

# Electrochemical DNA Biosensor for Sequences Related to the Human Papillomavirus Type 16 using Methylene Blue

Elaine VM Souza<sup>1,3\*</sup>, Gustavo A Nascimento<sup>1</sup>, Nataly Amorim de Santana<sup>1</sup>, Danielly S. Campos-Ferreira<sup>1</sup>, Juliana de Araújo Bibiano<sup>1</sup>, Mariana S Arruda<sup>1</sup>, Danyelly Brunaska<sup>1,2</sup> and José L Lima-Filho<sup>1,2</sup>

<sup>1</sup> Laboratory of Immunopathology Keizo Asami (LIKA), Federal University of Pernambuco (UFPE), AV. Moraes s/n, 50670-901 Recife, PE, Brazil

<sup>2</sup> Department of Biochemistry, Federal University of Pernambuco (UFPE), AV. 12 Professor Moraes Rego s/n, 50670-901 Recife, PE, Brazil

<sup>3</sup> Nursing Department, Universidade Federal de Alagoas-UFAL, Arapiraca 15 Campus, AV. Manoel Severino Barbosa, s/n, Bom Sucesso-Arapiraca-AL, CEP:57309-005, Brazil

## Abstract

Several studies show that infections with high-risk human papillomaviruses (HPV), mainly HPV type 16, can lead to the development of tumors as the cervical cancer. A rapid and precise diagnosis of the precancerous lesions by HPV is extremely important for a successful treatment. The present paper describes an electrochemical DNA biosensor for specific sequence detection of the E1 HPV 16, using the methylene blue (MB) as indicator hybridization. In order to develop the sensor, a 21-mer ssDNA probe related to E1 HPV gene was immobilized on the work electrode (WE). The obtained optimum conditions for immobilization of HPV E1.16P on the activated WE was with an immobilization time of 5 min and probe concentration of 8  $\mu$ M. The hybridization between the probe and its complementary sequence was studied by Differential Pulse Voltammetry (DPV). Some hybridization experiments with non-complementary oligonucleotides were carried out to study the selectivity of the biosensor. The results showed that this DNA biosensor could be used for detection E1 HPV gene due to selectivity of the electrochemical measurement system. A detection limit of probe immobilized on electrode to its complementary sequence was 1.49 nM.

**Keywords:** HPV; DNA biosensor; Methylene blue

## Introduction

The relationship between human papillomavirus (HPV) and the occurrence of tumors was first described in the cervix, and it is currently known that HPV is a necessary condition for the development of cancer in this region [1]. Virtually 100% of cervical cancers, the second commonest cancer in women worldwide, contain the HPV DNA sequences from a high-risk oncogenic genital HPV [2,3]. Several experimental evidences show that infections associated with HPV can lead to the development of tumors in a variety of sites as the anogenital region of male and female, breast, skin, nails, lungs and the head and neck [4,5].

The HPV can infect keratinocytes of skin and mucosa and it a double stranded DNA with approximately 8 kb [6]. Papillomaviruses are not classified by serotype but by genotype about 130 HPV types are known according to the sequence of the viral DNA [3] and new types are continually being found [7,8]. The HPV can be classified into high (e.g., HPV-16 and 18) or low-risk (e.g., HPV-11) [9], related with their potential to induce a cancer [10].

The early diagnostic of the presence of these viruses has been a very important tool to prevent a cancer development, and molecular diagnostic system has also improved within last few years. Molecular techniques are sensitive, reliable and effective [11], it is not yet possible to introduce these techniques in public health in developing countries because of their high cost. It is necessary to research and develop alternative methods for rapid and inexpensive detection of HPV in order to standardize the laboratory in public health for assisting the early diagnosis of cervical cancer [11].

