



Label-free nanostructured biosensor for *Schistosoma mansoni* detection in complex biological fluids

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ABSTRACT

Schistosomiasis is a neglected tropical disease with a worldwide prevalence. Neuroschistosomiasis is the most severe presentation of the disease and affects the central nervous system. In this work, *Schistosoma mansoni* detection was based on self-assembled layers of 3-mercaptopropyltrimethoxysilane (MPTS) and electro-synthesized gold nanoparticles (AuNPs). The DNA probe was chemisorbed onto AuNPs. The biosensor was characterized by electrochemical impedance spectroscopy (EIS) and cyclic voltammetry (CV). The impedimetric response of the MPTS-AuNPs-DNA_{probe} system indicates an effective modification of the electrode surface. Topographical atomic force microscopy images were used to characterize the self-assembled layers on the gold electrode surface. The proposed biosystem was able to recognize the *S. mansoni* genome sequence at different concentrations in samples of urine, cerebrospinal fluid, and serum. Several concentration ranges were evaluated: urine (27–50 pg μL^{-1}), cerebrospinal fluid (25–60 pg μL^{-1}), and serum (27–42 pg μL^{-1}). The limit detection (LOD) of the biosensor was 0.6 pg μL^{-1} . The developed label-free genosensor was able to detect small concentrations of *S. mansoni* DNA in complex biological fluids.

1. Introduction

Schistosomiasis has a critical role for public health with socio-economic impacts on impoverished communities [1]. Of note, chronic *S. mansoni* infection leads to significant residual morbidity. The disease remains a devastating neglected tropical disease that affects approximately 260 million people worldwide [2,3]. Also, about 800 million people are currently at risk of developing schistosomiasis worldwide [4,5].

The symptomatology of the acute phase is related to fever, diarrhea, headache, asthenia, and anorexia, or even asymptomatic [6]. The continuous infections can result in granulomatous reactions and fibrosis in the affected organs. Also, these manifestations can occur associated with lesions of the central nervous system, polyposis with bloody diarrhea, splenomegaly, portal hypertension, and pulmonary hypertension [7]. Neuroschistosomiasis (NS) can clinically manifest as meningomyelorradiculitis, myelitis, and radiculitis. In addition, cerebral, pseudotumoral, or spinal forms are also observed. Medullary

schistosomiasis is the most severe clinical manifestation of the *S. mansoni* infection [8].

The intestinal schistosomiasis diagnosis is usually performed through the Kato-Katz fecal smear technique (a standard method recommended by the World Health Organization for qualitative and quantitative determination). However, the Kato-Katz technique is limited due to its low sensitivity, particularly in mild infections [9,10]. Immunological assays are used for performing the routine laboratory tests, especially in the chronic-phase chagasic infection [11]. The immunological assays are intradermal reaction, complement fixation reaction, indirect immunofluorescence, enzyme-linked immunosorbent assay (ELISA), and capture ELISA [12]. In addition, the immunological assays are dependent on the search for *S. mansoni* antibodies [13,14].

However, antibodies-based tests present limitations due to the cross-reaction with other antigens (mainly of other helminths), difficulty of differentiating from the active infection of the previous contact with the parasite, and low sensitivity for asymptomatic cases [15]. Cerebrospinal fluid (CSF) is extensively used for diagnosis of patients with

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clinically suspected NS [16].

In this context, electrochemical genosensors have become an alternative for the development of new diagnostic methods [17,18]. Nanomaterials have been used for the development of biodevices due to their huge surface-to-volume ratio, controlled size, and differentiated physicochemical properties. In addition, nanomaterials are extensively used to immobilize the recognition elements (e.g., DNA or proteins) [19].

AuNPs are the most commonly used nanomaterial in biosensor development through biomolecules conjugation [19]. AuNPs have been used associated with different nanomaterials as graphene oxide (GO) and carbon nanotubes (CNT) to obtain hybrids nanoplateforms for DNA detection [19,20]. Hybrids nanoplateforms improve the sensor performance due to the large surface area and good electrical conductivity attributed to a synergistic effect of these nanomaterials [21–24]. Electrochemical techniques can monitor variations in the electrical properties (resistance or capacitance) of the electrode surface [25]. Potentiodynamic methods (e.g., potential cycling) are suitable for obtaining AuNPs on the electrode surface [26,27]. Thus, impedimetric nanosensors are potentially promising due to their potential for rapid DNA detection with low cost and sensitivity [25,28].

