STUDY OF THE PROPERTIES OF BIOTIN-STREPTAVIDIN SENSITIVE BIOFETS

Thomas Windbacher, Viktor Sverdlov, Siegfried Selberherr
Institute for Microelectronics, TU Wien, Gußhausstraße 27–29/E360, A-1040 Wien, Austria
Windbacher@iue.tuwien.ac.at, Sverdlov@iue.tuwien.ac.at, Selberherr@iue.tuwien.ac.at

Clemens Heitzinger, Norbert Mauser
Wolfgang Pauli Institute and Department of Mathematics, University of Vienna, Nordbergstrasse 15, A-1090 Wien, Austria
Clemens.Heitzinger@univie.ac.at, Norbert.Mauser@univie.ac.at

Christian Ringhofer
Department of Mathematics, Arizona State University, Tempe, AZ 85287, U.S.A.
Ringhofer@asu.edu

Keywords: BioFET, Field-effect biosensor, Biotin-streptavidin, Simulation, Multi-scale problem, Interface conditions.

Abstract: In this work the properties of a biotin-streptavidin BioFET have been studied numerically with homogenized boundary interface conditions as the link between the oxide of the FET and the analyte which contains the bio-sample. The biotin-streptavidin reaction pair is used in purification and detection of various biomolecules; the strong streptavidin-biotin bond can also be used to attach biomolecules to one another or onto a solid support. Thus this reaction pair in combination with a FET as the transducer is a powerful setup enabling the detection of a wide variety of molecules with many advantages that stem from the FET, like no labeling, no need of expensive read-out devices, the possibility to put the signal amplification and analysis on the same chip, and outdoor usage without the necessity of a lab.

1 INTRODUCTION

Today’s technology for detecting tumor markers, antigen-antibody complexes, and pathogens is time-consuming, complex, and expensive (Pirrung, 2002), (Shinwari et al., 2006). For instance, a typical procedure to detect a given DNA complex is to increase the concentration by RT (reverse transcription) or PCR (polymerase chain reaction), followed by a process step that will add a label to the DNA enabling detection by light or radiation. After all these steps the sample is applied to a microarray. The microarray consists of an array of spots, and every single spot is able to detect a different type of molecule. After the reaction has taken place the array is read by an expensive microarray reader.

Replacing the above sensing mechanism by an electrical detection has several benefits. First, the optical microarray reader becomes superfluous. Detection by FET (field-effect transistor) makes the integration of amplifying and analyzing circuits on the same chip possible, thus saving also equipment. The advanced development of semiconductor process technology allows mass production of such devices, decreasing the price dramatically. Various kinds of reaction pairs are possible and have been studied, like detection of DNA (Fritz et al., 2002), (Hahm and Lieber, 2004), (Gao et al., 2007), cancer markers (Zheng et al., 2005), proteins, e.g. biotin-streptavidin (Im et al., 2007), (Cui et al., 2001), (Gupta et al., 2008), (Stern et al., 2007), albumin (Park et al., 2008), and transferrin (Girard et al., 2006). In these papers different device types and materials were investigated and provided different solutions for each problem. In principle, every molecule that is charged in the solute and that can be bound to the surface layer can be detected by a BioFET. The field of applications is very wide and spans from DNA sequencing, point of care applications, to controlling environmental pollution and the spread of diseases. The BioFET can be easily integrated into the chip environment. By putting a microfluidic channel above the functionalized gate of the
BioFET the chip can be turned into a mini-laboratory - the lab on chip. This enables better control of the environmental parameters (e.g. local pH or detecting the amount of a special protein) and gives the possibility of local measurement (e.g. how a cell reacts to a stimulus), thus providing a complete lab-on-a-chip. However, there are still many problems to overcome and a lot of research is needed. For instance, an interesting way to avoid problems by poor isolation between device and solution has been shown by (Kim et al., 2006).

2 METHOD

A BioFET consists of several parts: a semiconductor transducer, a dielectric layer, a biofunctionalized surface, and the analyte (Figure 1). The semiconductor transducer is a conventional FET. The dielectric surface, and the analyte (Figure 1). The semiconductor device changes and thus the potential in the semiconductor is in an aqueous solution. If a target molecule is attached, so it is able to bind the desired molecule. The face contains immobilized biomolecule receptors at a stimulus), thus providing a complete lab-on-a-chip.

The binding of the target with the receptor happens at the Angstrom length scale, while the semiconductor is changed.

The binding of the target with the receptor happens at the Angstrom length scale, while the semiconductor device is in the micrometer length scale. Thus a proper way of combining the semiconductor-solution interface is crucial.