Currently there has been much interest in developing electrochemical DNA biosensors for rapid and specific diagnosis of bacterial and viral infections, genetic diseases and among other applications [12-16]. All reports have revealed that the electrochemical biosensors has advantages such as speed, simplicity, low cost, high sensitivity, and relatively simple instrumental [17,18] and the possibility of real-time

detection [19,20]. The DNA biosensors are effective in converting the event of hybridization of DNA in a signal that can be measured. DNA hybridization usually occurs between an oligonucleotide with a known sequence (probe) and an unknown complementary strand of nucleic acid from solution (target). The hybridization of the probe to the target sequence can be detected using appropriate hybridization indicators [21]. The hybridization indicator can bind DNA through reversible physical intercalation between base pairs or through electrostatic interaction [22]. Methylene blue (MB) is an aromatic heterocyclic that has been widely used as an electrochemical intercalator to monitor the DNA hybridization reaction, because MB binds specifically to guanine bases in ssDNA [23] and a lower current signal is observed following DNA hybridization. In this study, a simple method was proposed using MB to monitor the hybridization reaction. The electrochemical DNA biosensor based on the gene of HPV 16 was proposed. Due to the key role of HPV 16 in induction of cancer of the cervix, development low cost and small size device associated sensitive, specific and rapid detection methods will certainly provide important new tools for the diagnosis of disease.

## Experimental Materials

Electrochemical experiments were performed using AUTOLAB PGSTAT 30 (METROHM AUTOLAB, Netherlands) and GPES 4.7

**\*Corresponding author:** Souza EVM, 1 Laboratory of Immunopathology Keizo Asami (LIKA), Federal University of Pernambuco (UFPE), AV. Moraes s/n, 50670-901 Recife, PE, Brazil, Tel: +55 82 9931 0827; Fax: 82-3214 +55 1172; E-mail: [elainevms@yahoo.com.br](mailto:elainevms@yahoo.com.br)

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software package. The utilized two-electrode system was composed of a pencil graphite electrode (Total length 3 cm and 2.5 mm diameter) as the working electrode. In addition, reference electrode was screen-printed using Ag/AgCl ink (Electrodag-Acheson, USA) under gold wire and then dried at 60°C.

The experiments were performed in triplicate. All the synthetic oligonucleotides (targets and probes) were purchased from Integrated DNA Technologies (Belo Horizonte, Brazil) as lyophilized powder with the following sequences:

DNA probe\_HPVE1.16P (21-mer sequence): GCAAAGGCAG-CAATGTTAGCA

DNA target\_HPVE1.16T (21-mer sequence): TGCTAACATT-GCTGCCTTTGC

DNA non-complementary (24-mer sequence): CCAAAGCCTT TCAAAAAAAGATTTCCA

Mix sequence: (DNA target\_HPVE1.16R and DNA non-complementary).

The oligonucleotide stock solutions were prepared with 0.5 M acetate buffer (pH 5.0) and kept frozen. Tris base was obtained from Promega (USA), and sodium acetate was obtained from Sigma (USA). The UltraPure distilled Water was purchased from Invitrogen (USA), and reagents were of analytical grade.

## Bioinformatics

The HPV 16 E1 nucleotides sequence was collected from the National Center for Biotechnology Information (NCBI) database, corresponding to GenBank accession number NC\_001526. Subsequently, HPV16 E1 was aligned using the software CLC Combined Workbench v 3.6.1 software. Target sequence (HPVE1.16T) and non-complementary sequences were also constructed using this software.

## Preparation of the working electrode (WE)

The pencil graphite (type 4B, total length of 3 cm and diameter of 2.5 mm) was polished with emery impregnated disc to obtain a smooth surface. The body of the working electrode (WE) was coated well with silicone, resulting in a free area for immobilization of 28.50 mm<sup>2</sup>. The graphite electrode was washed with ultra-pure water to remove possible contaminants in the surface of the electrode. The polished surface of the working electrode was activated by applying a potential [24] of 1.8 V for 5 min in 0.5 M acetate buffer (pH 5.0).

## Immobilization of the oligonucleotides probe

Immobilization of the HPVE1.16P (probe) was conducted following electrochemical pre-treatment of the WE.

The probe was immobilized on the activated WE by applying 0.5 V (24) within different period of time (1-10 minute) in 0.5 M acetate buffer solution (pH 5.0) containing oligonucleotides. The electrode was then rinsed with ultrapure water.

## Hybridization

The hybridization was performed by immersing the probe-modified WE (WE\_HPVE.16P) in a solution (50 µl) containing a known amount of the target oligonucleotides in acetate buffer (0.5 M) pH 5.0 with 300 rpm stirring. The hybridization was allowed to proceed for 15 minutes at 57°C. The electrodes were washed with Tris-HCl buffer (pH 7.0) for removing non-hybridized sequences. The same protocol was applied for the hybridization of probe with non-complementary sequences.