In this connection, the present study evaluated the assembly process of the 3-mercaptopropyltrimethoxysilane (MPTS) and electrochemically synthesized AuNPs (Fig. 1) for *S. mansoni* genome detection. The organosilane (MPTS) was used to obtain compelling sol-gel derivative films to immobilize AuNPs since it interacts with the working electrode through Au–S bonds [29]. AuNPs were electrochemically synthesized, and the applied potential cycling controls the nucleation/growth process of the AuNPs [30]. Also, AuNPs offer a unique approach for immobilization of thiol-modified DNA. Therefore, the association of sol-gel derivative films with AuNPs allows the development of a genosensor for *S. mansoni* detection in CSF and other biological fluids.

2. Materials and methods

2.1. Materials

DNA samples of *S. mansoni* were assigned by the Department of Parasitology of the Aggeu Magalhães Institute (Recife, Brazil). RT-qPCR was useful for the previous characterization of all samples. 3-Mercaptopropyltrimethoxysilane (MPTS), Gold chloride (III) hydrate

($\text{HAuCl}_4 \cdot x\text{H}_2\text{O}$), and bovine serum albumin (BSA) were purchased from Sigma-Aldrich (St. Louis, MO, USA). Ferrocyanide and potassium ferrocyanide were obtained from VETEC (Brazil). DNA solutions were prepared in phosphate-buffered saline (PBS, pH 7.4). The other chemicals used in this work were reagent grade and used as received without further purification. Ultrapure water was obtained from a Millipore Milli-Q system.

2.2. Sample preparation

Controls and patient's samples were collected ($v = 1 \text{ mL}$) and their analyses consisted of general and specific cell counts, glucose, CSF, protein electrophoresis, and immunological determinations (ELISA) for schistosomiasis, herpes, syphilis, cysticercosis, toxoplasmosis, cytomegalovirus, Epstein-Barr virus, human immunodeficiency virus, varicella zoster, Cryptococcus, and human cell lymphotropic virus. The samples were also analyzed by nPCR [8]. Electrophoresis using genomic DNA from *S. mansoni* at different concentrations is shown in Fig. S1.

The urine samples ($v = 5 \text{ mL}$) were transferred to a 15-mL Falcon tube. DNA was extracted according to Silva using protocol 2 [31]. Schistosome DNA was purified using the illustrated tissue & cells genomic Prep Mini Spin Kit (GE Healthcare, UK) according to the manufacturer's instructions. DNA samples from patients were extracted and purified using the illustrated blood genomic Prep Mini Spin Kit (GE Healthcare, UK) according to the manufacturer's instructions. Thus, it was quantified in a spectrophotometer and stored at -20°C . The thiolated specific primer for *S. mansoni* was Schfo17 (5'-GTGCTGGTGGTTGACGAGTTC-3') [32].

2.3. Modification of the electrode surface

Initially, 50 mM MPTS solution was prepared by mixing MPTS with deionized water (0.5:100, v/v) and 600 μL of 0.1 mol L^{-1} HCl [33]. Then, the mixture was sonicated for 9 min followed by stirring for 45 min until obtaining a transparent homogeneous solution. The electrode surface was gently polished with sandpaper and deionized water. Subsequently, the bare gold electrode (BGE) was immersed in a solution of sodium hypochlorite for 10 min, washed with deionized water, and dried at room temperature. After, the electrode surface was modified with 4 μL of the homogeneous solution of MPTS by drop coating and allowed to dry at 25°C . Subsequently, 1 mM de $\text{HAuCl}_4 \cdot x\text{H}_2\text{O}$ in

