

Direct detection and discrimination of double-stranded oligonucleotide corresponding to hepatitis C virus genotype 3a using an electrochemical DNA biosensor based on peptide nucleic acid and double-stranded DNA hybridization

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Abstract Development of an electrochemical DNA biosensor for the direct detection and discrimination of double-stranded oligonucleotide (dsDNA) corresponding to hepatitis C virus genotype 3a, without its denaturation, using a gold electrode is described. The electrochemical DNA sensor relies on the modification of the gold electrode with 6-mercapto-1-hexanol and a self-assembled monolayer of 14-mer peptide nucleic acid probe, related to the hepatitis C virus genotype 3a core/E1 region. The increase of differential pulse voltammetric responses of methylene blue, upon hybridization of the self-assembled probe with the target ds-DNA to form a triplex is the principle behind the detection and discrimination. Some hybridization experiments with non-complementary oligonucleotides were carried out to assess whether the developed DNA sensor responds selectively to the ds-DNA target. Diagnostic performance of the biosensor is described and the detection

limit was found to be 1.8×10^{-12} M in phosphate buffer solution, pH 7.0. The relative standard deviation of measurements of 100 pM of target ds-DNA performed with three independent probe-modified electrodes was 3.1%, indicating a remarkable reproducibility of the detection method.

Keywords DNA biosensor · Hepatitis C · PNA probe · Triplex formation · Gold electrode

Introduction

The use of nucleic acid recognition layers in biosensor design represents a new and exciting area in analytical chemistry. The specificity of DNA hybridization devices (i.e., their ability to respond to the target sequence in the presence of non-complementary strands) depends primarily on the selected probe and secondarily upon the hybridization. Peptide nucleic acid (PNA) is an extremely good structural mimic of DNA (or RNA) in which the negatively charged sugar–phosphate backbone is replaced with a neutral, acyclic, and achiral pseudopeptide backbone composed of repeating *N*-(2-aminoethyl)glycine units linked by amide bonds. The nucleobases are attached to the backbone through methylene carbonyl linkages. PNA may be used in many of the same applications as traditional synthetic DNA or DNA analogs. However PNA chains exhibit superior hybridization properties towards complementary ss-DNA (duplex formation), as well as improved chemical and enzymatic stability, faster hybridization, tighter binding—especially at low ionic strength, and greater specificity compared with natural nucleic acids [1–6]. Furthermore PNA oligomers are able to form very stable duplex structures

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The cysteine-tethered PNA oligonucleotide was dissolved in 0.1% trifluoroacetic acid and kept frozen at -20°C . More diluted solutions of the PNA probe were prepared by using 0.5 M acetate buffer solution (pH 4.8).

The stock solutions of the DNA oligonucleotides (500 $\mu\text{g/ml}$) were prepared with TE buffer solution (10 mM Tris-HCl, 1 mM EDTA, pH 8.0) and kept frozen. More diluted solutions of the oligonucleotides were prepared by using 0.2 M phosphate buffer solution (pH 7.2). Other chemicals were of analytical reagent grade. The distilled, deionized, and sterilized water was used in all solution preparation. All the experiments were performed at room temperature in an electrochemical cell.

Preparation of double-stranded oligonucleotides

To prepare double-stranded oligonucleotides, equal concentrations (50 μM) of the complementary oligonucleotide pairs were effectively mixed in a microtube. The mixed solution was heated at 95°C in a water bath for 10 min and then was allowed to cool down gradually to room temperature.

Apparatus

All electrochemical experiments were performed in a 10-ml-volume cell, connected to an AUTOLAB PGSTAT 30 electrochemical analysis system and GPES 4.7 software package (Eco Chemie, the Netherlands). The utilized three-electrode system consisted of a polycrystalline gold working electrode (Bioanalytical Systems, UK) with a 3-mm diameter, a saturated calomel electrode (SCE) as the reference electrode, and a platinum wire as the auxiliary electrode. Supporting electrolyte was deoxygenated by purging with nitrogen gas for 10 min prior to measurements, and the cell was blanketed with nitrogen gas for the duration of the measurements.