Transport in a FET with a gate length of \( L_{\text{g}} \), is usually modeled via the drift-diffusion approach (Tang and Ieong, 1995), (Selberherr, 1984). The aqueous solution is described by the Poisson-Boltzmann equation.

\[
\varepsilon_0 \nabla \cdot (\varepsilon_{\text{Ana}} \nabla \psi(x,y)) = - \sum_{\sigma \in S} \sigma q \varepsilon_0 \varepsilon_{\text{Ana}} \varepsilon_{\text{Oxid}} \left( \frac{\partial}{\partial y} \frac{\partial \psi(x,y)}{\partial y} \right)
\]  

(1)

\( k_B \) denotes Boltzmann’s constant, \( T \) the temperature in Kelvin, and \( \sigma \in S \), where \( S \) contains the valences of the ions in the electrolyte. \( \varepsilon_0 \) describes the permittivity of vacuum, and \( q \) the elementary charge. \( \psi_p \) is the chemical potential. \( \varepsilon_{\text{Ana}} \) is the ion concentration in equilibrium, while \( \varepsilon_{\text{Ana}} \approx 80 \) is the relative permittivity of water.

The sum describes the carrier densities arising from the Boltzmann model. Assuming sodium-chloride as salt, which is a 1 : 1 salt, the expression given in (1) can be reduced to

\[
\varepsilon_0 \nabla \cdot (\varepsilon_{\text{Ana}} \nabla \psi(x,y)) = 2q \varepsilon_0 \varepsilon_{\text{Ana}} \sinh \left( \frac{q}{k_B T} (\psi(x,y) - \psi_p) \right).
\]

(2)

The charge on the surface due to chemical reaction of the \( H^+ \) and \( OH^- \) was modeled at \( pH = 7 \) with the site-binding model (Shinwari et al., 2006):

\[
Q_{Ox} = q N_s \left[ \frac{[H^+]_{\text{b}}}{K_a} e^{-\frac{q}{k_B T} \psi(x,y)} - \frac{K_b}{[H^+]_{\text{b}}} e^{\frac{q}{k_B T} \psi(x,y)} \right].
\]

(3)

\( N_s \) denotes the surface binding site density, while \( K_a \) and \( K_b \) are the equilibrium constants for charging the surface positively and negatively, respectively. \([H^+]_{\text{b}}\) describes the positive hydrogen ion concentration of the bulk and is corrected to the activity of the hydrogen concentration by the \( e^{\frac{q}{k_B T} \psi(x,y)} \) terms.

The biomolecules are modeled in a physics-based bottom-up approach. By calculating the charge and dipole moment for a single molecule (see for example Figure 2, (Poghossian et al., 2005)), a mean charge density and a mean dipole moment density of the boundary layer is obtained. This bridges the gap between the Angstrom length scale of the biomolecules and the micrometer dimensions of the FET (Heitzinger et al., 2008a), (Heitzinger et al., 2008b), (Ringhofer and Heitzinger, 2008), (Windbacher et al., 2008).

The link between the gate oxide and the aqueous solution is realized by two interface conditions, (Heitzinger et al., 2008a), (Heitzinger et al., 2008b), (Ringhofer and Heitzinger, 2008), (Heitzinger and Klimeck, 2007),

\[
\varepsilon_0 \varepsilon_{\text{Oxid}} \frac{\partial}{\partial y} \psi(0-,x) - \varepsilon_0 \varepsilon_{\text{Ana}} \frac{\partial}{\partial y} \psi(0+,x) = -C(x),
\]

(4)

\[
\psi(0-,x) - \psi(0+,x) = -\frac{D_x(x)}{\varepsilon_{\text{Ana}} \varepsilon_0}.
\]

(5)

The x-axis is parallel oriented to the oxide surface, while the y-axis points into the liquid. \( \psi(0-) \) describes the potential in the oxide, while \( \psi(0+) \) relates to the potential in the solute. The first equation describes the jump in the field, while the second introduces a dipole moment which causes a shift of the
Figure 2: Biotin-streptavidin complex (http://www.pdb.org) on the oxide surface. Two iso-surfaces for plus and minus $0.03 \text{eV}^{-1}$ are shown.

Figure 3: Potential profile at the interface (from left to right: semiconductor, oxide, solute).

potential taken into account by adjusting the potential in the analyte (Figure 3).

3 SIMULATION

Three different types of dielectric were simulated. SiO$_2$ as a reference, Al$_2$O$_3$, and Ta$_2$O$_5$ as possible high-k materials, with relative permittivities of 3.9, 10, and 25 respectively. As solute 1mMol sodium-chloride at pH = 7 was considered. The parameters for the site-binding model can be found in Table 1 (Landheer et al., 2005). For each dielectric the unprepared state (just water and salt), the prepared state (water, salt, and biotin), and the bound state when the chemical reaction has taken place (water, salt, and biotin-streptavidin) were calculated for two different mean distances between molecules ($\lambda = 10\text{nm}$, $\lambda = 15\text{nm}$). The data used for calculating charge and dipole moment of biotin and streptavidin are obtained from http://www.pdb.org (1SEW.pdb, Figures 2, 12). The potential distribution across the device is shown in Figure 4 and output curves were calculated for every parameter combination mentioned above, assuming a 100% binding efficiency. The potential of the reference electrode is set to 0.4V so that the FET will be in moderate inversion as proposed by (Deen et al., 2006).