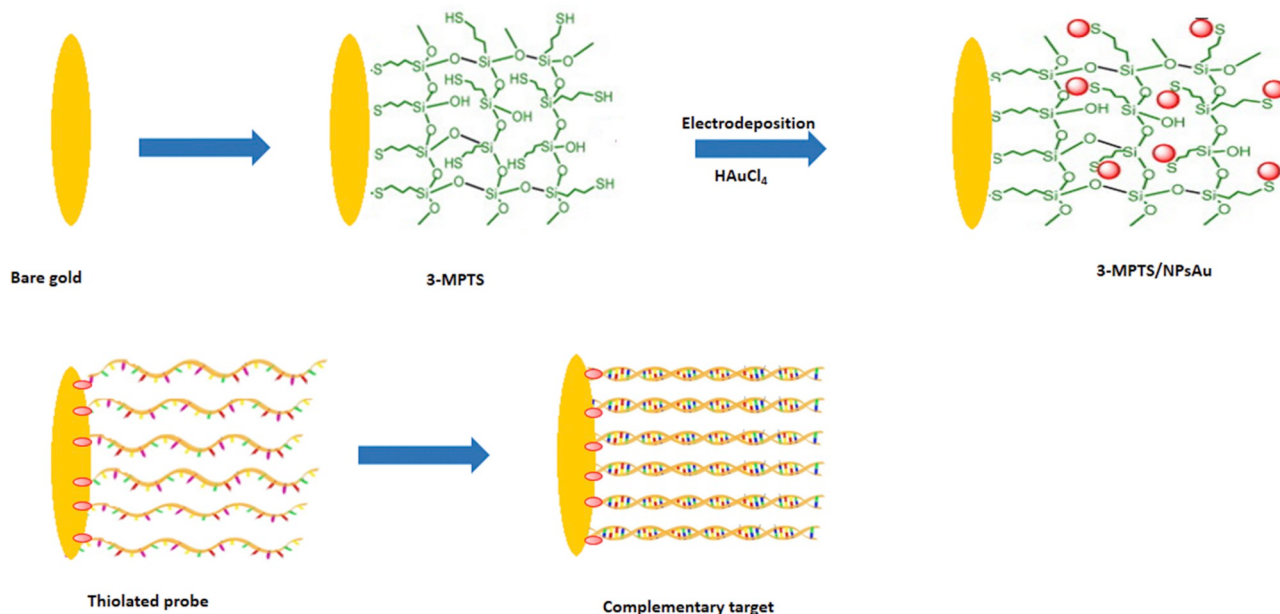


Fig. 1. Schematic representation of the fabrication process of the biosensor.