Procedures

Preparation of the working electrode

Gold disk electrode (AuE, 7 mm² area) was polished with wet alumina slurry for at least 10 min and rinsed repeatedly with water. For complete removal of fouling layer (self-assembled monolayers) on the AuE, the polished surface was dipped in 1:1 EtOH-H₂O solution containing 0.5 M NaBH₄ for 10 min [36, 38]. The electrode was treated by cycling of the potential between -0.3 and 1.5 V at a scan rate of 0.1 V s^{-1} until the cyclic voltammogram did not change, indicating that the electrode surface was clean. The electrode was then washed with water, and placed upside-down in a humidity chamber. A 6- μl droplet of 5 μM PNA

probe aqueous solution was deposited onto the pre-treated gold electrode for self-assembly. This probe self-assembled AuE was rinsed with water and then dipped in 1 mM aqueous solution of 6-mercapto-1-hexanol (MCH) at room temperature for 30 min. Evaluation of the PNA self-assembled monolayer formation was performed by using the electrochemical signal of the $\text{Fe}(\text{CN})_6^{3-}/\text{Fe}(\text{CN})_6^{4-}$ redox system.

Probe-DNA hybridization

Two types of hybridization were tested in this research, namely drop hybridization and pre-heated electrode drop hybridization. Drop hybridization was carried out by dropping 6 μl of target ds-DNA onto the PNA self-assembled monolayer (PNA-SAM) AuE for 20 h in a humidity chamber, and then the electrode was rinsed with PBS. In the pre-heated electrode drop hybridization the PNA self-assembled AuE was immersed in Tris-EDTA buffer at 85°C for 5 min and then a 6- μl droplet of target ds-DNA in PBS was deposited onto the PNA self-assembled AuE in a humidity chamber. Hybridization of the probe with non-complementary ds-DNA sequences was carried out following the same method as described above.

Labeling of the hybrid

MB was interacted with the probe or hybrid by immersing the modified electrode surface into a stirring 20 mM Tris-HCl buffer (pH 7.0) solution, containing 20 μM MB and 20 mM NaCl, for 5 min without applying any potential. After accumulation of MB, the electrode was rinsed with a 20 mM Tris-HCl buffer (pH 7.0) solution for 10 s.

Voltammetric measurements

The reduction signal of the accumulated MB was measured by using differential pulse voltammetry (DPV) in the 20 mM Tris-HCl buffer (pH 7.0) solution containing 20 mM NaCl and scanning the electrode potential between 0.10 and -0.50 V at a pulse amplitude of 25 mV. The raw data were treated by using the Savitzky and Golay filter (level 2) of the GPES software, followed by the GPES software moving average baseline correction using a 'peak width' of 0.01 between non-hybridized and hybridized status.

Results and discussion

Detection of target DNA as ds-oligonucleotide

Detection of ds-DNA target is accomplished by monitoring of the difference occurring between the DP voltammetric

responses of MB accumulated on the PNA-SAM AuE, in the absence and presence of target. The DP voltammograms for the probe-modified AuE before and after hybridization with 1 μM complementary ds-oligonucleotide confirmed that a significant increase in the MB signal from -200 ± 6.7 nA to -630 ± 19.3 nA was observed (not shown). The enhancement of the peak current is attributed to the hybridization of the PNA probe with the ds-oligonucleotide and indicates that more MB molecules interact with the PNA-SAM/ds-DNA hybrid.

Effect of some experimental variables

SAM formation time and probe concentration are important factors affecting the SAM formation on the electrode. The effect of these parameters was studied by cyclic voltammetric measurement in $\text{K}_3\text{Fe}(\text{CN})_6/\text{K}_4\text{Fe}(\text{CN})_6$ redox solution. Indeed as the SAM formation is extended the redox signal of $\text{K}_3\text{Fe}(\text{CN})_6/\text{K}_4\text{Fe}(\text{CN})_6$ is decreased. The results showed that 2 h and 25 μM are a suitable time and concentration for the probe self-assembly, respectively.