## MB reduction measurements

The probe-modified WE was immersing into the 500 µM solution of MB for 10 minutes without applying any potential on the electrode. WE was rinsed with 20 mM Tris-HCl buffer (pH 7.0), MB reduction sign was measured at DPV. The same protocols were applied for probe modified WE following hybridization with complementary or non-complementary sequences. All the experiments were performed in an electrode chemical cell.

## Electrochemical analysis

The electrochemical analysis were carried out by Differential Pulse Voltammetry (DPV) (from -0.9 up to -0.1 V) in Tris-HCl buffer pH 7.0, with modulation amplitude of 50 mV and scan rate of 20 mV/s. The results were treated using the GPES software moving average baseline correction, using a "peak width" of 0.01.

## Statistics analysis

Experimental data were analyzed by the software Statistica 8 using parametric tests. To evaluate statistical significance, the one-way ANOVA was used for the comparison of multi independent group data. A level of  $p < 0.05$  was assigned as significant.

## Results and Discussion

### Preliminary Investigation

The pencil graphite electrode is successfully applied as the working electrode [25,26]. Several studies demonstrate the utility of pencil electrodes and DNA probe for amperometric measurements of trace nucleic acids [13-15,27-29]. These transducers attract more attention than other methods due to their high sensitivity, small dimensions, low cost, and compatibility with microfabrication technology [28,30-32].

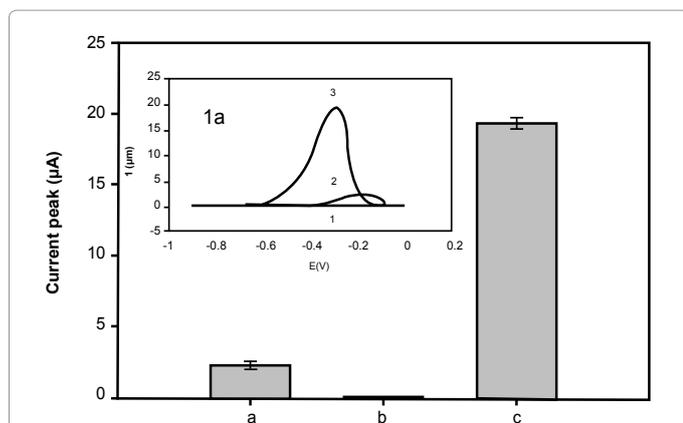
In the present work, we used this procedure for detection of oligonucleotides corresponding to E1 HPV 16 gene, in order to aid the diagnostic the HPV virus. The pretreatment of the carbon surface increases its roughness and hydrophilicity which is essential for DNA adsorption on the electrodes [14].

The activated WE without MB showed no significant current peak, while the activated WE with MB showed about  $2.25 \pm 0.21$  µA (Figure 1). However, the MB voltammetric signal showed a higher peak with the WE\_HPVE.16P ( $19.30 \pm 0.39$  µA), due to guanines free bases interaction of with MB. The increased was 84.44% when compared with activated WE with MB.

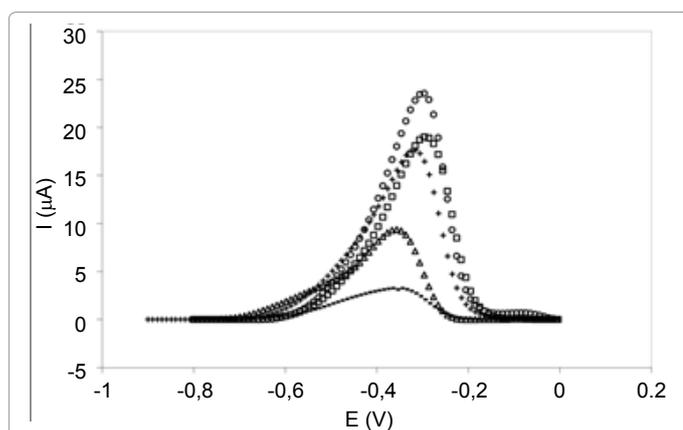
### Effect of probe concentration

In the electrochemical DNA biosensor was immobilized E1 HPV 16 probe (WE\_HPVE.16P) on the WE surface. The immobilization of the probe was with a potential 0.5 V in acetate buffer (0.5 M/pH 4.8) solution with increasing concentration of HPVE1 16P (1.5-20 µM). Previous studies demonstrate that this potential enhances stability of the immobilized probe, due to the electrostatic attraction between the positively carbon surface charged and the negative hydrophilic sugar-phosphate backbone charged with the bases oriented toward the solution ready to hybridize with the target [33].