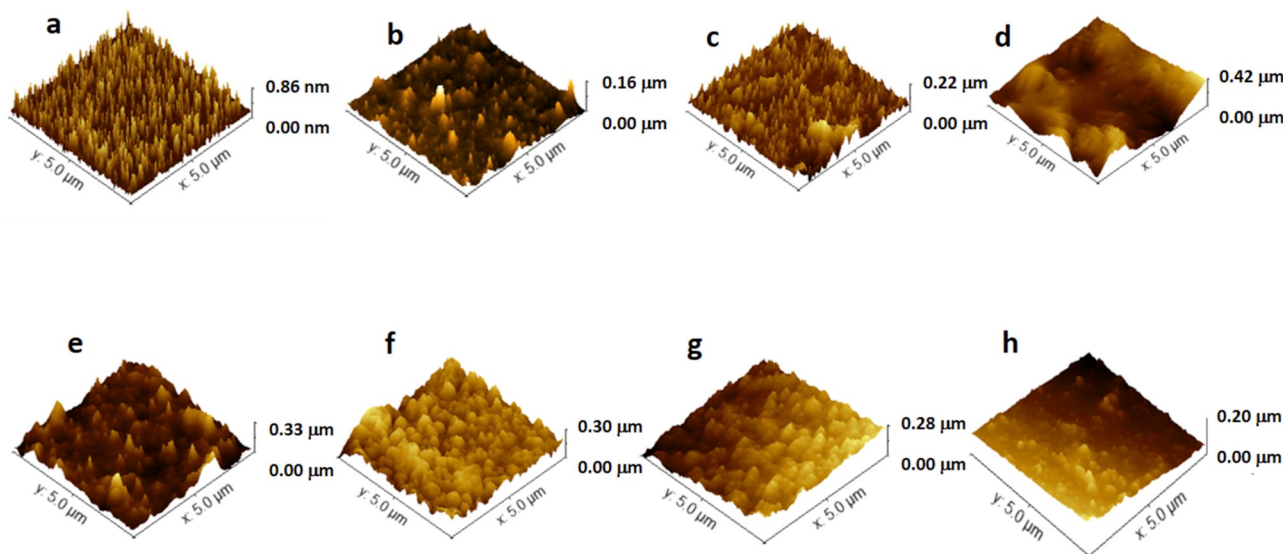


Fig. 2. AFM images for the stepwise modification process: **a)** MPTS_AuNPs; **b)** MPTS_AuNPs-DNA_{probe}; **c)** MPTS_AuNPs-DNA_{probe}-DNA_{worm}; **d)** MPTS_AuNPs-DNA_{probe}-DNA_{serum} Sample **e)** MPTS_AuNPs-DNA_{probe}-DNA_{urine} sample; **f)** MPTS_AuNPs-DNA_{probe}-DNA_{CSF} sample; **g)** MPTS_AuNPs-DNA_{probe}-negative control.

0.5 mol L⁻¹ H₂SO₄ was cycled 30 times in the range of potentials between +1.3 V and -0.2 V at a scan rate of 50 mV s⁻¹. AuNPs were directly deposited on the working electrode previously modified with MPTS. BSA was used in all experiments to avoid non-specific interactions.

2.4. Hybridization analysis

Target DNA was diluted in 10 mM PBS solution (pH 7.4), heated for 3 min at 40 °C (to denature the double-stranded DNA into single strands), and followed by hybridization with a complementary probe. Hybridization was performed by a simple drop-coating method using 2 μL of the target DNA obtained from CSF, urine, and serum at different concentrations. We also evaluated the electrochemical response of the biosensor after exposure to DNA samples containing ascorbic acid and cholesterol as interferents. In these assays, 2 μL of each sample was dropped on the sensor surface for 15 min. The selectivity study was carried out in 10 mM PBS solution (pH 7.4) containing 2 μL of the analyte and of each interferent. The stability was analyzed through the sensor responses after immersion in 5 mM NaOH solution and subsequent immobilization of the target DNA.

2.5. Electrochemical measurements

Cyclic voltammetry and electrochemical impedance measurements were performed using a PGSTAT 128N potentiostat/galvanostat (Autolab, The Netherlands) at room temperature. A three-electrode configuration was employed with an Ag/AgCl reference electrode (saturated KCl), a platinum wire electrode as the auxiliary electrode, and the gold electrode (BGE) as the working electrode. The impedance spectra were recorded within the 100 mHz and 100 kHz frequency range with an alternating voltage adjusted to an amplitude of 10 mV. The CV analysis was performed in the potential range between +0.7 V and -0.2 V at a scan rate of 50 mV s⁻¹. The electrolytic solution was 10 mM K₄[Fe(CN)₆]⁴⁻/K₃[Fe(CN)₆]³⁻ (1:1, v/v) in 20 mM PBS (pH 7.4). All electrochemical measurements were performed in triplicate (n = 3).

2.6. Atomic force microscopy measurements

AFM was used to perform structural analyses of the system in a non-contact mode in the air at room temperature (t = 25 °C) [34]. Cantilevers with an AFM silicon probe (Nanoworld, Japan, resonance

frequency = 300 kHz, spring constant = 42 Nm⁻¹) were used for all measurements. The images were collected with a scanning rate of 1.0 Hz in a scan area of 5 × 5 μm. The images were obtained from at least two macroscopically separated areas. The Gwyddion software was used to analyze the AFM images [35].

3. Results and discussion

3.1. Morphological characterization

Fig. 2a and b shows the BGE surface after modification with MPTS and electrodeposited AuNPs, respectively. Then, surfaces with a roughness of 0.86 nm (Figs. 2a) and 0.16 μm (Fig. 2b) were obtained, suggesting an efficient assembly of the layers. MPTS provides a 3D polymeric network with thiol tail groups that binds to gold surfaces. Also, this matrix provides a biocompatible environment that preserves the biological structure and activity after immobilization process. The polymeric network is essential to AuNPs assembly allowing the incorporation of a high amount of biomolecules that improve the sensor sensitivity. DNA probe immobilization resulted in changes of the surface roughness to 0.22 μm (Fig. 2c) in agreement with previous studies reported in the literature [33,36].

The specific interaction of the sensor and *S. mansoni* samples was evaluated. Significant changes in topography were observed for worm DNA (Fig. 2d), contaminated serum samples (Fig. 2e), urine (Fig. 2f), and LCR (Fig. 2g). The non-complementary target shows a homogeneous distribution on the electrode surface without the presence of aggregates (Fig. 2h). Thus, our results confirm the specificity of the sensor. AFM imaging is in good agreement with CV and EIS results.