The two hybridization procedures, namely drop hybridization and pre-heated electrode drop hybridization, were applied to the PNA-modified AuE. The results depicted in Fig. 1 indicate that the pre-heating of the electrode had a significant effect on the hybridization efficiency, because a 70% increase was observed in the MB signal, compared with the other method. Therefore, the pre-heated electrode drop hybridization method is recommended for future

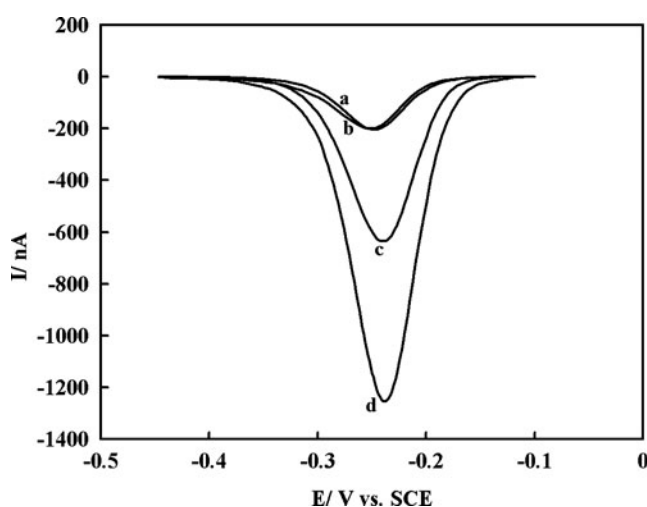


Fig. 1 Differential pulse voltammograms of accumulated MB on the PNA probe-modified AuE: before hybridization **a** without heating or **b** after 10 min heating at 85 $^{\circ}\text{C}$; and after hybridization using **c** non-heated electrode drop hybridization or **d** pre-heated electrode drop hybridization methods with 5 μM ds-HCV3a target oligonucleotide solution. Electrode conditions: a 6- μl droplet of 25 μM PNA solution dropped onto the pre-treated AuE

experiments. Note that the probe-modified AuE subjected to the heating procedure showed no significant change in the MB signal when compared with the non-heated probe-modified electrode. This indicates that the MB signal on the probe-modified electrode does not depend on the electrode heating and also no SAM peeling off occurred.

There are many reports about the effect of thiolated PNA probe density on hybridization efficiency and kinetics [39–41]. Probe density has a key role in the function of biosensor. In most cases, high probe densities are desired in order to increase the signal strength and thus to improve the detection limits. However, binding efficiency may be decreased at higher probe densities. The probe density can be controlled by altering the probe concentration on the gold surface during the probe film fabrication. We used various concentrations of the probe in order to obtain an optimal PNA probe density for DNA hybridization. The resulting MB DP voltammograms are shown in Fig. 2. As can be seen in the inset in Fig. 2, by increasing the probe concentration up to 5 μM , MB DP voltammetric signal is increased; but at higher PNA probe coverage (probe concentration greater than 7 μM), a decrease in the MB reduction signal was observed. We attribute this effect to a decrease in the number of triplexes formed on the electrode surface due to the steric and electrostatic hindrances arising from the more tightly packed PNA monolayer at higher surface coverage as described earlier [36, 39–41].

The influence of the hybridization time on the hybridization efficiency was also studied and results are shown in Fig. 3. As displayed in the inset in Fig. 3, by increasing the hybridization time up to 20 h, MB reduction signal increases and then levels off for more hybridization times. Therefore 20 h is suggested as the optimum time for the hybridization; however, as with other sensing procedures, the hybridization time may be reduced to 4–5 h, because under these conditions about 65–70% of maximum hybridization is achieved and is enough for the present biosensor to function correctly, while the selectivity and sensitivity of determination remain nearly unchanged.

Sensor performance

Selectivity

Several hybridization experiments were carried out to assess whether the developed DNA sensor responds selectively to the target ds-DNA. For this purpose, 1 nM of complementary ds-oligonucleotides (ds-HCV3a), two different non-complementary ds-oligonucleotides—ds-UGT and ds-HCV1a, and a mixture of them and a mixture of complementary dsHCV3a and two non-complementary ds-oligonucleotides were used. As seen in Fig. 4, the interaction between the non-complementary oligonucleo-

