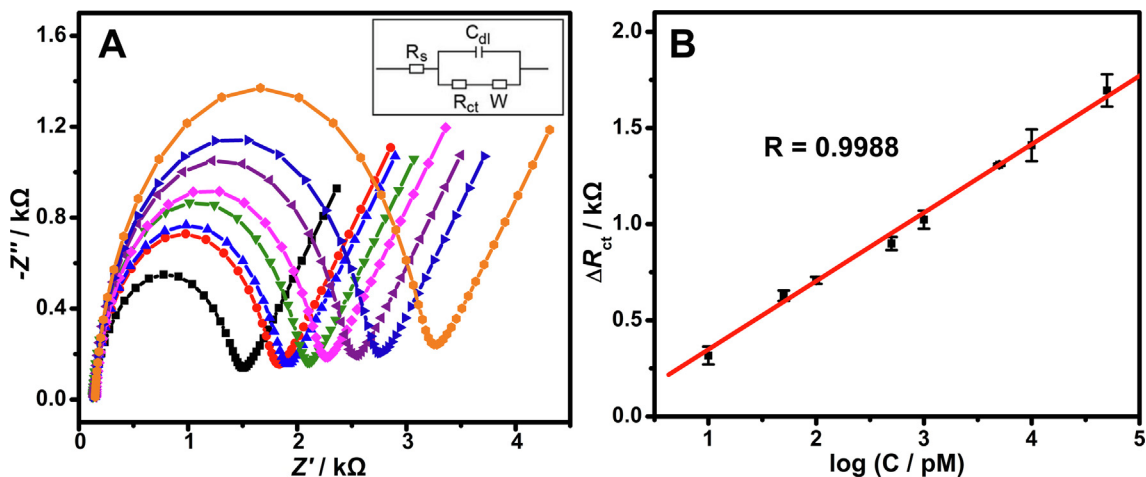


Fig. 3. EIS responses of the fabricated biosensor with different concentrations of miR-21 (10 fM to 50 pM) (A); The detection was analyzed in 5.0 mM of $[\text{Fe}(\text{CN})_6]^{3-/4-}$ containing 0.1 M of KCl (0.1 Hz–100 kHz); Open current potential: 0.24 V, amplitude: 0.01. Calibration plot of the ΔR_{ct} vs. the logarithm of miR-21 concentration (B), $n = 10$.

Table 2

Comparison of different methods for detection of miRNA.

Materials	Analytical technique	Linear range	Limit of detection	Reference
biotin-MB-AuNPs ^a	EIS	1–1000 pM	300 fM	[27]
CdTe nanocrystals and Au nanoclusters ^b	ECL ^c	100 fM–100 nM	21.7 fM	[38]
Electrode-free	Colorimetric	0–0.1 nM	32 fM	[39]
ss-DNA/MWCNT/MB/GCE ^d	DPV ^e	0.1–500 pM	84.3 fM	[40]
Au@NPFe ₂ O ₃ NC ^f	DPV	500 fM–5 nM	100 fM	[41]
Arched Probe Mediated Isothermal Exponential Amplification	DPV	20 fM–10 nM	5.36 fM	[42]
MBA-AuNPs/DA-AuNPs ^g	DPV	100 fM–10 pM	45 fM	[43]
Hairpin capture probe/target recycling/MB ^h	DPV	0.5 pM–2 nM	56 fM	[44]
MoS ₂ -Thionine-AuNPs ⁱ	SWV ^j	1 pM–10 nM	260 fM	[45]
GO-CA-PGEs ^k	EIS	355–2130 nM	41.2 nM	[46]
Dendrimer-SPE ^l	EIS	140–360 nM	74 nM	[47]
polypyrrole	EIS	5–80 pM	2 pM	[48]
cDNA/MCH/H1/H2/miR-21/GE	EIS	10 fM–50 pM	4.63 fM	This work

^a Molecular beacon probe was conjugated with gold nanoparticles.

^b CdTe nanocrystals and Au nanoclusters modified on glassy carbon electrode (GCE).

^c Electrochemiluminescence.

^d miRNA/single strands DNA (ss-DNA)/multi-walled carbon nanotube/molecular blue modified on GCE.

^e Differential pulse voltammetry.

^f Gold-loaded nanoporous iron oxide nanocubes.

^g 4-mercaptophenylboronic acid -capped gold nanoparticles /dopamine -capped gold nanoparticles.

^h Hairpin capture probe/target recycling/methyl blue.

ⁱ Thionine and gold nanoparticles co-functionalized MoS₂ nanosheet.

^j Square wave voltammetry.

^k Graphene oxide/covalent agents/pencil graphite electrodes.

^l Poly(amidoamine) dendrimer modified single-use screen printed carbon electrode.

Table 3

The proposed biosensor applied in 3 human serum samples.

Human serums	Added (fM)	Founded (fM)	Recovery (%)	RSD (%)
1	0	1.3	–	0.96
	100	119.9	118.61	1.60
	500	419.35	83.61	1.40
2	0	0.63	–	9.10
	100	108.64	108	3.47
	500	492.31	98.34	9.67
3	0	0.15	–	3.00
	100	112.47	112.33	1.16
	500	488.53	97.68	1.70

To investigate the selectivity of this proposed method, miR-199a, miR-155, and miR-144 sequences replaced the target miR-21 (Fig. 4A). A lower ΔR_{ct} value of miR-199a, miR-155, and miR-144 sequences was noticed when compared to the ΔR_{ct}

of the target miR-21. Moreover, the biosensor incubated with the mixed samples exhibited a similar ΔR_{ct} value when only miR-21 was present. These results show that the proposed electrochemical impedimetric biosensor is efficiently selective toward miR-21.