3.2. Electrochemical characterization

The electrochemical characterization of the modified electrode was performed through cyclic voltammetry and electrochemical impedance spectroscopy. From Fig. 3a, we can observe a decrease in the anodic peak current after the addition of MPTS (curve b) as compared to the unmodified electrode (curve a). Thus, we obtained an effective assembly of the MPTS layer on the electrode surface. The decreased current can be attributed to the increased steric hindrance of MPTS [37]. An increase of the anodic and cathodic peak currents was verified due to AuNPs electrodeposition, which results in a rise in the surface area and improvement of the conductivity (curve c). Further wane in

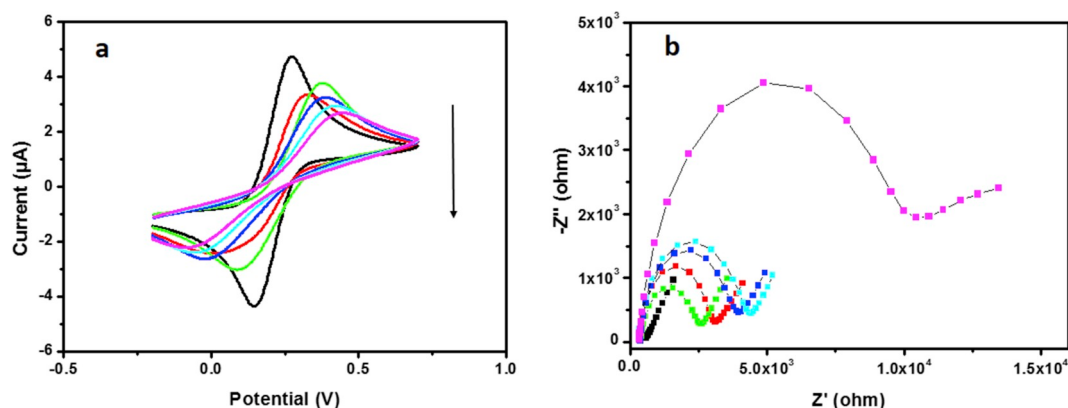


Fig. 3. (a) Cyclic voltammograms of the steps of electrode modification and bio-recognition processes: BGE (□), MPTS (—), MPTS_AuNPs (—), MPTS_AuNPs-DNA_{probe} (—), MPTS_AuNPs-DNA_{probe}-BSA (—), MPTS_AuNPs-DNA_{probe}-genome (—). Cyclic voltammetry scan was initiated at -0.2 V vs. Ag/AgCl in positive direction at 50 mV s⁻¹. (b) Nyquist plots for the stepwise modification process: BGE (□), MPTS (●), MPTS_AuNPs (●), MPTS_AuNPs-DNA_{probe} (●), MPTS_AuNPs-DNA_{probe}-BSA (●), MPTS_AuNPs-DNA_{probe}-genome (●).

the currents is observed after immobilization of the thiol-modified DNA probe on the MPTS-AuNPs system (curve d). BSA was used to block the remaining sites (curve e). In particular, after the hybridization process with the complementary genome, a further decrease in the anodic and cathodic peaks (curve f) was obtained due to additional blockage of the electron transfer to the electrode surface. Fig. 3b shows the impedance spectra of each stage of the sensor platform development in a frequency range between 100 mHz and 100 kHz. Significant differences in the impedance spectra were observed during the assembly of the sensor platform when compared to the unmodified gold electrode since each layer acts as a mass transfer block layer [34,38].

The electrochemical response of the genosensor was evaluated after exposure to different concentrations of target DNA (0.6, 1.3, 2.6, 5.2, 10.4, 20.9, 40.0, and 83.9 pg μL⁻¹). A nonconductive complex formation occurs after the recognition of the contaminated samples that difficult the passage of the faradaic current to the transducer [40]. The voltammetric response of the genosensor is proportional to the number of recognized molecules (Fig. 4a).

A modified Randles equivalent circuit was used [41] to fit the impedance data (Fig. 4b, solid lines). We obtained a good agreement between the equivalent circuit model and the experimental results. The circuit is composed of resistance of the solution (R_{sol}), charge transfer resistance (R_{ct}), constant phase element (CPE), and Warburg diffusion element (W). Ideally, W and R_{sol} represent the mass properties and diffusion characteristics of the redox probe in solution, respectively. CPE represents a non-ideal characteristic of double layer capacitance [42].