According to Figure 2, the concentrations 8 µM ( $19.3 \pm 0.36$  µA), 15 µM ( $21.9 \pm 4.25$  µA) and 20 µM ( $17.4 \pm 1.95$  µA) presented the highest signal and parametric statistical analysis (ANOVA-significant level  $p < 0.05$ ) demonstrated that the MB reduction signal was statistically similar. Therefore, concentration of 8 µM was suggested as good for



**Figure 1:** Histogram of MB reduce signal of the (a) activated WE with MB, (b) activated WE without MB and (c) activated probe-modified WE with MB (WE\_HPVE.16P). Differential pulse voltammetry method was used to analyses the current peaks in the following conditions: initial potential -0.9 V and end potential -0.1 V; modulation amplitude 50 mV and scan rate 20 mV/s. All results were plotted with the means of experiments performed.



**Figure 2:** Effect of HPV probe concentration on activated WE. Differential pulse voltammograms on the MB reduction signal in HPV sequences; (-) DNA probe\_HPVE1.16F 1.5 µM; ( $\Delta$ ) DNA probe\_HPVE1.16F 5 µM; ( $\square$ ) DNA probe\_HPVE1.16F 8 µM; ( $\circ$ ) DNA probe\_HPVE1.16F 15 µM; (+) DNA probe\_HPVE1.16F 20 µM. Experimental conditions were described in Figure 1. Experiments conditions were as cited in Figure 1.

the probe immobilization on the WE because of lower concentration and also presented better reproducibility than the concentration the 15 µM and 20 µM.

### Effect of immobilization time (probe)

The probe immobilization time (HPVE1.16P) was investigated on WE in the 8 µM concentration at different times (1-10 min). According to Figure 3, 1 minute presented a current peak of  $3.65 \pm 0.40$  µA, while the time of 3, 5 and 10 minutes showed respectively signal of  $8.24 \pm 0.22$  µA,  $19.30 \pm 0.39$  µA and  $8.80 \pm 0.53$  µA, respectively. The results showed that the MB reduction signal increased with probe immobilization time up to 5 min (1-5 min), and then decreased significantly with 10 min (Figure 3). This decrease is attributed to the massive accumulation of the probe on the electrode surface, occurring probably overlapping probes, leading to a lower availability of guanine bases for MB interaction. Therefore, 5 min was suggested as optimum time for probe immobilization (8 µM). This immobilization time of 5 minutes is also found in a study of the human interleukine-2 in the construction of biosensors for the identification of recombinant plasmid with the

gene sequence of interleukine-2 [28]. Results also were found in an electrochemical DNA biosensor with pencil graphite electrode for diagnosis of hepatitis B [31], detection of specific sequences of human papilloma virus (14), label-free electrochemical detection of the specific oligonucleotide sequence of dengue virus type 1 [15], electrochemical DNA biosensor for the detection of human papillomavirus E6 gene inserted in recombinant plasmid [34] and detection of Hepatitis C virus [35].

These results showed that probe concentration and the immobilization time affect the performance of the biosensor; therefore, these parameters are important for biosensor development.

### Hybridization Detection

Detection of the hybridization was accomplished using reduction of the accumulated MB, that discrimination of the hybrid (DNA-DNA) from the probe (WE\_HPVE.16P). The hybridization events were monitored by DPV utilizing 15 minutes with six different target concentrations (4, 7, 11 and 12 nM). MB reduction signal on the WE was around -0.3 V in 20 mM Tris-HCl buffer solution (pH 7.0). In this study, hybridization step was from the annealing with 300 rpm. The MB peak current from the probe-modified WE ( $19.3 \pm 0.39$  µA) before hybridization were bigger than the WE after hybridization in different concentrations. This happens due the interaction of the MB with single-stranded ssDNA by electrostatic binding [36].