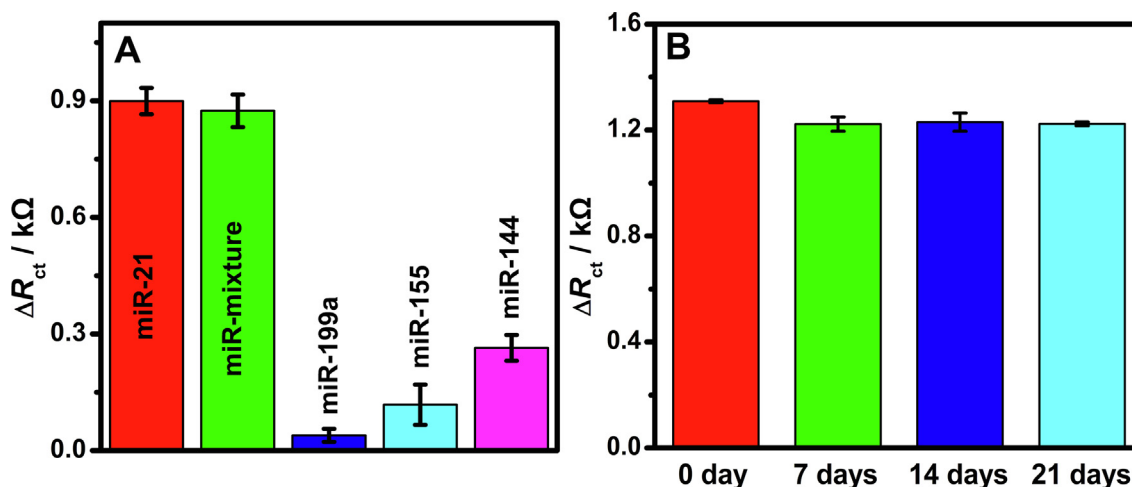


Fig. 4. EIS responses of the proposed biosensor for the detection of different miRNA sequences, including 5.0×10^{-7} μM of miR-21, 5.0×10^{-6} μM of miR-199a, 5.0×10^{-6} μM of miR-155, 5.0×10^{-6} μM of miR-144, and the mixture of the four kinds of miRNAs (A). The storage stability of the miR-21 biosensor at 4 °C (B), $n = 10$.

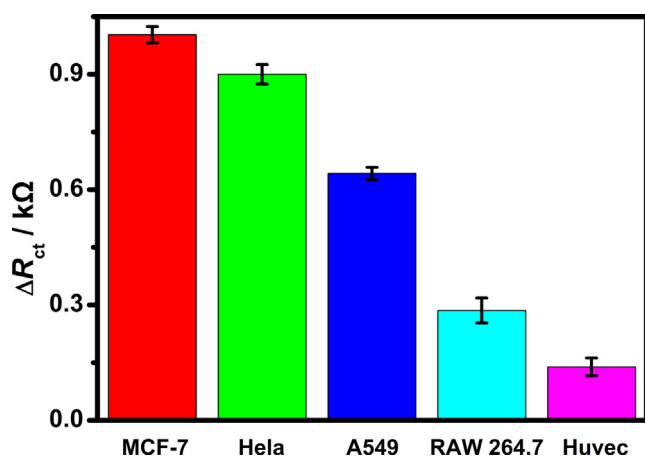


Fig. 5. EIS responses for detecting miR-21 produced from MCF-7, HeLa, A549, RAW 264.7, and HUVEC cells, $n = 10$.

To demonstrate the biosensor stability, EIS was employed to assess the self-assembly electrode change in the presence of 5.0×10^{-6} μM of miR-21 (Fig. 4B). The prepared sensing electrodes were kept at 4 °C for different times (0, 7, 14, and 21 days). As expected, biosensor ΔR_{ct} values stored for 7, 14, and 21 days decrease less than the initial response ΔR_{ct} value. This shows that the proposed impedimetric biosensor possesses satisfactory storage stability.

Further analysis was conducted on the feasibility of the impedimetric biosensor, which was utilized to determine the content of miR-21 derived from five different cancer cells (Fig. 5). A more significant EIS response was obtained from MCF-7, HeLa, and A549 cells than that obtained from the endogenous reference genes RAW 264.7 and HUVEC. This proves the over-expression of miR-21 in HeLa, MCF-7, and A549, which conforms to the reported work [46–50].

4. Conclusion

In conclusion, a facile electrochemical biosensor combining efficient HCR strategy and EIS method was first clarified for sensitive miR-21 determination. The interfacial charge transfer resistance (R_{ct}) difference was probed via EIS and Randles equivalent circuit.

HCR relies on DNA strand substitution to form alternate nucleic acid concatamers and avoid some fussy steps of protein enzymes. After amplifying via HCR, oligonucleotides with negatively charged repelling $[\text{Fe}(\text{CN})_6]^{3-/4-}$ ions can form spatial blockages. Many linear DNA concatamers can lead to a great increase in interfacial R_{ct} . R_{ct} positively correlated with miR-21 content enables direct and quantitative miR-21 detection. Then, the designed biochemical self-assembly strategy of this impedimetric biosensor is successfully applied to the analysis of the miR-21 expression level extracted from five different cells. This electrochemical biosensor design would help rapidly and simply detect miR-21, thereby developing a powerful clinical analysis platform.

CRediT authorship contribution statement

Tianjiao Meng: Methodology, Software, Writing - original draft. **Dan Zhao:** Writing - review & editing. **Huimin Ye:** Data curation, Writing - original draft. **Yue Feng:** Methodology, Data curation. **Huan Wang:** Writing - review & editing, Funding acquisition. **Yufan Zhang:** Investigation, Methodology, Data curation, Software, Supervision, Funding acquisition.

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.jcis.2020.05.118>.

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