The performance of the genosensor was evaluated by the relative

variation of R_{CT} (ΔR_{CT}), providing quantitative information of the DNA hybridization processes. The ΔR_{CT} was calculated according to the following equation:

$$\Delta R_{CT} (\%) = [(R_{CT}(\text{recog}) - R_{CT}(\text{sensor})) / R_{CT}(\text{sensor})] \times 100,$$

where R_{CT}(recog) is the value of the electron transfer resistance of the sensor system after the biorecognition process, and R_{CT}(sensor) is the value of the charge transfer resistance of the genosensor. The R_{CT} element depends on the insulating characteristic at the electrode/electrolyte interface. The filling of the recognition sites (Θ) was calculated using the following equation $\Theta = (1 - R_s) / R_H$, where R_s is the R_{CT} of the biosensor and R_H is the R_{CT} of the biosensor after exposure to different concentrations of the genomic target sequence. Θ as a function of the concentration of the genomic target sequence is shown in Fig. 5. The value increases with increasing target DNA concentration and is found in 0.74 (74%) at a concentration of 83.9 pg μL⁻¹.

A linear behavior was obtained from 0.6 to 10.4 pg μL⁻¹ (inset of Fig. 5b). However, after this range of concentrations, a saturation profile is observed. The changes in R_{CT} are higher than those of other impedance components. Thus, R_{CT} was used to evaluate the interfacial properties of the sensor system.

3.3. Detection of *S. mansoni* in different biological fluids (cerebrospinal fluid, urine, and serum)

CSF is an important laboratory sample for diagnosis in patients with suspected neuroschistosomiasis [13]. Therefore, we performed electrochemical analysis of CSF samples from patients diagnosed with

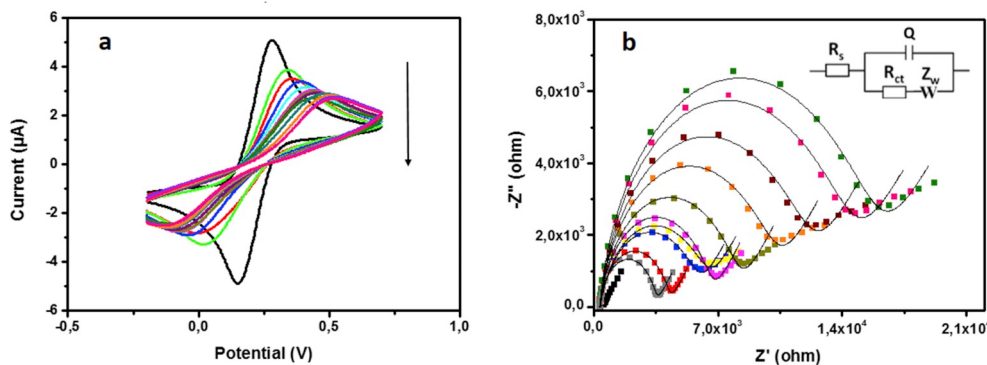


Fig. 4. (a) Cyclic voltammograms of the biosensor exposed to target genomic DNA: BGE (□), MPTS (—), MPTS_AuNPs (—), MPTS_AuNPs-DNA_{probe} (—), MPTS_AuNPs-DNA_{probe}-BSA (—), MPTS_AuNPs-DNA_{probe}-genome (0.6 pg μL⁻¹), Biosensor-genome (1.3 pg μL⁻¹), Biosensor-genome (2.6 pg μL⁻¹), Biosensor-genome (5.2 pg μL⁻¹), Biosensor-genome (10.4 pg μL⁻¹), Biosensor-genome (20.9 pg μL⁻¹), Biosensor-genome (40.0 pg μL⁻¹) and Biosensor-genome (83.9 pg μL⁻¹). (b) Nyquist plots of the biosensor after exposure to different concentrations of target genomic DNA and negative control: BGE (□), MPTS_AuNPs-DNA_{probe}-BSA (■), Biosensor-genome (0.6 pg μL⁻¹), Biosensor-genome (1.3 pg μL⁻¹), Biosensor-genome (2.6 pg μL⁻¹), Biosensor-genome (5.2 pg μL⁻¹), Biosensor-genome (10.4 pg μL⁻¹), Biosensor-genome (20.9 pg μL⁻¹), Biosensor-genome (40.0 pg μL⁻¹), Biosensor-genome (83.9 pg μL⁻¹) and Biosensor-negative control (■). The solid lines represent the simulation data. Inset: Randles equivalent circuit.