The strong affinity of MB to the free guanine bases exposed to the exterior of ssDNA explains this decreased of the MB signal [37] in the double helix of the DNA hybrids. Thus, ssDNA and dsDNA identifies through the difference of the electrochemical signals. Several articles demonstrate that, electroactive indicators, as the MB, have frequently been used to characterize ssDNA and dsDNA by DPV [38].

The decrease in the MB signal is attributed to less MB accumulation on the DNA hybrid that causes the inaccessibility of MB to the guanine bases. It also is due to a steric inhibition of the reducible groups of MB packed between the bulky double helix of DNA hybrids [30].

After hybridization, the peak current response of MB reduction decreased around 13.5%-41.6%, according to the target concentration. However, a complementary DNA probe sequence electrostatically immobilized onto a glassy carbon electrode modified with chitosan for dengue using ferrocenium shows a difference of around 26%, before and after of the hybridization [39].

The experiment with non-complementary oligonucleotides was carried out to assess whether the suggested DNA sensor is selective via hybridization to the target. The current peak of MB observed at the WE\_HPVE.16P after hybridization with non-complementary oligonucleotides was approximately equal the signal of the probe immobilized. The results demonstrated that the interaction between this sequence and immobilized probe did not lead to significant decrease in the MB reduction signal due to the absence of hybridization, being similar to the probe signal.

Electrochemical detection of HPV utilizing potential observed a slight decrease of the MB signal that is attributed to the small and irregular hybridization of non-complementary oligonucleotides with the probe sequences [14]. Irregular hybridization was not visualized in our study using in the hybridization process temperature.

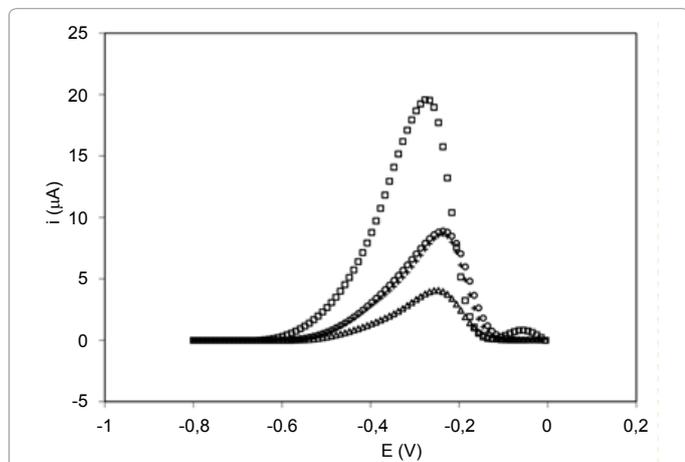
The experiments with mix of oligonucleotide sequence (complementary and non-complementary) demonstrated that the signal was equal to the WE after hybridization with complementary

sequence (Figure 4). This is attributed to the process of hybridization of the probe with the HPVE1. 16T sequence present in the mix. These observations indicated that only complementary sequence could form a stable double strand DNA on the probe modified and thus demonstrating the selectivity of the biosensor proposed.

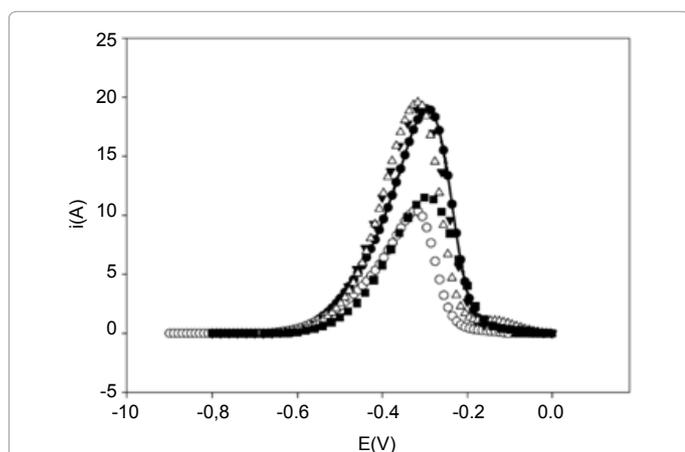
The biosensor for detection of human interleukine-2 gene [30], dengue [15] and bovine papillomavirus [39] also present selectivity between complementary and non-complementary sequences utilizing potential to hybridization