Table 1: The parameters needed for the site-binding model using different dielectric.

<table>
<thead>
<tr>
<th>Oxide</th>
<th>$pK_a$</th>
<th>$pK_b$</th>
<th>$N_s$ [cm$^{-2}$]</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>SiO$_2$</td>
<td>-2</td>
<td>6</td>
<td>$5 \cdot 10^{14}$</td>
<td>(Bousse, 1982)</td>
</tr>
<tr>
<td>Al$_2$O$_3$</td>
<td>6</td>
<td>10</td>
<td>$8 \cdot 10^{14}$</td>
<td>(Bousse, 1982)</td>
</tr>
<tr>
<td>Ta$_2$O$_5$</td>
<td>2</td>
<td>4</td>
<td>$10 \cdot 10^{14}$</td>
<td>(Bousse et al., 1991)</td>
</tr>
</tbody>
</table>

4 RESULTS

Figures 5, 6, and 7 show a decrease in the output current for biotin attached to the surface in comparison to the unprepared surface. This downward shift for the bound state in comparison to the unbound state is due to the increase of negative charges at the interface, which is also confirmed by the difference between the curves for $\lambda = 10\text{nm}$ and $\lambda = 15\text{nm}$, since for 10nm the molecules are more dense than by 15nm.

As can be seen in the Figures 5, 6, and 7 the bigger the $\varepsilon_r$ of the dielectric the bigger is the output current. Thus high-k materials deliver stronger output signals. According to (Deen, 2007) however, higher $\varepsilon_r$ dielectric constants may lead to higher trap densities and thus to a decreased signal-to-noise ratio. Therefore a trade-off between bigger output signal and signal-to-noise ratio has to be met. Figure 8 shows the output
In the biochemical community there is an ongoing discussion, if the orientation of the biomolecule is relevant for sensing. Several papers have shown contradictory results (Oh et al., 2005), (Wacker et al., 2004), (Kusnezow et al., 2003), (Peluso et al., 2003), (Turkova, 1999). All these papers are based on optical detection. Although more study is needed, we mention that for optical detection it is more important to choose the linking molecule in a way that the reaction is not hindered by steric effects (receptors block each other) or the binding sites are blocked or even broken by the crosslinker. In the case of a BioFET, however, a field-effect as working principle is used. Thus it is important to have a linker that is as short as possible, to be close to the surface. To increase the signal-to-noise ratio, the linker should have as little charge as possible. For example, in order to detect streptavidin,
biotin is used as a binding agent. A biotin molecule is attached to the surface with a neutral linker. Streptavidin then binds to biotin thus forming a bound state.

The charge difference between the unbound state of a biotin alone, which is negatively charged with a single elementary charge and the bound state of biotin-streptavidin, which is negatively charged with five elementary charges, is large enough for detection. We also note that due to the tetrameric nature of streptavidin it has four sites to bind biotin as shown in Figure 12. Therefore, the linker binding biotin to the surface should be short enough in order to prevent binding several biotin molecules to a single molecule of streptavidin.

5 CONCLUSIONS

The model shows a strong dependence on surface charges and indicates a detectable shift in the threshold voltage depending on their orientation related to the surface. The bound state (streptavidin-biotin) negatively charged with five elementary charges compared to the unbound state (biotin) negatively charged with one elementary charge leads to a reduced conductivity, when hybridization has taken place. Also the shift of the threshold voltage and output characteristics due to different molecule orientations (0°...perpendicular to surface, 90°...lying flat on surface) can be seen. This shows the usefulness of the simulation method for the design of efficient BioFETs.

Figure 9: Potential profile for biotin-streptavidin at $\lambda = 10\,\text{nm}$ from left (semiconductor) to right (oxide).

Figure 10: Small signal resistance for SiO$_2$, Al$_2$O$_3$, and Ta$_2$O$_5$ for calculation without dipole moment, angle 0° (perpendicular to surface), and angle 90° (parallel to surface) at biotin only.

Figure 11: Small signal resistance for SiO$_2$, Al$_2$O$_3$, and Ta$_2$O$_5$ for calculation without dipole moment, angle 0° (perpendicular to surface), and angle 90° (parallel to surface) at bound state (biotin-streptavidin).

Figure 12: Scheme of the tetrameric protein streptavidin and biotin.
REFERENCES