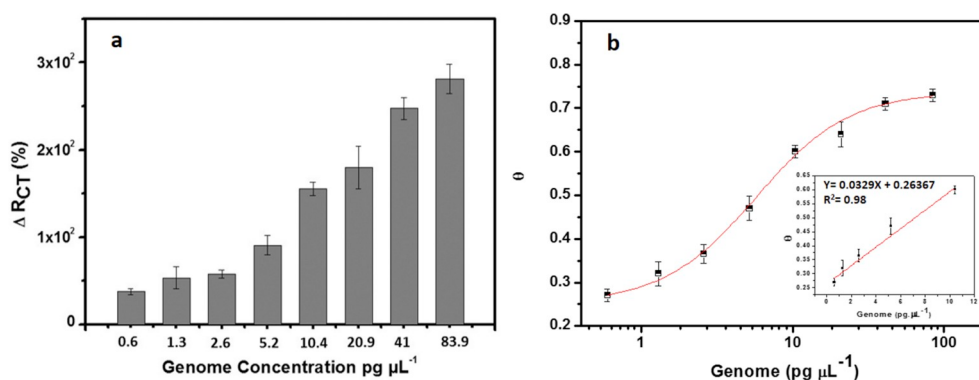


Fig. 5. (a) $\Delta R_{CT}\%$ and (b) θ as a function of different concentrations of the genomic target sequence.

Table 1

Equivalent circuit values obtained from fitted impedance results.

Modified electrode	Genomic target sequence ($\text{pg } \mu\text{L}^{-1}$)	$R_{CT}/k\Omega$	$Q/\mu\text{F}$	n
Bare gold electrode	–	0.16	6.66	0.51
MPTS-AuNPs-DNA probe	–	3.84	2.08	0.86
Sensor system-DNA target	0.6	5.37	3.06	0.83
	1.3	5.85	3.06	0.83
	2.6	6.18	1.08	0.86
	5.2	7.64	1.23	0.86
	10.4	9.59	1.74	0.86
	20.9	11.40	1.63	0.87
	41.0	13.70	1.51	0.88
	83.9	15.10	1.44	0.89
Negative control	–	4.06	0.94	0.91
Sensor system-urine sample	50.0	9.0	0.90	0.91
	40.0	7.18	0.99	0.89
	36.0	6.42	1.06	0.90
	30.0	5.37	0.98	0.90
	27.0	4.94	1.10	0.91
Sensor system-CSF sample	60.0	11.3	2.78	0.85
	47.0	8.57	2.93	0.79
	34.0	6.11	1.92	0.85
	30.0	5.39	2.04	0.83
	25.0	4.50	3.03	0.82
Sensor system-serum sample	42.0	7.58	0.65	0.92
	39.0	6.92	0.55	0.93
	35.0	6.27	0.60	0.93
	31.0	5.63	0.93	0.90
	27.0	4.91	1.10	0.90

neuroschistosomiasis. Table 1 shows the $R_{CT}\%$ values of the samples containing DNA from the adult worm of *S. mansoni*. PCR was used to validate the concentrations of the genomic DNA from *S. mansoni*. There was no significant difference in DNA concentration detection by both measurement methods. The obtained values were used to get a reference standard curve with a limit of detection (LOD) of $0.6 \text{ pg } \mu\text{L}^{-1}$.

Table 2

Comparison of the analytical performances of the proposed genosensor with other biosensors for the detection of *Schistosoma* sp.