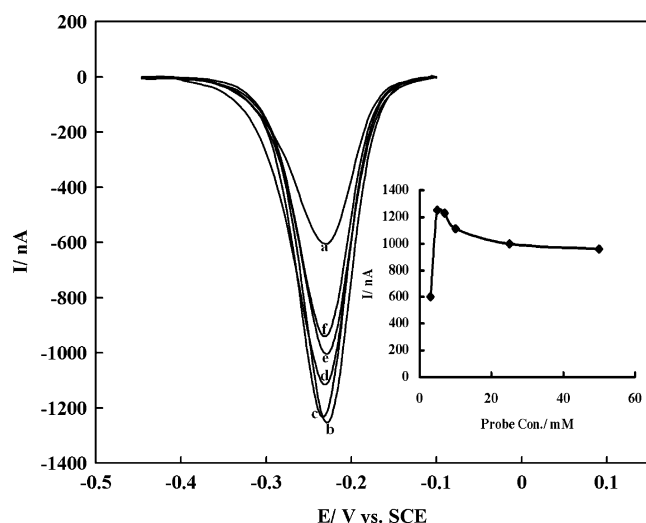


Fig. 2 Differential pulse voltammograms of accumulated MB on PNA probe-modified AuE using various probe concentrations, namely **a** 3, **b** 5, **c** 7, **d** 10, **e** 25, and **f** 50 μ M, after hybridization with 5 μ M ds-HCV3a target oligonucleotide solution. *Inset* variation of the MB peak current versus probe solution concentration

tides and self-assembled probe did not lead to a significant increase in the MB reduction signal (curves **c**, **d**, and **e**) due to the absence of significant hybridization between the probe and the non-complementary ds-DNAs (compare curves **c**, **d**, and **e** with curve **a**). As shown also in Fig. 4 the MB signals for the hybridized complementary ds-DNA alone (curve **b**) and in the presence of two non-complementary ds-DNA (curve **f**) are almost the same (compare curve **b** with curve **f**). This

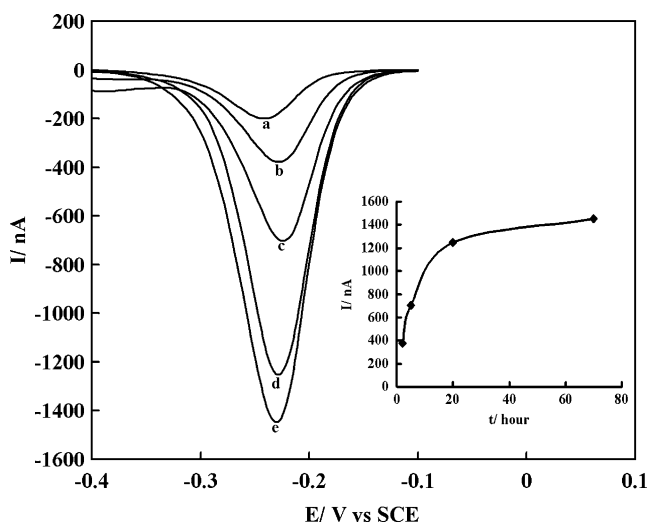


Fig. 3 Differential pulse voltammograms of accumulated MB on the PNA probe-modified AuE before (**a**) and after hybridization with 5 μ M ds-HCV3a target oligonucleotide at various hybridization times: **b** 2, **c** 5, **d** 20, and **e** 72 h. *Inset* variation of the MB peak current versus hybridization time. Electrode conditions: a 6- μ l droplet of 5 μ M PNA solution was dropped onto the pre-treated AuE

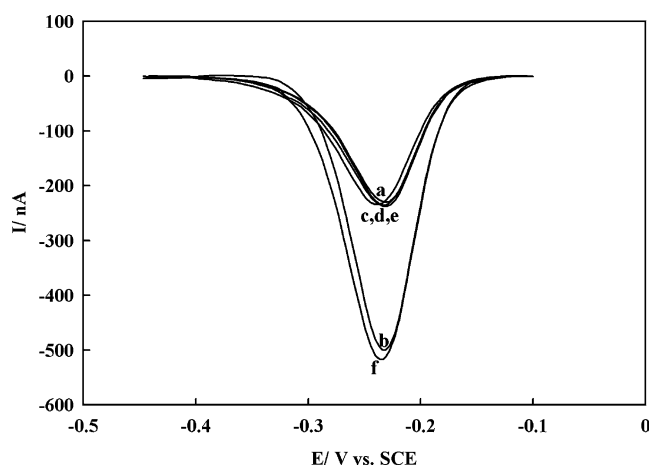


Fig. 4 Differential pulse voltammograms of accumulated MB on the PNA probe-modified AuE after hybridization with: **a** blank buffer solution; 1 nM of **b** ds-HCV3a complementary oligonucleotides and **c** ds-UGT; **d** ds-HCV1a; **e** mixed solution of ds-UGT and ds-HCV1a as ds-non-complementary oligonucleotides; and **f** after hybridization with ds-HCV3a in a mixture solution of ds-HCV3a and two non-complementary oligonucleotides each with same concentration of 1 nM. Electrode conditions: as in Fig. 3