### Quantitative analysis of target

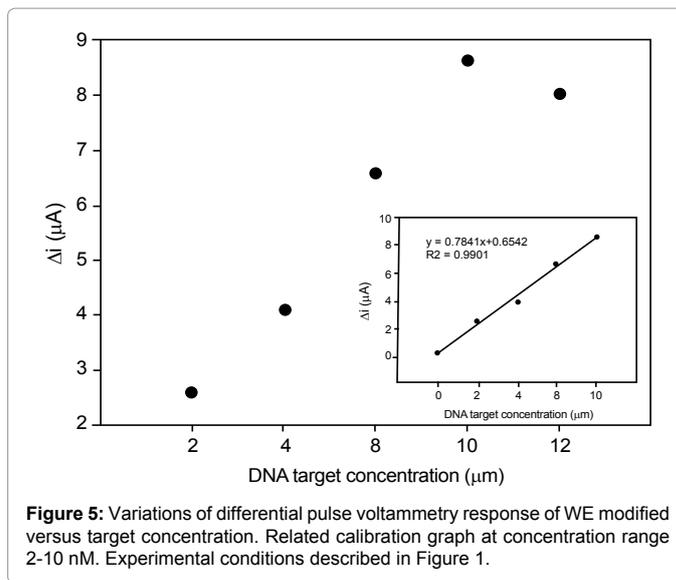
The biosensor sensitivity was investigated by varying the concentration of the specific target, the difference between MB signals on the probe-modified WE in the absence and presence of target ( $\Delta I$ ) is increased with increasing the target concentration up to 10 nM (Figure 5). As seen in inset of figure 5, the calibration graph was linear



**Figure 3:** Effect of period of time on probe immobilization. Differential pulse voltammograms on the MB reduction signal in HPV sequences; ( $\Delta$ ) 1 minute; (+) 3 minutes; ( $\square$ ) 5 minutes ( $\circ$ ) 10 minutes. Experimental conditions were described in Figure 1. Experiments conditions were as cited in Figure 1.



**Figure 4:** Differential pulse voltammograms on the MB reduction signal in HPV sequences; ( $\bullet$ ) DNA probe\_HPVE1.16F 8  $\mu$ M before hybridization; ( $\circ$ ) DNA probe\_HPVE1.16F 8  $\mu$ M after hybridization with oligonucleotides target\_HPVE1.16R 8 nM; ( $\Delta$ ) DNA probe\_HPVE1.16F 8  $\mu$ M after hybridization with DNA noncomplementary 8 nM; ( $\blacksquare$ ) DNA probe\_HPVE1.16F 8  $\mu$ M after hybridization with mix sequences 8 nM (DNA targeted\_HPVE1.16R and DNA non-complementary), ( $\blacktriangledown$ ) DNA probe\_HPVE1.16F 8  $\mu$ M after hybridization with 0.5 M acetate buffer (pH 5.0). Experimental conditions were described in Figure 1.



**Figure 5:** Variations of differential pulse voltammetry response of WE modified versus target concentration. Related calibration graph at concentration range 2-10 nM. Experimental conditions described in Figure 1.

between 2-10 nM with correlation coefficient of 0.9901. The regression equation was  $\Delta I (\mu A) = 0.7841x + 0.6542$ , with a detection limit of 1.49 nM. The relative standard deviation over three independently modified probe electrodes measured at 8 nM of target was around 0.41% indicating a remarkable reproducibility of the detection method. Such detection limit compares favorably with those reported for the other electrochemical hybridization methods, such as, 5.78 nM detection of human interleukine-2 gene [30].

### Conclusion

The effective immobilization of probe oligonucleotides related to HPV 16 on the electrode surface treaty was achieved. The optimum conditions for the construction of the working electrode were in the concentration of 8  $\mu$ M and time of 5 minutes.

DNA biosensors utilizing graphite electrode for the detection of hybridization are useful for identifying specific nucleotide sequences (HPV) with selectivity. The hybridization of oligonucleotides on graphite was with annealing temperature using a MB reduction signal. The detection limit for this biosensor with complementary DNA target was of 1.49 nM within a hybridization time of 15 minutes.

New researches in biosensor techniques and hybridization methodology at the molecular level may contribute to future advancements in specific HPV diagnosis.

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