Type of biosensor	Analyte	Detect technique	Linear range	Limit	Reference
Piezoelectric immunosensor	<i>S. japonicum</i>	quartz crystal	$10\text{--}200 \text{ ng mL}^{-1}$	5 ng mL^{-1}	[45]
Screen-Printed Biosensor	<i>S. mansoni</i>	cyclic and differential pulse voltammetry	$0.038\text{--}20 \text{ ng mL}^{-1}$	0.038 ng mL^{-1}	[46]
Electrochemical biosensor	<i>Schistosoma</i> egg antigen	anodic stripping voltammetry (ASV)	$0.02\text{--}1 \text{ } \mu\text{g mL}^{-1}$	$0.01 \text{ } \mu\text{g mL}^{-1}$	[47]
Electrochemical immunoassay	<i>S. japonicum</i>	stripping voltammetry	$2 \text{ ng mL}^{-1} - 15 \text{ } \mu\text{g mL}^{-1}$	1.3 ng mL^{-1}	[48]
Spectroelectrochemical immunoassay	Soluble Egg Antigen (SEA)	UV/Vis spectrometry and cyclic voltammetry	–	$3.31 \times 10^{-5} \text{ ng mL}^{-1}$	[49]
Electrochemical immunoassay	<i>S. japonicum</i>	linear sweep voltammetry (LSV)	1:5000–1:100	1:6457	[50]
Metalloimmunoassay	<i>S. japonicum</i> antibody	anodic stripping voltammetry (ASV)	$6.4 \text{ ng mL}^{-1} - 100 \text{ } \mu\text{g mL}^{-1}$	3.0 ng mL^{-1}	[51]
Genosensor	<i>S. mansoni</i>	Cyclic voltammetry and Impedance spectroscopy electrochemical	$27\text{--}50 \text{ pg } \mu\text{L}^{-1}$ (urine)	6.0 ng mL^{-1}	This work
			$25\text{--}60 \text{ pg } \mu\text{L}^{-1}$ (CSF) $27\text{--}42 \text{ pg } \mu\text{L}^{-1}$ (serum)		

Also, *S. mansoni* can be detected in other biological fluids, such as urine and serum. Urine is not a conventional sample for diagnosis of *S. mansoni* but is promisor for DNA research. Urine has the great advantage of not being a sample of invasive collection and being easily obtainable. In this study, CSF, urine, and serum samples were analyzed under the same parameters. A significant increase in the R_{CT} values was obtained after the recognition of positive samples.

The sensitivity of the genosensor was evaluated through a calibration curve obtained from impedimetric responses obtained after its exposure to the target sample. The biosensor has demonstrated excellent sensitivity for detecting the genomic target at low concentrations in CSF, urine, and serum. Table 1 shows the fitted impedance parameters for different samples. The proposed biosensor is effective at low sample volumes ($2 \text{ } \mu\text{L}$). The comparison of the proposed genosensor with other previous biosensors is shown in Table 2. Thus, the present biosensor demonstrated the feasibility for rapid molecular detection of *S. mansoni* compared with other methods.

3.4. Selectivity and stability of the genosensor

The adsorption of interfering molecules on the electrode surface is a factor that can modify the electrochemical response [43] and suppress the electrode response [44]. The interferents tested in this work were ascorbic acid (AA) and cholesterol. Fig. S2 shows the impedimetric response of the genosensor in the presence of the complex mixtures (analyte + interfering substances). The results indicate that the response of the genosensor was not significantly affected in the presence of substances usually found in biological fluids such as AA and cholesterol.

The reusability of the biosensor was evaluated immersing the genosensor in a solution of 5 mM NaOH for 5 min to unbound the genome. This process was repeated two more times on the same day (Figs. S3a and S3b). After 48 h of storage under refrigeration, the analytical tests were carried out under the same experimental conditions (Figs. S3c and

S3d). Thus, it was possible to confirm the stability of the genosensor. Therefore, the proposed sensor is a useful platform for rapid and specific detection of *S. mansoni* in different biological fluids as CSF, urine, and serum.

4. Conclusion

A novel electrochemical sensing platform based on a self-assembled layer of 3-mercaptopropyltrimethoxysilane and electrochemically synthesized AuNPs for diagnosis of schistosomiasis was successfully developed. The impedimetric and voltammetric analyses were effective to evaluate the process of modifying the electrode surface. The developed genosensor showed excellent performance in the detection of target genomic sequences at low concentrations in CSF, urine, and serum with a limit of detection (LOD) of $0.6 \text{ pg } \mu\text{L}^{-1}$. The developed system is presented as a promising alternative for the label-free detection of *Schistosoma* in CSF, urine, and serum.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.talanta.2019.05.111>.

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