result indicates that only the complementary ds-DNA could form a triplex on the probe self-assembled AuE. Thus the biosensor can detect fully matched ds-DNA in a mixed solution of complementary and non-complementary ds-oligonucleotides, and high selectivity of this new strategy (ds-DNA complementary detection) is confirmed.

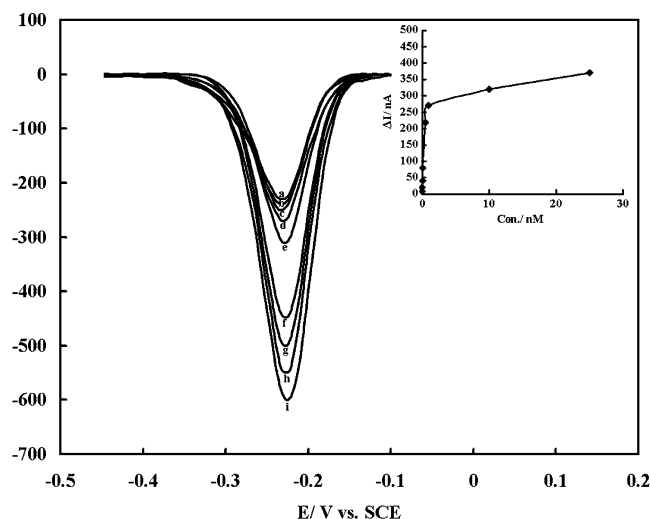


Fig. 5 Differential pulse voltammograms of accumulated MB on the PNA probe-modified AuE after hybridization with ds-HCV3a target oligonucleotide of different concentrations: **a** 0, **b** 10, **c** 25, **d** 50, **e** 100, **f** 500, **g** 1,000, **h** 10,000, **i** 25,000 pM. *Inset* variation of the difference between the MB reduction signals of the PNA probe-modified AuE in the presence and absence of ds-HCV3a target sample (ΔI) versus target ds-DNA concentration. Electrode conditions: as in Fig. 4

Sensitivity

The MB reduction signal on the probe-modified Au electrode in the presence of ds-HCV3a target is increased with increasing the ds-target concentration (Fig. 5). As seen in the inset in Fig. 5 the difference between the MB reduction signal of the probe-modified AuE in the presence and in the absence of ds-target sample (ΔI) increases with increasing ds-target concentration and levels off at ca. 1 nM. Thus, in the presence of this concentration of the ds-target, the maximum capacity of the probe available on the electrode surface is involved in the hybridization event. The calibration graph is linear between 10 and 100 pM with regression equation $y \Delta I = 30.6 + 0.79 X \text{ (pM)}$ and correlation coefficient of 0.999. The detection limit, calculated by means of the following equation: $y_{\text{LOD}} \Delta I = y_B + 3S_{y/x}$ and the regression equation was about 1.8×10^{-12} M, where y_B is the signal of the blank (here intercept of calibration graph) and $S_{y/x}$ is the standard deviation of the blank (here standard deviation of the calibration graph). The relative standard deviation of measurements of 100 pM of ds-HCV3a performed with three independent probe-modified electrodes was 3.1%, indicating a remarkable reproducibility of the detection method.

Conclusion

The gold electrode modified with a SAM composed of the PNA probe and MCH was prepared and employed for the direct detection of ds-oligonucleotide corresponding to HCV3a via triplex formation. The enhancement of the peak current arising from the hybridization of the PNA probe with the target HCV3a ds-oligonucleotide indicates that more MB molecules intercalate within the triple-helix of PNA–ds-DNA. This provides the possibility of monitoring the hybridization of the PNA probe with the target ds-DNA. The present strategy could be considered as a fundamental platform for the development of further DNA detection techniques.

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